

Pairing Nature with Scientific Discoveries

Product Catalog Volume IV



Greetings from Vivantis Technologies

It is our great pleasure to bring you our latest version of Vivantis catalog.

Sincerely,

EL Law.
CEO of Vivantis Technologies



Contact



MALAYSIA (Headquarters) Telephone

+6 03 8025 1603

Fax

+6 03 8025 1637

Email

info@vivantechnologies.com, international@vivantechnologies.com

Mail

Vivantis Technologies Sdn. Bhd.
Revongen Corporation Center
No.12A, Jalan TP 5,
Taman Perindustrian UEP,
47600 Subang Jaya,
Selangor Darul Ehsan, Malaysia.

Website

www.vivantechnologies.com



"Nature is man's teacher" - Alfred Billings Street

There is a vast ocean of knowledge, likely infinite, to be learned from the natural world that surrounds us. As an indispensable tool, Science helps us to understand and appreciate the works of nature, from animal behaviour to weather patterns and more. It allows us to delve in the past, connects us with the present and helps us in predicting the future. The union of science and nature with creativity and imagination forms the scientific enterprise and fuels the scientific endeavour.

From the arduous effort of scientists in the past and present, we know that knowledge gained from studying one aspect of nature is applicable to other parts though seemingly unrelated. If one looks hard enough, one would notice the intricate patterns and connections nature has to offer, which ultimately contributes to human advancement. Science can be considered a continuous process for gathering knowledge through careful observations of nature. Scientists assume that whilst total and absolute truth may not be within our reach, increasingly precise estimates can be made about the world around us and how it works.

Nowadays, scientific discoveries are not complete without technology in the equation. Technology has played a vital role in driving and pushing the scientific boundaries to achieve discoveries we never dreamed of becoming reality. Nature is constantly inspiring technological advances and our coexistence on this planet should not be taken for granted. We should respect, appreciate and harvest nature's bounty wisely.



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Acquaint Us:

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info@vivantechnologies.com international@vivantechnologies.com

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Vivantis reserves the right to make changes to these Terms & Conditions, at any time without notice.

New Products

2X OneStep Taq ReverseTrans **PCR** Master Mix

Description

2X OneStep Tag ReverseTrans PCR Master Mix offers rapid and sensitive end-point detection of RNA templates in a single step. 2X OneStep Tag ReverseTrans PCR Master Mix is an optimized ready-to-use 2X concentrated RNA amplification mixture containing M-MuLV Reverse Transcriptase, RNase Inhibitors, Tag DNA Polymerase, reaction buffer and dNTPs. It contains all the components required for routine RNA amplification except template and primers. M-MuLV Reverse Transcriptase has the absence of RNase H activities that enhance the synthesis of long cDNAs and amplification of long transcripts. 2X OneStep Tag ReverseTrans PCR Master Mix allows cDNA synthesis and PCR to be performed using only gene-specific primers.

Features

- Saves time and reduces contamination due to reduced number of tests and pipetting steps
- Stable at 4°C for 6 months, allowing immediate reaction setup without the time-consuming thawing of reagent
- Suitable for all routine RNA amplification applications



Ordering Information

Catalog No	Description	Pack Size
RTMM01	2X OneStep Taq ReverseTrans PCR Master Mix	100 applications

2X ViRed OneStep Taq ReverseTrans **PCR Master Mix**

Description

2X ViRed OneStep Tag ReverseTrans PCR Master Mix offers rapid and sensitive end-point detection of RNA templates in a single step. 2X ViRed OneStep Tag ReverseTrans PCR Master Mix is an optimized ready-to-use 2X concentrated RNA amplification mixture containing M-MuLV Reverse Transcriptase, RNase Inhibitors, Tag DNA Polymerase, reaction buffer and dNTPs. It contains all the components required for routine RNA amplification except template and primers. M-MuLV Reverse Transcriptase has the absence of RNase H activities that enhances the synthesis of long cDNAs and amplification of long transcripts. 2X ViRed OneStep Tag ReverseTrans PCR Master Mix allows one-step RT-PCR using only gene-specific primers. 2X ViRed OneStep Tag ReverseTrans PCR Master Mix contains the inert red dye and stabilizers that allow direct loading of final PCR products onto gels for electrophoresis. The red color dye migrates at approximately 400bp on 1% agarose gel in 1X TBF Buffer.

Features

- Suitable for all routine RNA amplification applications
- · Reduces set-up time and buffer-dye mixing
- Minimizes potential contamination due to reduced number of tests and pipetting steps
- Easy confirmation of complete mixing
- No additional loading dye needed direct loading of final products



Catalog No	Description	Pack Size
RTMM02	2X ViRed OneStep <i>Taq</i> ReverseTrans PCR Master Mix	100 applications

New Products

Tricolor Broad Range Prestained **Protein Ladder**

140 100 75 60 For migration pattern & approximate molecular weight usina Bis-Tris 8-15% in MOPS Buffer and MES Buffer, kindly refer to website for more information Glycine

Description

Tricolor Broad Range Prestained Protein Ladder contains 13 proteins that resolve into sharp, tight bands in the range of 5-245kDa.lt can be used to monitor molecular weight separation during electrophoresis, estimate molecular weights of proteins of interest, and evaluate western transfer efficiency. It contains 2 reference bands ~25 and ~75 kDa coupled with blue chromophore as well as red dye and green dye for easy identification. It can be used on PVDF and nvlon membrane.

Features

Broad Range: 5-245kDa Convenient: Ready to use

Easy Identification: ~25 and ~75kDa reference bands coupled with

blue chromophore, red dye and green dye

Quality Control

Tested in SDS-polyacrylamide gel electrophoresis and western

Storage

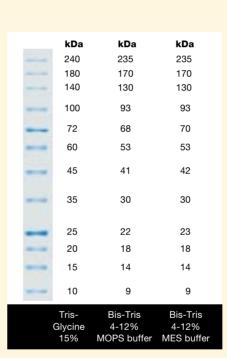
- Stable at -20°C for 2 years
- Stable at 4°C for 6 months



Ordering Information

Catalog No	Description	Pack Size
PR0624	Tricolor Broad Range Prestained Protein Ladder	2 x 250 µl

Whole Blue Range Prestained **Protein Ladder**



Description

Whole Blue Range Prestained Protein Ladder contains 12 proteins that resolve into sharp, tight bands in the range of 10-240kDa.lt can be used to monitor molecular weight separation during electrophoresis, estimate molecular weights of proteins of interest, and evaluate western transfer efficiency. It contains 2 reference bands ~25 and ~72 kDa coupled with blue chromophore for easy identification. It can be used on PVDF and nylon membrane.

Features

Broad Range: 10-240kDa Convenient: Ready to use

Easy Identification: ~25 and ~75kDa reference bands coupled with

blue chromophore

Quality Control

Tested in SDS-polyacrylamide gel electrophoresis and western

Storage

- Stable at -20°C for 2 years
- Stable at 4°C for 6 months



ordoring initialities		
Catalog No	Description	Pack Size
PR0623	Whole Blue Range Prestained Protein Ladder	2 x 250 µl

Viva qGreen I Fluorescent Dye (20x in DMSO)

Description

Viva qGreen I Fluorescent Dye (equivalent to SYBR® Green Dye) is a sensitive green fluorescent nucleic acid dye used for detection of double stranded DNA. The dye is widely used in non-specific detection of amplification in quantitative real-time PCR (qPCR) experiments. The detection is monitored by measuring the increase in fluorescence throughout the cycles.

(equivalent to SYBR® Green Dve)

Features

Easy and affordable

Probes are not required, reduce assay setup and running cost; given that PCR primers are well designed and reaction is well characterized.

Higher sensitivity

Increased fluorescence when bound to any double-stranded DNA.

Stable during storage and under PCR condition, able to withstand repeated freeze-thaw cycles.

Versatile applications

Can be used as a general double stranded DNA binding dye for common DNA quantification, melt curve analysis, etc.

Compatible with most system

Compatible with major brands of qPCR instruments & enzyme

No.	. Color	Name	Туре	Ct	Ct Comment
1		100ng	Standard	5.47	Mean Ct: 5.537
2		100ng	Standard	5.59	
3		100ng	Standard	5.55	
4		10ng	Standard	8.07	Mean Ct: 8.077
5		10ng	Standard	8.03	
6		10ng	Standard	8.13	
7		1ng	Standard	11.63	Mean Ct: 11.653
8		1ng	Standard	11.87	
9		1ng	Standard	11.46	
10		0.1ng	Standard	15.09	Mean Ct: 14.880
11		0.1ng	Standard	14.58	
12		0.1ng	Standard	14.97	
13		0.01ng	Standard	18.34	Mean Ct: 17.567
14		0.01ng	Standard	17.76	
15		0.01ng	Standard	16.66	100

Sample of DNA: Bacteria DNA Test: qPCR test with VIva qGreen I

Figure: Sensitivity of the Viva qGreen I Fluorescent Dye based real-time PCR assay. Amplification plot (cycle number versus fluorescence) of known copies of DNA standard (100ng - 0.01ng) was plotted with three replicates.

Note: SYBR® Green is a registered trademark of Molecular Probes, Inc.

Ordering Information

Catalog No	Description	Pack Size
SD1101	Viva qGreen I Fluorescent Dye 20X in DMSO (equivalent to SYBR® Green Dye)	1ml / pack size
SD1103	Viva qGreen II Fluorescent Dye 20X in Water (equivalent to EvaGreen® Dye)	1ml / pack size

Viva qGreen II Fluorescent Dye

(20x in Water)

(equivalent to EvaGreen® Dve)

Description

Viva qGreen I Fluorescent Dye (equivalent to SYBR® Green Dye)

Redistribution of Dye Molecule

No fluorescence change

Viva gGreen II Fluorescent Dye

(equivalent to EvaGreen® Dye)

Release on Demand

Drop in fluorescence

Figure: Viva qGreen I Fluorescent Dye

quickly rebinds to the regions that remain

double stranded, there is no drop in fluorescence. Viva qGreen II Fluorescent

Dve does not redistribute from the melted regions of single-stranded DNA back to double-stranded DNA, resulting in a reduction of fluorescence. This difference

gives the Viva qGreen II Fluorescent Dye the higher sensitivity in detecting

mplification due to "release on demand

DNA binding mechanism.

Melting Step

OH CHI

OHOHO

HOHICHHAI!

66 Melting Step

Viva qGreen II Fluorescent Dye (equivalent to EvaGreen® Dye) is one of the most sensitive dyes to detect double stranded DNA in quantitative real-time PCR (qPCR) experiments as well as high-resolution DNA melt curve analysis, yielding robust and reproducible results.

Safer

The dye is noncytotoxic & nonmutagenic for safe handling and easy disposal down to drain, completely impermeable to cell membrane.

Higher sensitivity

Low PCR inhibitory and high concentration of dye used for maximal signal and high resolution DNA melt analysis.

Extremely stable

Stable during storage and under PCR condition. No dye decomposition in PCR buffer at 95-100°C for 48 hours. Highly stable under alkaline or acidic condition and able to withstand repeated freeze-thaw cycles

Versatile applications

Used as a general double stranded DNA binding dye for DNA quantification, melt curve analysis and more.

Excellent for qPCR and isothermal application

Brighter and more sensitive than Viva gGreen I Fluorescent Dye (equivalent to SYBR® Green) for detecting amplification due to novel 'release on demand' DNA binding mechanism.

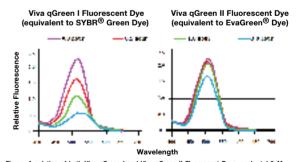


Figure: A solution of both Viva qGreen I and Viva qGreen II Fluc concentration in Tris Buffer was incubated at 99°C. The absorption followed over a period of 3 hours.

Note: EvaGreen® is a registered trademark of Biotium. Inc

Catalog No	Description	Pack Size
SD1101	Viva qGreen I Fluorescent Dye 20X in DMSO (equivalent to SYBR® Green Dye)	1ml / pack size
SD1103	Viva qGreen II Fluorescent Dye 20X in Water (equivalent to EvaGreen® Dye)	1ml / pack size



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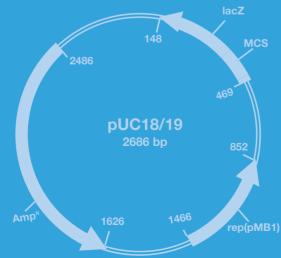
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Vivantis 10-Go™ RE Kits

Icons Description

Restriction Endonucleases

Vivantis Restriction Endonucleases

List of Vivantis Restriction Endonucleases

List of Commercial Isoschizomers

Corresponding to Vivantis Restriction

Endonucleases

Alphabetical List of Recognition Specificities

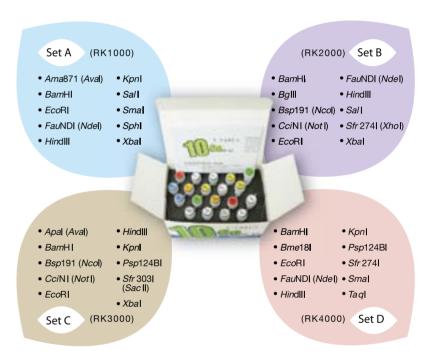
of Vivantis Restriction Endonucleases



Vivantis 10-Go™ RE Kit

Giving scientists time to tango!

A typical freezer in a molecular biology laboratory consists of a host of the quintessential restriction enzymes from various sources. Next come the laborious hunt for different buffer charts to culminate in a compromise of buffers for double digestions. Vivantis presents a solution to this task! Our 10-Go kits consist of selected enzymes for your use, coupled with recommended optimal buffers for double digestions. This is complemented with an unbelievable price performance ratio with the offer of "buy 1, get more". Our 10-Go kits present sets of restriction enzymes to provide you with hassle free tools for your molecular biology applications, saving both time and money!



Catalog No	Description	Pack Size
RK1000	10-Go™ RE Kit Set A	50 - 500u
RK2000	10-Go™ RE Kit Set B	50 - 500u
RK3000	10-Go™ RE Kit Set C	50 - 500u
RK4000	10-Go™ RE Kit Set D	50 - 500u



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Description

Restriction endonucleases are produced in bacteria as a defense mechanism against invasion of foreign DNA derived from viruses. Restriction endonucleases are able to hydrolyze both strands of DNA within or very near to its recognition site. These enzymes generally require divalent metal cation (Mg²⁺) for their activity. Most of the restriction endonucleases recognize hexanucleotide (6) target sites, but others recognize 4, 5 or even 8 nucleotides sequences. Depending on their cleavage position, restriction endonucleases produce either sticky (5' or 3' overhang) or blunt ends.

Restriction endonucleases are categorized into 3 types: Type I, II and III. Both types I and III are seldom used by molecular biologists due to its specific requirement for ATP and the fact that it cleaves DNA at a substantial distance from its recognition site. Type II enzymes, however, do not require ATP and generally cleave within or near its recognition site. Due to its ability to cleave at specific sites to produce defined fragments, these enzymes are now necessary tools in molecular biology. Currently, there are more than 3000 type II restriction endonucleases, exhibiting over 200 different specificities, many of which are now commercially available.

Color-Tag Buffer System

Our Color-Tag buffers are supplied at 10X concentration together with the restriction endonucleases. The buffers are stored in color-coded tubes corresponding in color to the cap of its restriction endonuclease storage tube. This Color-Tag buffer system ensures convenience and highest performance. Our buffers have been subjected to stringent analysis for maximum use across our entire line of restriction endonucleases.

Buffers should be stored at -20°C for long term storage. Buffers kept at +4°C should be aliquoted. Our restriction endonucleases perform 100% of its certified activity in the recommended buffer provided.

Factors Influencing Restriction Endonuclease Performance

Star Activity

'Star Activity' is a term used for an altered cleavage which occurs when a restriction endonuclease is under non-standard conditions. In cases like this, restriction endonuclease cleavage sequences are similar but not identical to their defined recognition sequence. Normally this can be seen with high enzyme concentrations and buffers that deviate from the recommended conditions. In most cases, star activity may be caused by high glycerol concentration in the reaction mixture or presence of other organic solvents, such as ethanol or low ionic strength or high pH values in reaction buffer or substitution of cofactor Mg²⁺ with other divalent cations such as Mn²⁺.

Buffer System of Restriction Endonucleases



1X Buffer V1 10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, and 100µg/ml BSA.



1X Buffer V2 10mM Tris-HCI (pH7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100μg/ml BSA.



1X Buffer V3 50mM Tris-HCI (pH7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100ua/ml BSA.



1X Buffer V4 10mM Tris-HCI (pH8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100μg/ml BSA.



1X Buffer V5 30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA.



0.5X Buffer UB 12.5mM Tris-acetate (pH7.6 at 30°C), 5mM Mg-acetate, 50mM K-acetate, 3.5mM 2-mercaptoethanol, and 25µg/ml BSA.

1.0X Buffer UB 25mM Tris-acetate (pH7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol, and 50µg/ml BSA.

1.5X Buffer UB 37.5mM Tris-acetate (pH7.6 at 30°C), 15mM Mg-acetate, 150mM K-acetate, 10.5mM 2-mercaptoethanol, and 75µg/ml BSA.

2.0X Buffer UB 50mM Tris-acetate (pH7.6 at 30°C), 20mM Mg-acetate, 200mM K-acetate, 14mM 2-mercaptoethanol. and 100µg/ml BSA.

Dam and Dcm Methylation

Restriction endonucleases are sensitive to different types of modified bases occurring Restriction Endonucleases (cont'd) in the DNA sequences. Methylation of cytosine to 5'-methylcytosine (mC), adenine to N6-methyladenine (mA) in, or adjacent to the site recognized by a restriction endonuclease may prevent hydrolysis. All restriction endonucleases produced by Vivantis have been examined for sensitivity to dam and dcm. An icon marks known methylation effects in the individual listing of restriction endonucleases.

Quality Control Test

A series of quality control tests have been performed to ensure our restriction endonucleases are well suited for research purposes.

Unit Determination

1u is defined as the amount of enzyme that is required to digest 1µg of substrate DNA to completion in 1 hour in a total reaction volume of 50µl, with appropriate assay conditions (salt concentration, pH and temperature). Please note that the activity of the restriction endonuclease is substrate-dependent. The enzyme should be titrated to determine its actual activity when working with a new substrate.

Overdigestion Assay

The absence of detectable levels of non-specific nucleases is demonstrated by incubating different amounts of restriction endonuclease for 16 hours with 1µg of substrate DNA under optimum assay conditions. The banding pattern generated must be identical to the normal banding pattern produced at 1 hour digestion of the enzyme being tested.



Agarose Gel Photo (0.7%) Lambda DNA digested with 10u of Aatll for 1 hour (a), 16 hours (b)

Ligation and Recutting Assay

This assay is to demonstrate the absence if detectable levels of phosphatase and exonucleases. DNA fragments are produced by an excessive over-digestion of substrate DNA with each restriction endonuclease. These fragments are then ligated with T4 DNA Ligase. The ligated fragments are then recut with the same restriction endonuclease. Ligation can only be occur if the 5' and 3' termini are left intact, and only those molecules with a perfectly restored recognition site can be recleaved. A normal banding pattern after cleavage indicates that both the 5' and 3' termini are intact and that the enzyme preparation is free of detectable phophatases and exonucleases.

5'[32P]-Labeled Oligonucleotide Assay

The labeled oligonucleotide assay allows for the identification of exonuclease contaminants during restriction endonuclease preparation and ensures the purity and quality of our restriction endonucleases. This test is performed by setting up a restriction endonuclease reaction for 3 hours with 1µl of restriction endonuclease incubated with a 5'-[32P] labeled synthetic oligonucleotide (single-and doublestranded) that has no recognition site for the test restriction endonuclease. The products are separated on a polyacrylamide gel and the results are analyzed by an imaging system. The absence of degradation products signifies the purity of the restriction endonuclease.

Buffer System of



1X Buffer AccB1I 10mM Tris-HCI (pH7.5 at 25°C), 10mM MgCl, 100mM KCl, and 100µg/ml BSA.

1X Buffer Ama87I 10mM Tris-HCI (pH8.5), 10mM MgCI, 150mM NaCl, 100µg/ml BSA.

1X Buffer Ball 20mM Tris-HCI (pH8.5), 10mM MgCl, 200mM NaCl, and 1mM DTT.

1X Buffer Bsp131 10mM Tris-HCI (pH7.6), 10mM MaCI, 200mM KCl, and 100µg/ml BSA.

1X Buffer DrallI 10mM Tris-HCI (pH7.6), 10mM MgCI, 200mM KCl, and 100µg/ml BSA.

1X Buffer EcoRI 50mM Tris-HCI (pH7.5 at 30°C). 10mM MgCl, 100mM NaCl, 0.02% Triton X-100, and 100µg/ml BSA.

1X Buffer EcoRV 10mM Tris-HCI (pH8.5 at 30°C). 10mM MgCl, 100mM NaCl, and 100ua/ml BSA. 1X Buffer Mboll 33mM Tris-acetate (pH7.6 at 30°C), 10mM Mg-acetate, 66mM K-acetate, and 1mM DTT.

1X Buffer Ssol 10mM Tris-HCI (pH7.6 at 30°C), 10mM MgCl, 100mM KCl, and 100µg/ml BSA.

1X Buffer Bbv 20mM Tris-HCI (pH8.5), 10mM MgCl, 20mM NaCl and 1mM DTT. Incubate at 37°C.

1X Buffer Bsp 20mM Tris-HCI (pH8.5), 10mM MgCl, 200mM NaCl and 1mM DTT.



Agarose Gel Photo (0.7%) Lambda DNA digested with EcoRI (a), fragments ligated with T4 DNA Ligase (b), ligated DNA redigested with EcoRI (c)



List of Vivantis Restriction Endonucleases

Vivantis Restriction Endonuclease	Prototype	Recognition Sequence / Cleavage Site 5'→3'	Product No	Page
Aat II	Aat II	GACGT↓C	RE1100	033
Acc 16I	Mst I	TGC↓GCA	RE1102	033
Acc 651	Kpn I*	G [↓] GTACC	RE1104	034
Acc B1I	Hgi CI	G [↓] GYRCC	RE1106	034
Acc B7I	Pf/ MI	CCANNNN↓NTGG	RE1108	035
Acc BSI	Bsr BI	CCGCTC (-3 / -3)	RE1110	035
Ac/ I	Ac/I	AA↓CGTT	RE1112	036
Acs l	Apol	R↓AATTY	RE1114	036
Afi I	Bsi YI	CCNNNNN↓NNGG	RV1116	037
Ahl l	Spel	A↓CTAGT	RE1118	038
Alul	A/u l	AG↓CT	RE1120	038
Ama 87 I	Ava l	C [↓] YCGRG	RE1122	039
Apa I	Apa l	gggcc↓c	RE1124	040
AsiG I	Age l	A↓CCGGT	RE1126	041
Asp A2I	AvrII	C↓CTAGG	RE1128	041
AspLEI	Hha I	gcg↓c	RE1130	042
Asp S9I	Asu l	G↓GNCC	RE1132	042
Asu HPI	Hph I	GGTGA (8/7)	RE1134	043
Asu NHI	Nhe I	G↓CTAGC	RE1136	043
Bam H I	Bam HI	G↓GATCC	RV1138	044
Bbv 12I	Hgi Al	gwgcw↓c	RE1140	045
Bgl l	Bg/ l	GCCNNNN [↓] NGGC	RV1142	046
Bg/ II	Bg/ II	A↓GATCT	RV1144	046
Bmc Al	Scal	AGT↓ACT	RV1146	047
Bme181	Avall	g↓gwcc	RE1148	047
Bme RI	Eam11051	GACNNN↓NNGTC	RV1150	048
Bmi I	Nla IV	GGN↓NCC	RV1152	048
Bmr FI	ScrFI	CC↓NGG	RV1154	049
Bmt I	NheI*	GCTAG↓C	RE1156	049
Bpu 10 I	<i>Bpu</i> 10I	CCTNAGC (-5/-2)	RE1158	050
Bpu 14 I	Asu II	TT [↓] CGAA	RE1160	051
Ври МІ	Cau II	cc↓sgg	RV1162	051
Bse 1I	Bsr l	ACTGG (1/-1)	RE1166	053
Bse118I	Cfr 10I	R↓CCGGY	RE1168	053
Bse 21I	Sau I	CC [↓] TNAGG	RE1170	054

List of Vivantis Restriction Endonucleases (cont'd)

Vivantis Restriction Endonuclease	Prototype	Recognition Sequence / Cleavage Site 5'→3'	Product No	Page
Bse 3DI	BsrD I	GCAATG (2/0)	RE1172	054
Bse8I	Bsa BI	GATNN [↓] NNATC	RE1174	055
Bse PI	BsePI	g↓cgcgc	RE1176	055
Bse X3I	Xma III	c∮ggccg	RE1178	056
Bsh VI	Cla I	AT↓CGAT	RV1180	056
Bsn l	HaeIII	gg↓cc	RV1182	057
Bso 31I	Eco31I	GGTCTC (1/5)	RE1184	058
Bsp13I	BspMII	T↓CCGGA	RE1186	058
Bsp1720I	Espl	GC [↓] TNAGC	RE1188	059
Bsp 19 I	Ncol	C↓CATGG	RE1190	059
Bss MI	Mbol	↓GATC	RV1192	060
Bss NI	Acy l	GR↓CGYC	RV1194	061
Bss NAI	Snal	GTA [↓] TAC	RE1196	061
Bss T1I	Sty I	c↓cwwgg	RE1198	062
Bst 2UI	EcoRII*	cc↓wgg	RE1202	063
Bst 4CI	Tsp 4CI	ACN↓GT	RE1204	063
Bst 6I	Ksp 632I	CTCTTC (1/4)	RE1206	064
BstAUI	Bsp1407I	T [↓] GTACA	RE1208	064
BstBAI	Bsa AI	YAC [↓] GTR	RE1210	065
Bst DEI	Dde l	C↓TNAG	RE1212	065
Bst DSI	Dsa l	C↓CRYGG	RE1214	066
Bst ENI	Eco NI	CCTNN [↓] NNNAGG	RE1216	066
Bst F5I	Fok ! *	GGATG (2/0)	RE1218	067
Bst FNI	Fnu DII	cg↓cg	RE1220	067
Bst H2I	Haell	RGCGC↓Y	RE1222	068
Bst HHI	Hha I	gcg↓c	RE1224	068
Bst MAI	Bsm Al	GTCTC (1/5)	RE1226	069
Bst MBI	Mbo l	↓GATC	RE1228	069
Bst MCI	Mcrl	CGRY↓CG	RE1230	070
Bst NSI	Nsp l	RCATG↓Y	RE1232	070
Bst PAI	Psh Al	GACNN↓NNGTC	RE1234	071
Bst SNI	Sna BI	TAC [↓] GTA	RE1236	071
Bst V2I	Bbv II	GAAGAC (2/6)	RE1238	072
Bst XI	Bst XI	CANNNNN↓NTGG	RE1240	072
Bst X2I	XhoII	R [↓] GATCY	RE1242	073

List of Vivantis Restriction Endonucleases (cont'd)

Bit MI	Vivantis Restriction Endonuclease	Prototype	Recognition Sequence / Cleavage Site 5→3	Product No	Page
Dini	Btu MI	Nru I	TCG↓CGA	RV1246	074
Drall	Cci NI	Not I	gc [†] ggccgc	RE1248	074
Drall	Din I	Nar l *	ggc↓gcc	RV1252	075
Dise Dird GACNNINN NOTE RE1372 077	Dra l	Aha III	TTT [↓] AAA	RE1254	076
Saci	Dra III	Dra III	CACNNN [↓] GTG	RE1256	076
EcoRI	Dse DI	Drd l	GACNNNN↓NNGTC	RE1372	077
EcoRV	EcolCRI	Sac I*	GAG [↓] CTC	RE1258	078
FauNDI Ndel	EcoRI	Eco RI	G [↓] AATTC	RE1260	078
Foil Accl GT	EcoRV	EcoRV	GAT↓ATC	RE1262	079
Fok Fok GGATG (9/13) RE1270 081	Fau NDI	Nde l	CA [↓] TATG	RE1266	080
FriOI HgiJII GRGCY¹C RE1272 082 HindII HindII GTY¹RAC RE1274 083 HindIII HindIII A¹AGCTT RV1276 083 HinII HindIII A¹AGCTT RV1276 083 HinII HinII G¹ANTC RE1278 084 HpaII HpaII GT¹AC RE1280 084 HpaII HpaII C¹CGG RE1280 084 HpaII HpaII C¹CGG RE1282 085 HspAI Hhal¹ G¹CGG RE1284 085 KpnI KpnI GTCGC RE1284 085 KpnI KpnI GTCGC RE1286 086 Ksp22I BcII T¹GATCA RE1288 086 MboII MboII GAAGA (8/7) RE1290 087 MhII SduI GAGCGT RE1292 088 MmII MnII A¹CGCGT RE1292 088 MroNI<	Fb/ l	Accl	GT↓MKAC	RE1268	081
HindII HindII GTY IRAC RE1274 083 HindIII HindIII AIAGCTT RV1276 083 HindIII HindIII AIAGCTT RV1276 083 HindI HindIII AIAGCTT RV1276 083 HindI HindIII AIAGCTT RV1276 083 HindI HindIII GIANTC RE1278 084 HpaI GTT IAAC RE1280 084 HpaI HpaII CICGG RE1282 085 HspAI Hhal* GICGC RE1284 085 KpnI KpnI GGTACIC RV1286 086 Ksp22I BcII TIGATCA RE1288 086 MboIII MboII GAAGA (8/7) RE1290 087 MinII SduI GDGCHIC RE1292 088 MiuI MiuI AICGCGT RE1294 088 MiuI MiuI AICGCGT RE1294 088 MinII MinI CCTC (7/6) RE1296 089 MroNI Nael* GICCGGC RE1298 089 MroXI XmnI GAANNINNTTC RE1300 090 Msp1 HpaII CICGG RE1302 090 Msp20I BaII TGGICCA RE1302 090 Msp20I BaII TGGICCA RE1304 091 MspA1I NspBII CMGICCA RE1306 091 PceI StuI AGGICCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCTIC RE1312 094 PspCI PmaCI CACIGTG RE1314 095	Fok l	Fok l	GGATG (9/13)	RE1270	081
HindIII HindIII A AGCTT RV1276 083 HinfI HinfI G¹ANTC RE1278 084 HpaI HpaI G¹T¹AAC RE1280 084 HpaII HpaII C¹CGG RE1282 085 HspAI Hhal¹ G¹CGC RE1284 085 KpnI KpnI GGTAC¹C RV1286 086 Ksp22I BcII T¹GATCA RE1288 086 MboII MboII GAAGA (8/7) RE1290 087 MhII SduI GDGCH¹C RE1292 088 Miul Miul A¹CGCGT RE1292 088 Miul Miul A¹CGCGT RE1294 088 MroNI Miul A¹CGGGT RE1294 088 MroNI Nael¹ G¹CCGGC RE1296 089 MroXI XmnI GAANN¹NNTTC RE1300 090 Msp1 HpaII C¹CGG RE1302 090 Msp20	Fri OI	Hgi JII	GRGCY↓C	RE1272	082
Hinfl Hinfl GIANTC RE1278 084 Hpall Hpal GTT↓AAC RE1280 084 Hpall Hpall C↓GGG RE1282 085 HspAl Hhal¹ G¹CGC RE1284 085 Kpnl Kpnl GGTAC↓C RV1286 086 Ksp221 Bcil T↓GATCA RE1288 086 MbcIl Mboll GAAGA (8/7) RE1290 087 Mhil Sdul GDGCH↓C RE1292 088 Miul Miul A↓CGCGT RE1294 088 Miul Miul A¹CGCGT RE1294 088 Minl Moll CCTC (7/6) RE1296 089 MroNI Nael¹ G¹CCGGC RE1298 089 MroXI Xmnl GAANN↓NNTTC RE1300 090 Msp1 Hpall C↓CGG RE1302 090 Msp201 Bail TGG↓CCA RE1304 091 MspA11 NspBII CMG↓CKG RE1306 091 Pcel Stul AGG¹CCT RE1308 093 Pctl Bsml GAATGC (1/-1) RE1310 093 Psp124BI Sacl GAGCT↓C RE1312 094 PspC1 PmaCl CAC↓GTG RE1314 095	HindII	HindII	GTY [↓] RAC	RE1274	083
Hpal Hpal GTT↓AAC RE1280 084 Hpall Hpall C¹CGG RE1282 085 HspAl Hhal* G¹CGC RE1284 085 KpnI KpnI GGTAC¹C RV1286 086 Ksp22l BcI T¹GATCA RE1288 086 MbcII MbcII GAAGA (8/7) RE1290 087 MhII Sdul GDGCH¹C RE1292 088 Mull Mull A¹CGCGT RE1292 088 MnII Mull A¹CGCGT RE1294 088 MnII MnII CCTC (7/6) RE1296 089 MroNI Nael* G¹CCGGC RE1298 089 MroXI XmnI GAANN¹NNTTC RE1300 090 MspI HpaII C¹CGG RE1302 090 Msp20I BaI TGG¹CA RE1304 091 MspA1I NspBII CMG¹CCA RE1306 091 Pcel <th>HindIII</th> <th>HindIII</th> <th>A↓AGCTT</th> <th>RV1276</th> <th>083</th>	HindIII	HindIII	A↓AGCTT	RV1276	083
Hpall Hpall C¹CGG RE1282 085 HspAl Hhal* G¹CGC RE1284 085 KpnI GGTAC¹C RV1286 086 KpnI GGTAC¹C RV1286 086 KpnI GGTAC¹C RV1286 086 MboII T¹GATCA RE1288 086 MboII MboII GAAGA (8/7) RE1290 087 MhiI SduI GDGCH¹C RE1292 088 Miul Miul A¹CGCGT RE1292 088 MnII Miul A¹CGCGT RE1294 088 MnoII MnII CCTC (7/6) RE1296 089 MroNI Nael* G¹CCGGC RE1298 089 MroXI XmnI GAANN¹NNTTC RE1300 090 MspI HpaII C¹CGG RE1302 090 Msp20I BaII TGG¹CCA RE1304 091 MspA1I NspBII CMG¹CKG RE1306 <th< th=""><th>Hinfl</th><th>Hinfl</th><th>G↓ANTC</th><th>RE1278</th><th>084</th></th<>	Hinfl	Hinf l	G↓ANTC	RE1278	084
HspAI Hhal* G¹CGC RE1284 085 KpnI GGTAC¹C RV1286 086 Ksp22I BcI T¹GATCA RE1288 086 MboII MboII GAAGA (8/7) RE1290 087 MhII SduI GDGCH¹C RE1290 087 MhII Miul A¹CGCGT RE1292 088 Mail Miul A¹CGCGT RE1294 088 Mail Miul A¹CGCGT RE1294 088 Mail Mail CCTC (7/6) RE1296 089 MroNI Nael* G¹CGGC RE1298 089 MroXI Xmal GAANN¹NNTTC RE1300 090 MspI HpaII C¹CGG RE1302 090 Msp20I BaiI TGG¹CCA RE1304 091 MspA1I NspBII CMG¹CCA RE1306 091 PceI StuI AGG¹CCT RE1300 093 PcI BsmI <th>Hpal</th> <th>Hpal</th> <th>GTT↓AAC</th> <th>RE1280</th> <th>084</th>	Hpa l	Hpa l	GTT↓AAC	RE1280	084
KpnI KpnI GGTAC	Hpa ll	Hpa ll	c↓cgg	RE1282	085
Ksp22I BcII T↓ GATCA RE1288 086 MboII MboII GAAGA (8/7) RE1290 087 MhII SduI GDGCH↓C RE1292 088 MIUI MIUI A↓ CGCGT RE1294 088 MnII MnII CCTC (7/6) RE1296 089 MroNI Nael⁺ G↓ CCGGC RE1298 089 MroXI XmnI GAANN↓NNTTC RE1300 090 MspI HpaII C↓ CGG RE1302 090 Msp20I BaII TGG↓ CCA RE1304 091 MspA1I NspBII CMG↓ CKG RE1306 091 PceI StuI AGG↓ CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT↓ C RE1312 094 PspCI PmaCI CAC GGT RE1314 095	Hsp AI	Hha l*	g↓cgc	RE1284	085
Mboll Mboll GAAGA (8/7) RE1290 087 Mh/I SduI GDGCH↓C RE1292 088 Miul Miul A↓CGCGT RE1294 088 Mn/I Mn/I CCTC (7/6) RE1296 089 MroNI Nael* G↓CCGGC RE1298 089 MroXI XmnI GAANN↓NNTTC RE1300 090 MspI HpaII C↓CGG RE1302 090 Msp20I Bail TGG↓CCA RE1304 091 MspA1I NspBII CMG↓CKG RE1306 091 PceI StuI AGG↓CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT↓C RE1312 094 PspCI PmaCI CAC↓GTG RE1314 095	Kpn l	Kpn l	GGTAC↓C	RV1286	086
Mh/I SduI GDGCH↓C RE1292 088 MIuI MiuI A↓CGCGT RE1294 088 Mn/I Mn/I CCTC (7/6) RE1296 089 MroNI NaeI* GJCCGGC RE1298 089 MroXI XmnI GAANN↓NNTTC RE1300 090 MspI HpaII CJCGG RE1302 090 Msp20I BaII TGGJCCA RE1304 091 MspA1I NspBII CMGJCKG RE1306 091 PceI StuI AGGJCTC RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCTJC RE1312 094 PspCI PmaCI CACJGTG RE1314 095	Ksp 22I	Bc/I	T [↓] GATCA	RE1288	086
Miul Miul A↓CGCGT RE1294 088 MnII MnII CCTC (7/6) RE1296 089 MroNI Nael* G↓CCGGC RE1298 089 MroXI XmnI GAANN↓NNTTC RE1300 090 MspI HpaII C↓CGG RE1302 090 Msp20I Ball TGG↓CCA RE1304 091 MspA1I NspBII CMG↓CKG RE1306 091 PceI Stul AGG↓CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT↓C RE1312 094 PspCI PmaCI CAC↓GTG RE1314 095	Mbo ll	Mbo ll	GAAGA (8/7)	RE1290	087
Mn/I Mn/I CCTC (7/6) RE1296 089 MroNI Nael* G¹ CCGGC RE1298 089 MroXI XmnI GAANN¹ NNTTC RE1300 090 MspI HpaII C¹ CGG RE1302 090 Msp20I Ba/I TGG¹ CCA RE1304 091 MspA1I NspBII CMG¹ CKG RE1306 091 PceI Stul AGG¹ CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT¹ C RE1312 094 PspCI PmaCI CAC¹ GTG RE1314 095	Mh/ l	Sdu l	GDGCH [↓] C	RE1292	088
MroNI Nael* $G^{\downarrow}CCGGC$ RE1298 089 MroXI XmnI GAANN $^{\downarrow}$ NNTTC RE1300 090 MspI HpaII $C^{\downarrow}CGG$ RE1302 090 Msp20I Ball TGG $^{\downarrow}CCA$ RE1304 091 MspA1I NspBII CMG $^{\downarrow}CKG$ RE1306 091 PceI Stul AGG $^{\downarrow}CCT$ RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT $^{\downarrow}C$ RE1312 094 PspCI PmaCI CAC $^{\downarrow}$ GTG RE1314 095	Mlu l	Mlu l	A↓CGCGT	RE1294	088
MroXI XmnI GAANN↓NNTTC RE1300 090 MspI HpaII C↓CGG RE1302 090 Msp20I Ball TGG↓CCA RE1304 091 MspA1I NspBII CMG↓CKG RE1306 091 PceI StuI AGG↓CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT↓C RE1312 094 PspCI PmaCI CAC↓GTG RE1314 095	Mn/ l	MnII	CCTC (7/6)	RE1296	089
MspI HpaII C^{\downarrow} CGG RE1302 090 Msp20I Ball TGG $^{\downarrow}$ CCA RE1304 091 MspA1I NspBII CMG $^{\downarrow}$ CKG RE1306 091 PceI StuI AGG $^{\downarrow}$ CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT $^{\downarrow}$ C RE1312 094 PspCI PmaCI CAC $^{\downarrow}$ GTG RE1314 095	Mro NI	Nae l*	g↓ccggc	RE1298	089
$Msp20I$ $BalI$ $TGG^{\downarrow}CCA$ $RE1304$ 091 $MspA1I$ $NspBII$ $CMG^{\downarrow}CKG$ $RE1306$ 091 $PceI$ $StuI$ $AGG^{\downarrow}CCT$ $RE1308$ 093 $PctI$ $BsmI$ $GAATGC (1/-1)$ $RE1310$ 093 $Psp124BI$ $SacI$ $GAGCT^{\downarrow}C$ $RE1312$ 094 $PspCI$ $PmaCI$ $CAC^{\downarrow}GTG$ $RE1314$ 095	MroXI	Xmn l	GAANN↓NNTTC	RE1300	090
MspA1I NspBII CMG↓CKG RE1306 091 PceI StuI AGG↓CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT↓C RE1312 094 PspCI PmaCI CAC↓GTG RE1314 095	Msp l	Hpa ll	c↓cgg	RE1302	090
PceI StuI AGG ↓ CCT RE1308 093 PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT ↓ C RE1312 094 PspCI PmaCI CAC ↓ GTG RE1314 095	Msp 20I	Ba/I	TGG↓CCA	RE1304	091
PctI BsmI GAATGC (1/-1) RE1310 093 Psp124BI SacI GAGCT ↓ C RE1312 094 PspCI PmaCI CAC ↓ GTG RE1314 095	Msp A1I	Nsp BII	CMG [↓] CKG	RE1306	091
Psp124BI SacI GAGCT ↓ C RE1312 094 PspCI PmaCI CAC ↓ GTG RE1314 095	Pce l	Stu l	AGG↓CCT	RE1308	093
PspCI PmaCI CAC GTG RE1314 095	Pct l	Bsm l	GAATGC (1/-1)	RE1310	093
	Psp124BI	Sacl	GAGCT↓C	RE1312	094
PspEI BstEII G GTNACC RE1316 095	PspCI	PmaCl	CAC↓GTG	RE1314	095
	PspEI	Bst EII	G [↓] GTNACC	RE1316	095

List of Vivantis Restriction Endonucleases (cont'd)

Vivantis Restriction Endonuclease	Prototype	Recognition Sequence / Cleavage Site 5'→3'	Product No	Page
PspOMI	Apa l*	g↓ggccc	RE1318	096
Pst l	Pst l	CTGCA [↓] G	RE1320	096
Pvu ll	PvuII	CAG [↓] CTG	RE1322	097
Rsa l	Rsal	GT↓AC	RE1324	097
Rsr 2I	Rsr II	cd† dwccd	RE1374	098
Sall	Sal l	G [↓] TCGAC	RV1326	099
Sbf l	Sse 8387 I	CCTGCA [↓] GG	RE1328	100
Sfa NI	Sfa NI	GCATC (5/9)	RE1376	101
Sfi l	Sfi I	GGCCNNNN [↓] NGGCC	RE1330	101
Sfr 274I	Xho l	C↓TCGAG	RE1332	102
Sfr 303I	SacII	ccec†ee	RE1334	102
Smal	Smal	ccc∮ggg	RE1336	103
Smi l	Swal	ATTT [↓] AAAT	RE1338	103
Smi MI	Ms/I	CAYNN [↓] NNRTG	RE1378	104
Sph l	Sph l	GCATG [↓] C	RV1340	104
Sse 9I	Tsp EI	[↓] AATT	RE1342	105
Ssp l	Ssp l	AAT [↓] ATT	RE1344	105
Taq l	Taq l	T↓CGA	RE1346	106
Tru 9I	Mse l	T ↓ TAA	RE1350	106
Tth 1111	Tth 111	GACN [↓] NNGTC	RE1356	107
Vha 464I	Af/ II	C [‡] TTAAG	RE1358	107
Vne l	Apa LI	G↓TGCAC	RE1360	108
Vsp l	Vspl	AT [‡] TAAT	RE1362	108
Xbal	Xbal	T [↓] CTAGA	RV1364	109
Xma l	Sma l*	c↓ccggg	RV1366	110
Zral	Aat II*	GAC↓GTC	RE1368	111
Zsp 2l	Ava III	ATGCA↓T	RE1370	111

Base Nomenclature

B: C/G/T M: A/C Y : C/T D: A/G/T H: A/C/T R: A/G V : A/C/G S: C/G W: A/T K: G/T

N: A/C/G/T



^{*} Indicates neoschizomer

List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Aat I	AGG↓CCT	Pce I	RE1308	093
Aat II	GACGT↓C	Aat II Zra I* (GAC [‡] GTC)	RE1100 RE1368	033 111
Acc16I	TGC↓GCA	Acc 16I	RE1102	033
Acc 651	G [↓] GTACC	Acc 651 KpnI* (GGTAC [‡] C)	RE1104 RV1286	034 086
Acc B1I	G [‡] GYRCC	Acc B1I	RE1106	034
Acc B7I	CCANNNN [↓] NTGG	Acc B7I	RE1108	035
AccBSI	CCGCTC (-3/-3)	AccBSI	RE1110	035
Acc I	GT↓MKAC	Fb1 1	RE1268	081
Acc II	cg↓cg	Bst FNI	RE1220	067
AccIII	T↓CCGGA	Bsp 13I	RE1186	058
Acl	AA↓CGTT	Ac/I	RE1112	036
Acs I	R [↓] AATTY	Acs I	RE1114	036
Acvl	CAC↓GTG	PspCI	RE1314	095
Acy I	GR↓CGTC	Bss NI	RV1194	061
Ade I	CACNNN [↓] GTG	Dra III	RE1256	076
Afa l	GT↓AC	Rsa I	RE1324	097
Afi I	CCNNNNN [↓] NNGTC	Afi I	RV1116	037
Af/ II	C [↓] TTAAG	Vha 464I	RE1358	107
Age I	A↓CCGGT	Asi GI	RE1126	041
Ahd I	GACNNN↓NNGTC	Bme R I	RV1150	048
Ahl l	A↓CTAGT	Ahl I	RE1118	038
Ajn I	[↓] ccwgg	Bst2UI* (CC [↓] WGG)	RE1202	063
Alul	AG↓CT	Alu I	RE1120	038
Alw 21I	gwgcw↓c	Bbv12I	RE1140	045
Alw 26I	GTCTC (1/5)	Bst MAI	RE1226	069
Alw 441	G↓TGCAC	Vnel	RE1360	108
Ama 87I	C [↓] YCGRG	Ama 871	RE1122	039
Aor 13HI	T↓CCGGA	Bsp13I	RE1186	058
Apa l	gggcc↓c	Apal PspОМ I* (G [↓] GGCCC)	RE1124 RE1318	040 096
Apa LI	G↓TGCAC	Vnel	RE1360	108
Аро І	R↓AATTY	Acsl	RE1114	036
Asel	AT [↓] TAAT	Vspl	RE1362	108
Asi G I	A↓CCGGT	Asi GI	RE1126	041
Asp I	GACN [↓] NNGTC	Tt h 1111	RE1356	107

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Asp 7001	GAANN [↓] NNTTC	MroXI	RE1300	090
Asp 718I	G [↓] GTACC	Acc 651 Kpn1* (GGTAC [↓] C)	RE1104 RE1286	034 086
Asp A2I	C [↓] CTAGG	Asp A2I	RE1128	041
Asp EI	GACNNN NNGTC	Bme RI	RV1150	048
Asp LEI	gcg↓c	Asp LEI Bst HHI Hsp AI* (G[↓]CGC)	RE1130 RE1224 RE1284	042 068 085
Asp S9I	G↓GNCC	Asp S9I	RE1132	042
Ass l	AGT↓ACT	Bmc Al	RV1146	047
Asu C21	cc↓sgg	Ври МІ	RV1162	051
Asu HPI	GGTGA (8/7)	Asu HPI	RE1134	043
Asu ll	TT↓CGAA	Bpu 14I	RE1160	051
Asu NHI	G [↓] CTAGC	Asu NHI Bmt I* (GCTAG[‡]C)	RE1136 RE1156	043 049
Ava I	C [↓] YCGRG	Ama 871	RE1122	039
Ava II	G↓GWCC	Bme18I	RE1148	047
Avi II	TGC↓GCA	Acc16I	RE1102	033
Avr II	C [↓] CTAGG	Asp A2I	RE1128	041
Axy I	CC↓TNAGG	Bse 21I	RE1170	054
Bal l	TGG↓CCA	Msp 20I	RE1304	091
Bam HI	G [↓] GATCC	Bam HI	RV1138	044
Ban I	G↓GYRCC	Acc B1I	RE1106	034
Ban ii	GRGCY↓C	Fri OI	RE1272	082
Ban III	AT↓CGAT	Bsh VI	RV1180	056
Bbe I	ggcgc↓c	DinI* (GGC ↓ GCC)	RE1252	075
Bbr PI	CAC↓GTG	Psp CI	RE1314	095
Bbs I	GAAGAC (2/6)	Bst V2I	RE1238	072
Bbu l	GCATG↓C	Sph I	RV1340	104
Bbv12I	gwgcw↓c	Bbv 12 I	RE1140	045
Bcl	T↓GATCA	Ksp 221	RE1288	086
Bcn I	cc↓sgg	Ври М І	RV1162	051
Bcul	A↓CTAGT	Ahl I	RE1118	038
Bfr l	C [↓] TTAAG	Vha 464 l	RE1358	107
Bfr BI	ATG [↓] CAT	Zsp2I* (ATGCA↓T)	RE1370	111
Bfu CI	[↓] GATC	Bss MI Bst MBI	RV1192 RE1228	060 069

List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases (cont'd)

	Recognition Sequence / Cleavage Site 5→3Endonuclea		Product No	Page
Bgl l	GCCNNNN↓NGGC	Bg/ I	RV1142	046
Bgl ll	A [↓] GATCT	Bgl ll	RE1144	046
Blf l	T↓CCGGA	Bsp13I	RV1186	058
Bln l	C↓CTAGG	Asp A2I	RE1128	041
Blp I	GC↓TNAGC	Bsp 1720 I	RE1188	059
Bmc Al	AGT↓ACT	Bmc Al	RV1146	047
Bme 1390 I	CC↓NGG	Bmr FI	RV1154	049
Bme 18I	G [‡] GWCC	Bme 18I	RE1148	047
Bme RI	GACNNN↓NNGTC	Bme RI	RV1150	048
Bmi I	GGN↓NCC	Bmi I	RV1152	048
Bmr FI	CC↓NGG	Bm FI	RV1154	049
Bmt I	GCTAG [‡] C	AsuNHI* (G [‡] CTAGC) Bmt1	RE1136 RE1156	043 049
Bmy I	GDGCH↓C	Mh/ I	RE1292	088
Box I	GACNN [↓] NNGTC	Bst PAI	RE1234	071
Bpi l	GAAGAC (2/6)	Bst V2I	RE1238	072
Bpt I	cc↓wgg	Bst 2UI	RE1202	063
Bpu 10 I	CCTNAGC (-5/-2)	Bpu 10I	RE1158	050
Bpu 1102I	GC↓TNAGC	Bsp 1720 l	RE1188	059
Bpu 14I	TT↓CGAA	Bpu 14I	RE1160	051
Ври АІ	GAAGAC (2/6)	Bst V2I	RE1238	072
Ври МІ	cc↓sgg	Bpu M I	RV1162	051
Bsa l	GGTCTC (1/5)	Bso 31 I	RE1184	058
Bsa 29 I	AT↓CGAT	Bsh VI	RV1180	056
Bsa Al	YAC↓GTR	Bst BAI	RE1210	065
Bsa BI	GATNN [↓] NNATC	Bse 8I	RE1174	055
Bsa HI	GR [↓] CGYC	Bss NI	RV1194	061
Bsa M I	GAATGC (1/-1)	Pct I	RE1310	093
Bsc 41	CCNNNNN [↓] NNGG	Afi l	RV1116	037
Bse 1I	ACTGG (1/-1)	Bse 1I	RE1166	053
Bse 118I	R↓CCGGY	Bse118I	RE1168	053
Bse 21I	CC↓TNAGG	Bse 211	RE1170	054
Bse 3DI	GCAATG (2/0)	Bse 3DI	RE1172	054
Bse 81	GATNN [↓] NNATC	Bse 8I	RE1174	055
BseAl	T↓CCGGA	Bsp13I	RE1186	058
Bse BI	cc↓wgg	Bst 2UI	RE1202	063

	Recognition Sequence / Cleavage Site 5→3Endonuclease	Vivantis Restriction	Product No	Page
BseC I	AT [↓] CGAT	Bsh VI	RV1180	056
Bse GI	GGATG (2/0)	Bst F5I Fok I* (GGATG (9/13))	RE1218 RE1270	067 081
Bse JI	GATNN↓NNATC	Bse 81	RE1174	055
Bse LI	CCNNNNN↓NNGG	Afi l	RV1116	037
Bse MI	GCAATG (2/0)	Bse3DI	RE1172	054
Bse NI	ACTGG (1/-1)	Bse 11	RE1166	053
Bse PI	g↓cgcgc	Bse PI	RE1176	055
Bse X3I	c∮ggccg	Bse X3I	RE1178	056
Bsh 1236 I	cg↓cg	Bst FNI	RE1220	067
Bsh 1285 I	CGRY↓CG	Bst MCI	RE1230	070
Bsh FI	gg↓cc	Bsn l	RV1182	057
Bsh NI	G [↓] GYRCC	Acc B1I	RE1106	034
Bsh TI	A [↓] CCGGT	Asi GI	RE1126	041
Bsh VI	AT↓CGAT	Bsh VI	RV1180	056
Bsi EI	CGRY↓CG	Bst MCI	RE1230	070
Bsi HKAI	gwgcw↓c	Bbv 12 I	RE1140	045
Bsi HKCI	C↓YCGRY	Ama 87 I	RE1122	039
Bsi SI	c↓cgg	Hpa ll Msp l	RE1282 RE1302	085 090
Bsi YI	CCNNNNN [↓] NNGG	Afi l	RV1116	037
Bs/I	CCNNNNN [↓] NNGG	Afi l	RV1116	037
Bsm Al	GTCTC (1/5)	Bst MAI	RE1226	069
Bsm l	GAATGC (1/-1)	Pct I	RE1310	093
Bsn l	gg↓cc	Bsn l	RV1182	057
Bso 31 I	GGTCTC (1/5)	Bso 31 I	RE1184	058
Bso BI	C↓YCGRG	Ama 871	RE1122	039
Bsp 119I	TT↓CGAA	Bpu 14I	RE1160	051
Bsp 120 I	g [†] ggccc	Apal* (GGGCC [↓] C) Psp OMI	RE1124 RE1318	040 096
Bsp 1286 I	GDGCH [↓] C	Mh/ I	RE1292	088
Bsp13I	T↓CCGGA	Bsp13I	RE1186	058
Bsp 1407I	T [‡] GTACA	Bst AUI	RE1208	064
Bsp 143 I	[↓] GATC	Bss MI	RV1192	060
		Bst MBI	RE1228	069
Bsp143II	RGCGC↓Y	Bst H2I	RE1222	068
Bsp 1720 I	GC↓TNAGC	Bsp 1720 I	RE1188	059



List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases (cont'd)

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Bsp19I	C↓CATGG	Bsp19I	RE1190	059
Bsp 681	TCG↓CGA	Btu MI	RV1246	074
Bsp ANI	GG↓CC	Bs nl	RV1182	057
Bsp DI	AT↓CGAT	Bsh VI	RV1180	056
Bsp E I	T↓CCGGA	Bsp13I	RE1186	058
Bsp LI	GGN↓NCC	Bmi l	RV1152	048
Bsp MAI	CTGCA [↓] G	Pst I	RE1320	096
Bsp T104I	TT↓CGAA	Bpu 14l	RE1160	051
Bsp T107I	G↓GYRCC	Acc B1I	RE1106	034
Bsp TI	C↓TTAAG	Vha 464I	RE1358	107
Bsp TNI	GGTCTC (1/5)	Bso 31 I	RE1184	058
Bsp XI	AT [↓] CGAT	Bsh VI	RV1180	056
Bsr l	ACTGG (1/-1)	Bse 1I	RE1166	053
Bsr Bl	CCGCTC (-3/-3)	Acc BSI	RE1110	035
Bsr DI	GCAATG (2/0)	Bse 3DI	RE1172	054
Bsr Fl	R↓CCGGY	Bse 118I	RE1168	053
Bsr GI	T [↓] GTACA	Bst AUI	RE1208	064
Bsr SI	ACTGG (1/-1)	Bse 11	RE1166	053
Bss Al	R↓CCGGY	Bse 118I	RE1168	053
Bss HI	C↓TCGAG	Sfr 274 I	RE1332	102
Bss HII	g↓cgcgc	Bse PI	RE1176	055
Bss KI	[↓] CCNGG	BmrFI* (CC [↓] NGG)	RV1154	049
Bss MI	[↓] GATC	Bss MI Bst MBI	RV1192 RE1228	060 069
Bss NI	GR↓CGYC	Bss N I	RV1194	061
Bss NAI	GTA [↓] TAC	Bss NAI	RE1196	061
Bss T1I	c↓cwwgg	Bss T1I	RE1198	062
Bst 1107 I	GTA [↓] TAC	Bss NAI	RE1196	061
Bst 2UI	cc↓wgg	Bst 2UI	RE1202	063
Bst 4Cl	ACN↓GT	Bst 4CI	RE1204	063
Bst 6 I	CTCTTC (1/4)	Bst 61	RE1206	064
Bst 98I	C↓TTAAG	Vha 464 I	RE1358	107
Bst ACI	GR↓CGYC	Bss NI	RV1194	061
Bst AUI	T [↓] GTACA	Bst AUI	RE1208	064
Bst BAI	YAC↓GTR	Bst BAI	RE1210	065
Bst Bl	TT↓CGAA	Ври 14I	RE1160	051

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Bst DEI	C↓TNAG	Bst DEI	RE1212	065
Bst DSI	C [↓] CRYGG	Bst DSI	RE1214	066
Bst EII	G [↓] GTNACC	Psp EI	RE1316	095
Bst ENI	CCTNN [↓] NNNAGG	Bst ENI	RE1216	066
Bst F51	GGATG (2/0)	Bst F51 Fok I* (GGATG(9/13))	RE1218 RE1270	067 081
Bst FNI	cg↓cg	Bst FNI	RE1220	067
Bst H2I	RGCGC↓Y	Bst H2I	RE1222	068
Bst HHI	GCG↓C	AspLEI BstHHI Hsp AI* (G[↓]CGC)	RE1130 RE1224 RE1284	042 068 085
Bst KTI	GAT [↓] C	BssMI* (↓GATC) BstMBI* (↓GATC)	RV1192 RE1228	060 069
Bst MAI	GTCTC (1/5)	Bst MAI	RE1226	069
Bst MBI	↓GATC	Bss MI Bst MBI	RV1192 RE1228	060 069
Bst MCI	CGRY↓CG	Bst MCI	RE1230	070
Bst NI	cc↓wgg	Bst 2UI	RE1202	063
Bst NSI	RCATG [↓] Y	Bst NSI	RE1232	070
Bst OI	cc↓wgg	Bst 2UI	RE1202	063
Bst PAI	GACNN↓NNGTC	Bst PAI	RE1234	071
Bst PI	G [↓] GTNACC	Psp EI	RE1316	095
Bst SCI	[↓] CCNGG	BmrFI* (CC ↓ NGG)	RV1154	049
Bst SNI	TAC [↓] GTA	Bst SNI	RE1236	071
Bst UI	cg↓cg	Bst FNI	RE1220	067
Bst V2I	GAAGAC (2/6)	Bst V2I	RE1238	072
Bst XI	CCANNNNN [↓] NTGG	Bst XI	RE1240	072
Bst X2I	R↓GATCY	Bst X2I	RE1242	073
Bst YI	R↓GATCY	Bst X21	RE1242	073
Bst ZI	c [†] ggccg	Bse X3I	RE1178	056
Bst Z17I	GTA [↓] TAC	Bss NAI	RE1196	061
Bsu 15I	AT↓CGAT	Bsh VI	RV1180	056
Bsu 36I	CC↓TNAGG	Bse 21 I	RE1170	054
Bsu RI	GG↓CC	Bsn l	RV1182	057
Bsu TUI	AT ↓ CGAT	Bsh VI	RV1180	056
Btg l	C↓CRYGG	Bst DSI	RE1214	066
Btu M I	TCG↓CGA	Btu MI	RV1246	074
Cci NI	gc↓ggccgc	Cci NI	RE1248	074

List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases (cont'd)

Cell	Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
BEI-HHI RE1224 088	Ce/II	GC [↓] TNAGC	Bsp1720I	RE1188	059
Christ G¹GNCC AsrS9I RE1132 042 Chr42I CCGC¹GG Sh 303I RE1334 102 Chr9I C¹CGGG Sh 303I RE1334 102 Chr9I C¹CGGG Sh 1 (CCC¹GGG) RE1334 102 Chr1 AT¹CGAT Bh VI RV1180 056 Cap 45I TT¹CGAA Bh VI RV1180 056 Cap AI A¹CCGGT AsrGI RE1324 097 Drall GGC¹GCC Dint RV1252 075 Drall GGC¹GCC Dint RV1252 075 Drall TTT¹AAA Drall RE1225 076 Drall GACNNN¹ANGTC Drall RE1256 076 Drall GACNNN¹ANGTC Brac BI RV1150 048	Cfol	gcg↓c	Bst HHI	RE1224	068
Cfr42I CCGC ¹ GG Sfr303I RE1334 102 Cfr9I C ¹ CCGGG Smal* (CCC ¹ GGG) RE1338 103 CbI AT ¹ CGAT BshVI RV1366 110 CbI AT ¹ CGAT BshVI RV1180 056 CbI TT ¹ CGAA BsuVI RV1180 056 CbI GGI RE1160 051 051 Cbi GGI RE1160 051 051 Cbi GGI RE1126 041 051 Cbi Did RE1324 097 097 Cbi Did RE1324 097 097 Cbi Did RE1324 097 097 095 097 096 097 097 096 097 097 096 097 097 096 097 099 099 099 099 099 099 099 099 099 099 099 099 099 099 099 <	Cfr 10I	R↓CCGGY	Bse 118I	RE1168	053
Cir 9 C ¹ CCGGG Smal* (CCC ¹ GGG) RE1336 103 205 110 103 105	Cfr 13I	G [↓] GNCC	Asp S9I	RE1132	042
Xmal RV1366 110	Cfr 42I	ccgc↓gg	Sfr 303I	RE1334	102
Csp 451 TT¹ CGAA Bou14I RE1160 051 Csp 6I G¹TAC Rsal* (GT¹AC) RE1324 097 Csp AI A¹ CCGGT As/GI RE1324 097 Csp AI A¹ CCGGT As/GI RE1324 097 Doel C¹ TNAG Bst DEI RE1126 041 Doel GGC¹GC Dral RV1522 075 Dorll GGC¹GC Dral RV1522 075 Dorll GGC¹GC Bss M¹ RV1192 060 Best MBI RV1192 060 RE1228 075 Doral TTT¹ AAA Dral RE1254 076 Drall CACNNN¹ ANGTC Drall RE1256 076 Drall GACNNN¹ NNGTC Drall RV1150 048 Dse DI GACNNN¹ NNGTC Bre RI RV1150 048 Dse DI GACNNN¹ NNGTC Bre RI RV1150 048 Eam¹ 104I CTCTTC (1/4) Bst 6I RE1206	Cfr 91	c↓ccggg			
CSp 6I G↓ TAC Rsal* (GT↓AC) RE1324 097 CSp AI A↓ CCGGT As/GI RE1126 041 DdeI C↓ TNAG Bst DEI RE1212 065 DinI GGC GCC DinI RV1252 075 DpnII ↓ GATC Bss M I RV1925 075 DpnIII ↓ GATC Bss M I RV1925 069 Droll TTT ↓ AAA Drall RE1284 069 Droll CACNNN ↓ GTG DrallI RE1256 076 Droll GACNNN ↓ NNGTC Des DI RE1372 077 DrI GACNNN ↓ NNGTC Bme RI RV1150 048 Dse DI GACNNN ↓ NNGTC Des D I RE1372 077 Eagl C ↓ GGCCG Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst 6I RE1206 064 Eam1105I GACNNN ↓ NNGTC Bme RI RV1150 048 Eal CTCTTC (1/4) Bst 6I RE1206	Clal	AT [↓] CGAT	Bsh VI	RV1180	056
CSp AI	Csp 451	TT [↓] CGAA	Bpu 14I	RE1160	051
Ddel C¹TNAG BstDEI RE1212 065 DinI GGC¹GCC DinI RV1252 075 DpnIII ¹GATC Bss M I RV1192 060 RE1228 069 060 RE1228 069 Dral TTT¹AAA Dral RE1254 076 DralIII CACNNN¹GTG DralIII RE1256 076 Dral GACNNN¹NNGTC Des DI RE1372 077 Dril GACNNN¹NNGTC BmeRI RV1150 048 Dse DI GACNNN¹NGTC BmeRI RV1150 048 Dse DI GACNNN¹NGTC Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst6I RE1206 064 Eam1105I GACNNN¹NGTC BmeRI RV1150 048 Earl CTCTTC (1/4) Bst6I RE1206 064 EciHKI GACNNN¹NGTC BmeRI RV1150 048 EciHKI GACNNN¹NGTC BmeRI RV1150 048 Eci136II GAG*CT BseX3I RE1258 078 PSp124B I*(GAGCT¹C) RE1312 094 Eci3II RE178 056	Csp 6I	G [↓] TAC	Rsal* (GT ↓ AC)	RE1324	097
Diric GGC GCC Diric RV1252 075	Csp AI	A↓CCGGT	Asi GI	RE1126	041
Doril	Dde l	C↓TNAG	Bst DEI	RE1212	065
Bst MBI RE1228 069	Din l	ggc↓gcc	Din l	RV1252	075
Draili CACNNN¹ GTG Draili RE1256 076 Drol GACNNNN¹ NNGTC Des DI RE1372 077 Dril GACNNN¹ NNGTC Bme RI RV1150 048 Dse DI GACNNN¹ NNGTC Des D I RE1372 077 Eagl C¹ GGCCG Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst 6I RE1206 064 Eam1105I GACNNN¹ NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bst 6I RE1206 064 EclHKI GACNNN¹ NNGTC Bme RI RV1150 048 EclHKI GACNNN¹ NNGTC Bme RI RV1150 048 Ecl136II GAG¹ CTC EcolCRI RE1258 078 R81312 094 EclXI C¹ GGCCG Bse X3I RE1178 056 056 Eco105I TAC¹ GTA Bsi SNI RE1236 071 071 071 072 072 072 072 072 072 <td>Dpnll</td> <td>[↓] GATC</td> <td></td> <td></td> <td></td>	Dpn ll	[↓] GATC			
Drol GACNNNN↓NNGTC Des DI RE1372 077 Dril GACNNN↓NNGTC Bme RI RV1150 048 Dse DI GACNNNN↓NNGTC Des DI RE1372 077 Eagl C↓GGCCG Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst 6I RE1206 064 Eam1105I GACNNN↓NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bst 6I RE1206 064 EclHKI GACNNN↓NNGTC Bme RI RV1150 048 EclHKI GAGNNN↓NNGTC Bme RI RV1150 048 Ecl136II GAG↓CTC EcolCRI RE1258 078 Psp124B I*(GAGCT↓C) RE1312 094 EclXI C↓GGCCG BseX3I RE1178 056 Eco130I TAC↓GTA Bs/SNI RE1236 071 Eco130I C↓CWWGG BsST1I RE1198 062 Eco147I AGG↓CT PceI RE1308 09	Dral	TTT [↓] AAA	Dral	RE1254	076
Dril GACNNN¹ NNGTC Bme RI RV1150 048 Dse DI GACNNNN¹ NNGTC Des D I RE1372 077 Eagl C¹ GGCCG Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst 6I RE1206 064 Eam1105I GACNNN¹ NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bst 6I RE1206 064 Ec/HKI GACNNN¹ NNGTC Bme RI RV1150 048 Ec/136II GAG¹ CTC Ecol CRI RE1258 078 Psp124B I¹ (GAGCT¹ C) RE1312 094 Ec/XI C¹ GGCCG Bse X3I RE1178 056 Eco105I TAC¹ GTA Bst SNI RE1236 071 Eco105I TAC¹ GTA Bst SNI RE1236 071 Eco130I C¹ CWWGG Bss T1I RE1198 062 Eco147I AGG¹ CCT PceI RE1308 093 Eco24I GRGCY¹ C FrIOI RE1262	Dra lli	CACNNN [↓] GTG	Dra lli	RE1256	076
Dse DI GACNNNN I NNGTC Des DI RE1372 077 Eagl C I GGCCG Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst 6I RE1206 064 Eam1105I GACNNN NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bst 6I RE1206 064 Ecilki GACNNN NNGTC Bme RI RV1150 048 Ecilki GACNNN NNGTC Bme RI RV1150 048 Ecilia GACNNN NNGTC Bme RI RV1150 048 Ecilia GACNNN I NNGTC Bme RI RV1150 048 Ecilia GAGCCT Be SS X3I RE1178 056 Ecilia C I GGCCG Bss X3I RE1198 062 Eco130I C I C	Drd l	GACNNNN [↓] NNGTC	Des DI	RE1372	077
Eagl C↓GGCCG Bse X3I RE1178 056 Eam1104I CTCTTC (1/4) Bst 6I RE1206 064 Eam1105I GACNNN↓NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bst 6I RE1206 064 Ec/HKI GACNNN↓NNGTC Bme RI RV1150 048 Ec/136II GAG CTC EcolCRI Psp124B I*(GAGCT↓C) RE1258 078 Ec/136II GAG GCCG Bse X3I RE1178 056 EcolXI C↓GGCCG Bse X3I RE1178 056 Ecol05I TAC J GTA Bst SNI RE1236 071 Eco130I C↓CWWGG Bss T1I RE1198 062 Eco147I AGG CCT PceI RE1308 093 Eco24I GRGCY C FriOI RE1272 082 Eco31I GGTCTC (1/5) Bsc 31I RE1184 058 Eco47I GJ GWCC Bme18I RE1148 047	Dri l	GACNNN [↓] NNGTC	Bme RI	RV1150	048
Eam1104I CTCTTC (1/4) Bsi 6I RE1206 064 Eam1105I GACNNN NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bsi 6I RE1206 064 EclHKI GACNNN NNGTC Bme RI RV1150 048 Ecl136II GAG CTC EcolCRI Psp 124B I*(GAGCT C) RE1258 Psp 124B I*(GAGCT C) 078 EclXI C GGCCG Bse X3I RE1178 056 Eco105I TAC GTA Bsi SNI RE1236 071 Eco130I C CWWGG Bss T1I RE1198 062 Eco147I AGG CCT PceI RE1308 093 Eco24I GRGCY C FriOI RE1272 082 Eco31I GGTCTC (1/5) Bso31I RE1184 058 Eco32I GAT ATC EcoRV RE1262 079 Eco47I G GROCC Bme18I RE1148 047	Dse DI	GACNNNN [↓] NNGTC	Des D I	RE1372	077
Eam1105I GACNNN↓NNGTC Bme RI RV1150 048 Earl CTCTTC (1/4) Bst 6I RE1206 064 Ec/HKI GACNNN↓NNGTC Bme RI RV1150 048 Ec/136II GAG↓CTC EcolCRI Psp124B I*(GAGCT↓C) RE1258 078 Ec/XI C↓GGCCG BseX3I RE1178 056 Eco105I TAC↓GTA Bst SNI RE1236 071 Eco130I C↓CWWGG Bss T1I RE1198 062 Eco147I AGG↓CCT PceI RE1308 093 Eco24I GRGCY↓C FriOI RE1272 082 Eco31I GGTCTC (1/5) Bso31I RE1184 058 Eco32I GAT↓ATC EcoRV RE1262 079 Eco47I G↓GWCC Bme18I RE1148 047	Eag l	c∮ ggccg	Bse X3I	RE1178	056
Earl CTCTTC (1/4) Bst 6I RE1206 064 Ec HK GACNNN↓NNGTC Bme RI RV1150 048 Ec 136 GAG↓CTC EcolCR RE1258 078 Psp124B I*(GAGCT↓C) RE1312 094 Ec X C↓GGCCG Bse X3I RE1178 056 Eco105 TAC↓GTA Bst SNI RE1236 071 Eco130 C↓CWWGG Bss T1 RE1198 062 Eco147 AGG↓CCT Pce RE1308 093 Eco24 GRGCY↓C Fr O RE1272 082 Eco31 GGTCTC (1/5) Bso31 RE1184 058 Eco32 GAT↓ATC EcoRV RE1262 079 Eco47 G↓GWCC Bme 18 RE1148 047	Eam 1104I	CTCTTC (1/4)	Bst 61	RE1206	064
Ec/HKI GACNNN↓NNGTC Bme RI RV1150 048 Ec/136II GAG↓CTC EcolCRI Psp124B I*(GAGCT↓C) RE1258 RE1312 078 RE1312 Ec/XI C↓GGCCG Bse X3I RE1178 056 Eco105I TAC↓GTA Bst SNI RE1236 071 Eco130I C↓CWWGG Bss T1I RE1198 062 Eco147I AGG↓CCT PceI RE1308 093 Eco24I GRGCY↓C FriOI RE1272 082 Eco31I GGTCTC (1/5) Bso31I RE1184 058 Eco32I GAT↓ATC EcoRV RE1262 079 Eco47I G↓GWCC Bme18I RE1148 047	Eam 1105 I	GACNNN [↓] NNGTC	Bme RI	RV1150	048
Ec/136II $GAG^{\frac{1}{4}}CTC$ EcolCRI $PSp124B$ I*(GAGCT $^{\frac{1}{4}}C$) RE1258 RE1312 094 Ec/XI $C^{\frac{1}{4}}GGCG$ $BSeX3I$ RE1178 056 Eco105I $TAC^{\frac{1}{4}}GTA$ $BStSNI$ RE1236 071 Eco130I $C^{\frac{1}{4}}CWWGG$ $BSST1I$ RE1198 062 Eco147I $AGG^{\frac{1}{4}}CCT$ $PCeI$ RE1308 093 Eco24I $GRGCY^{\frac{1}{4}}C$ $FriOI$ RE1272 082 Eco31I $GGTCTC(1/5)$ $BSo31I$ RE1184 058 Eco32I $GAT^{\frac{1}{4}}ATC$ $EcoRV$ RE1262 079 $Eco47I$ $G^{\frac{1}{4}}GWCC$ $Bme18I$ RE1148 047	Ear l	CTCTTC (1/4)	Bst 61	RE1206	064
Psp124B I*(GAGCT $^{\downarrow}$ C) RE1312 094 Ecl XI C $^{\downarrow}$ GGCCG Bse X3I RE1178 056 Eco105I TAC $^{\downarrow}$ GTA Bst SNI RE1236 071 Eco130I C $^{\downarrow}$ CWWGG Bss T1I RE1198 062 Eco147I AGG $^{\downarrow}$ CCT PceI RE1308 093 Eco24I GRGCY $^{\downarrow}$ C FriOI RE1272 082 Eco31I GGTCTC (1/5) Bso31I RE1184 058 Eco32I GAT $^{\downarrow}$ ATC Eco RV RE1262 079 Eco47I G $^{\downarrow}$ GWCC Bme18I RE1148 047	Ec/ HKI	GACNNN [↓] NNGTC	Bme RI	RV1150	048
Eco105I TAC $^{\downarrow}$ GTA Bst SNI RE1236 071 Eco130I C $^{\downarrow}$ CWWGG Bss T1I RE1198 062 Eco147I AGG $^{\downarrow}$ CCT PceI RE1308 093 Eco24I GRGCY $^{\downarrow}$ C Fri OI RE1272 082 Eco31I GGTCTC (1/5) Bso 31I RE1184 058 Eco32I GAT $^{\downarrow}$ ATC Eco RV RE1262 079 Eco47I G $^{\downarrow}$ GWCC Bme18I RE1148 047	Ecl136II	GAG↓CTC			
Eco130I C\$\frac{1}{2}\$ CWWGG BssT1I RE1198 062 Eco147I AGG\$\frac{1}{2}\$ CCT PceI RE1308 093 Eco24I GRGCY\$\frac{1}{2}\$ C FriOI RE1272 082 Eco31I GGTCTC (1/5) Bso31I RE1184 058 Eco32I GAT\$\frac{1}{2}\$ ATC EcoRV RE1262 079 Eco47I G\$\frac{1}{2}\$ GWCC Bme18I RE1148 047	Ecl XI	c↓ggccg	Bse X3I	RE1178	056
$Eco147I$ $AGG^{\downarrow}CCT$ $PceI$ RE1308 093 $Eco24I$ $GRGCY^{\downarrow}C$ $FriOI$ RE1272 082 $Eco31I$ $GGTCTC(1/5)$ $Bso31I$ RE1184 058 $Eco32I$ $GAT^{\downarrow}ATC$ $EcoRV$ RE1262 079 $Eco47I$ $G^{\downarrow}GWCC$ $Bme18I$ RE1148 047	Eco 105 I	TAC [↓] GTA	Bst SNI	RE1236	071
Eco 24I GRGCY ↓ C Fri OI RE1272 082 Eco 31I GGTCTC (1/5) Bso 31I RE1184 058 Eco 32I GAT ↓ ATC Eco RV RE1262 079 Eco 47I G ↓ GWCC Bme18I RE1148 047	Eco 130 I	c↓cwwgg	Bss T1I	RE1198	062
Eco 31I GGTCTC (1/5) Bso 31I RE1184 058 Eco 32I GAT I ATC Eco RV RE1262 079 Eco 47I GI GWCC Bme18I RE1148 047	Eco 147 I	AGG↓CCT	Pcel	RE1308	093
Eco 32I GAT I ATC Eco RV RE1262 079 Eco 47I GI GWCC Bme18I RE1148 047	Eco 24 I	GRGCY↓C	FriOI	RE1272	082
Eco 47I G G G G Bme 18I RE1148 047	Eco 31I	GGTCTC (1/5)	Bso 31I	RE1184	058
	Eco 321	GAT↓ATC	Eco RV	RE1262	079
Eco 52I C [↓] GGCCG Bse X3I RE1178 056	Eco 47 I	G [↓] GWCC	Bme18I	RE1148	047
	Eco 52 I	c↓ggccg	Bse X3I	RE1178	056

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Eco 72I	CAC↓GTG	Psp CI	RE1314	095
Eco 81 I	CC↓TNAGG	Bse 211	RE1170	054
Eco 881	c↓ycgrg	Ama 87I	RE1122	039
Eco 911	G [↓] GTNACC	Psp EI	RE1316	095
EcolCRI	GAG [↓] CTC	EcolCRI Psp124BI* (GAGCT↓C)	RE1258 RE1312	078 094
Eco NI	CCTNN [↓] NNNAGG	Bst ENI	RE1216	066
Eco O65I	G [‡] GTNACC	PspEI	RE1316	095
Eco RI	G [↓] AATTC	Eco RI	RE1260	078
Eco RII	↓ccwgg	Bst 2UI* (CC ↓ WGG)	RE1202	063
Eco RV	GAT↓ATC	Eco RV	RE1262	079
Eco T14I	c [↓] cwwgg	Bss T1I	RE1198	062
Eco T22I	ATGCA↓T	Zsp 21	RE1370	111
Eco T38I	GRGCY↓C	Fri OI	RE1272	082
Egel	GGC↓GCC	Din l	RV1252	075
Ehe l	ggc↓gcc	Din l	RV1252	075
Erh l	c↓cwwgg	Bss T1 I	RE1198	062
Fau NDI	CA [↓] TATG	Fau NDI	RE1266	080
Fbal	T↓GATCA	Ksp 221	RE1288	086
Fb/I	GT [↓] MKAC	Fb/I	RE1268	081
Fokl	GGATG (9/13)	Fokl BstF5 I* (GGATG(2/0))	RE1270 RE1218	081 067
Fri O I	GRGCY↓C	Fri OI	RE1272	082
Fspl	TGC↓GCA	Acc 16I	RE1102	033
Fun ll	G↓AATTC	EcoRI	RE1260	078
Haell	RGCGC↓Y	Bst H2I	RE1222	068
Hae III	GG↓CC	Bsnl	RV1182	057
Нар іі	c∤cgg	Hpa ll Msp l	RE1282 RE1302	085 090
Hha l	gcg↓c	Asp LEI Bst HHI Hsp AI* (G [‡] CGC)	RE1130 RE1224 RE1284	042 068 085
Hin 11	GR↓CGYC	Bss NI	RV1194	061
Hin 61	g↓cgc	Asp LEI* (GCG ↓ C) Bst HHI* (GCG ↓ C) Hsp AI	RE1130 RE1224 RE1284	042 068 085
Hincll	GTY [↓] RAC	Hin dll	RE1274	083
HindII	GTY [↓] RAC	HindII	RE1274	083
HindIII	A↓AGCTT	HindIII	RE1276	083

List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases (cont'd)

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Hinfl	G↓ANTC	Hinfl	RE1278	084
Hin P1 I	g [†] cgc	Asp LEI* (GCG [↓] C) Bst HHI* (GCG [↓] C) Hsp AI	RE1130 RE1224 RE1284	042 068 085
Hpa l	GTT↓AAC	Нра і	RE1280	084
Hpa ll	c↓cgg	Hpa ll Msp l	RE1282 RE1302	085 090
Hph l	GGTGA (8/7)	Asu HPI	RE1134	043
Hpy CH4III	ACN↓GT	Bst 4Cl	RE1204	063
Hpy F3I	C↓TNAG	Bst DEI	RE1212	065
Hsp 921	GR [↓] CGYC	Bss NI	RV1194	061
Hsp Al	g↓cgc	Asp LEI* (GCG ↓ C) Bst HHI* (GCG ↓ C) Hsp AI	RE1130 RE1224 RE1284	042 068 085
Kas l	g [↓] gcgcc	DinI* (GGC ↓ GCC)	RV1252	075
Kpn 2I	T↓CCGGA	Bsp 13I	RE1186	058
Kpn l	GGTAC [‡] C	Acc 65I* (G [↓] GTACC) Kpn I	RE1104 RV1286	034 086
Ksp 221	T [↓] GATCA	Ksp 22I	RE1288	086
Ksp 6321	CTCTTC (1/4)	Bst 61	RE1206	064
Ksp Al	GTT↓AAC	Нра і	RE1280	084
Ksp l	ccdc†dd	Sfr 303I	RE1334	102
Kzo 91	[↓] GATC	Bss MI Bst MBI	RV1192 RE1228	060 069
Mam l	GATNN [↓] NNATC	Bse 81	RE1174	055
Mbi l	CCGCTC (-3/-3)	AccBSI	RE1110	035
Mbol	[↓] GATC	Bss MI Bst MBI	RV1192 RE1228	060 080
Mbo ll	GAAGA (8/7)	Mbo ll	RE1290	087
Mf/ l	R↓GATCY	Bst X2I	RE1242	073
Mhl	GDGCH↓C	Mh/ I	RE1292	088
MIsI	TGG↓CCA	Msp 201	RE1304	091
Mlu l	A↓CGCGT	Mlu l	RE1294	088
MIu NI	TGG↓CCA	Msp 201	RE1304	091
Mly 113I	gg↓cgcc	DinI* (GGC ↓ GCC)	RV1252	075
MnI I	CCTC (7/6)	Mn/ I	RE1296	089
Mph 1103I	ATGCA↓T	Zsp 21	RE1370	111
Mrol	T↓CCGGA	Bsp13I	RE1186	058
Mro NI	g↓ccggc	Mro NI	RE1298	089

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Mro XI	GAANN↓NNTTC	Mro XI	RE1300	090
Msc l	TGG↓CCA	Msp 201	RE1304	091
Mse l	T [↓] TAA	Tru 91	RE1350	106
Msp l	c∮cgg	Hpa ll Msp l	RE1282 RE1302	085 090
Msp 20 I	TGG↓CCA	Msp 201	RE1304	091
Msp A1I	CMG [↓] CKG	Msp A1I	RE1306	091
Msp CI	C↓TTAAG	Vha 464 I	RE1358	107
Msp R9I	CC↓NGG	Bmr FI	RV1154	049
Mval	cc↓wgg	Bst 2UI	RE1202	063
Mva 1269 I	GAATGC (1/-1)	Pct l	RE1310	093
Mvn l	cg↓cg	Bst FNI	RE1220	067
Nael	GCC↓GGC	Mro NI* (G↓CCGGC)	RE1298	089
Nar l	gg↓cgcc	DinI* (GGC ↓ GCC)	RV1252	075
Ncil	cc↓sgg	Ври МІ	RV1162	051
Ncol	CvCATGG	Bsp19I	RE1190	059
Ndel	CA [↓] TATG	Fau NDI	RE1266	080
Nde ll	[↓] GATC	Bss MI Bst MBI	RV1192 RE1228	069 069
Ngo MIV	g↓ccggc	Mro NI	RE1298	089
Nhel	G↓CTAGC	Asu NHI	RE1136	043
		Bmtl* (GCTAG [↓] C)	RE1156	049
Nla IV	GGN↓NCC	Bmi l	RV1152	048
Not l	GC [↓] GGCCGC	Cci NI	RE1248	074
Nru l	TCG↓CGA	Btu MI	RV1246	074
Nsb l	TGC↓GCA	Acc 16I	RE1102	033
Nsi l	ATGCA↓T	Zsp 21	RE1370	111
Nspl	RCATG ↓ Y	Bst NSI	RE1232	070
Nsp III	C↓YCGRG	Ama 871	RE1122	039
Nsp V	TT↓CGAA	Bpu 14I	RE1160	051
Pael	GCATG↓C	Sph l	RV1340	104
Pae R7I	C↓TCGAG	Sfr 274I	RE1332	102
Paul	g [†] cgcgc	Bse PI	RE1176	055
Pce l	AGG↓CCT	Pce l	RE1308	093
Pct l	GAATGC (1/-1)	Pct l	RE1310	093
Pdi l	gcc↓ggc	MroNI* (G↓CCGGC)	RE1298	089
Pdm l	GAANN [↓] NNTTC	Mro XI	RE1300	090

List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases (cont'd)

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Pf/ BI	CCANNNN↓NTGG	Acc B7I	RE1108	035
PfI FI	GACN↓NNGTC	Tth 111I	RE1356	107
Pf/ MI	CCANNNN [↓] NTGG	Acc B7I	RE1108	035
Pho I	gg↓cc	Bsnl	RV1182	057
Pin Al	A↓CCGGT	Asi GI	RE1126	041
Pma Cl	CAC↓GTG	Psp CI	RE1314	095
Pmi ll	CAC↓GTG	Psp CI	RE1314	095
Psh Al	GACNN↓NNGTC	Bst PAI	RE1234	071
Psh BI	AT [↓] TAAT	Vspl	RE1362	108
Psp 124BI	GAGCT↓C	Psp 124BI	RE1312	094
Psp 1406 I	AA↓CGTT	Acl	RE1112	036
Psp 61	↓ ccwgg	Bst2UI* (CC↓WGG)	RE1202	063
Psp CI	CAC [↓] GTG	Psp CI	RE1314	095
Psp GI	↓ccwgg	Bst2UI* (CC↓WGG)	RE1202	063
Psp EI	G [‡] GTNACC	Psp EI	RE1316	095
Psp N4I	GGN [↓] NCC	Bmi l	RV1152	048
Psp OMI	e† geccc	Apal* (GGGCC	RE1124 RE1318	040 096
Psp PI	G [↓] GNCC	Asp S91	RE1132	042
Pst I	CTGCA [↓] G	Pst l	RE1320	096
Psu l	R↓GATCY	Bst X21	RE1242	073
Psy l	GACN↓NNGTC	Tth 111I	RE1356	107
Rsal	GT↓AC	Rsa l	RE1324	097
Rsr II	cg↓gwccg	Rsr 2l	RE1374	098
Rsr 2l	cg↓gwccg	Rsr 2l	RE1374	098
Sacl	GAGCT↓C	EcoICRI* (GAG [↓] CTC) Psp124BI	RE1258 RE1312	078 094
Sac ll	ccec†ee	Sfr 303I	RE1334	102
Sal 1	G↓TCGAC	Sall	RV1326	099
Sau 3AI	[↓] GATC	Bss MI	RV1192	060
Sau 96I	G [†] GNCC	Bst MBI Asp S9I	RE1228 RE1132	069 042
Sbf l	CCTGCA [↓] GG	Sbfl	RE1328	100
Scal	AGT↓ACT	Bmc Al	RV1146	047
ScrFI	CC↓NGG	Bmr FI	RV1154	049
Sdal	CCTGCA [↓] GG	Sbf l	RE1328	100
Sdul	GDGCH↓C	Mh/ I	RE1292	088

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Sfa NI	GCATC (5/9)	Sfa NI	RE1376	101
Sfi l	GGCCNNNN [↓] NGGCC	Sfi l	RE1330	101
Sfo l	ggc↓gcc	Din l	RV1252	075
Sfr 274 I	C↓TCGAG	Sfr 274I	RE1332	102
Sfr 303 I	ccgc↓gg	Sfr 303 I	RE1334	102
Sfu l	TT [↓] CGAA	Bpu 14 I	RE1160	051
Sgr BI	ccgc↓gg	Sfr 303I	RE1334	102
Sinl	G [‡] GWCC	Bme 18I	RE1148	047
Slal	C↓TCGAG	Sfr 274I	RE1332	102
Smal	ccc† ggg	Smal Xmal* (C [‡] CCGGG)	RE1336 RV1366	103 110
Smi l	ATTT [↓] AAAT	Smi l	RE1338	103
Smi MI	CAYNN ↓ NNRTG	Smi M I	RE1378	104
Sna BI	TAC↓GTA	Bst SNI	RE1236	071
Spa HI	GCATG↓C	Sph l	RV1340	104
Spel	AvCTAGT	Ah/ I	RE1118	038
Sph l	GCATG↓C	Sph l	RV1340	104
Sse 8387I	CCTGCA↓GG	Sbfl	RE1328	100
Sse 91	[↓] AATT	Sse 91	RE1342	105
Sse BI	AGGvCCT	Pce l	RE1308	093
Ssp BI	T [‡] GTACA	Bst AUI	RE1208	064
Sspl	AAT [↓] ATT	Ssp l	RE1344	105
Sst l	GAGCT↓C	EcoICRI* (GAG ↓ CTC)	RE1258	078
		Psp 124BI	RE1312	094
Strl	C↓TCGAG	Sfr 274I	RE1332	102
Stu l	AGG [↓] CCT	Pce l	RE1308	093
Sty l	C [↓] CWWGG	Bss T1I	RE1198	062
Sty D4I	↓ CCNGG	BmrFI* (CC ↓ NGG)	RV1154	049
Swal	ATTT ↓ AAAT	Smi I	RE1338	103
Taa l	ACN [↓] GT	Bst 4Cl	RE1204	063
Taq l	T [↓] CGA	Taq l	RE1346	106
Tas l	↓ AATT	Sse 91	RE1342	105
Tel I	GACN NNGTC	Tth1111	RE1356	107
Tli I	C↓TCGAG	Sfr 274I	RE1332	102
Tru 11	T [↓] TAA	Tru 91	RE1350	106
Tru 91	T↓TAA	Tru 91	RE1350	106

List of Commercial Isoschizomers Corresponding to Vivantis Restriction Endonucleases (cont'd)

Commercial Restriction Endonuclease	Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonuclease	Product No	Page
Tsp 509 I	↓ AATT	Sse 91	RE1342	105
Tsp EI	[↓] AATT	Sse 91	RE1342	105
Tth 111 I	GACN [↓] NNGTC	Tth 111 I	RE1356	107
Van 91 I	CCANNNN [↓] NTGG	Acc B7I	RE1108	035
Vha 464 I	C↓TTAAG	Vha 464 I	RE1358	107
Vnel	G [↓] TGCAC	Vnel	RE1360	108
Vpa K11BI	G [†] GWCC	Bme 18I	RE1148	047
Vsp l	AT [↓] TAAT	Vspl	RE1362	107
Xag l	CCTNN [↓] NNNAGG	Bst ENI	RE1216	066
Xap I	R [↓] AATTY	Acs l	RE1114	036
Xbal	T [‡] CTAGA	Xbal	RV1364	109
Xce l	RCATG [↓] Y	Bst NSI	RE1232	070
Xho l	C↓TCGAG	Sfr 274I	RE1332	102
Xho ll	R↓GATCY	Bst X21	RE1242	073
Xma CI	c↓ccggg	Smal* (CCC [↓] GGG) Xmal	RE1336 RV1366	103 110
Xmal	c↓ccggg	Smal* (CCC	RE1336 RV1366	103 110
Xma JI	C [↓] CTAGG	Asp A2I	RE1128	041
Xmi l	GT∮MKAC	Fb/ I	RE1268	081
Xmn l	GAANN [↓] NNTTC	Mro XI	RE1300	090
Zhol	AT [↓] CGAT	Bsh VI	RV1180	056
Zral	GAC↓GTC	AatII* (GACGT [↓] C) ZraI	RE1100 RE1368	033 111
Zrm l	AGT [↓] ACT	Bmc Al	RV1146	047
Zsp 2 I	ATGCA↓T	Zsp 2 I	RE1370	111

Base Nomenclature

B: C/G/T M: A/C
D: A/G/T Y: C/T
H: A/C/T R: A/G
V: A/C/G S: C/G
K: G/T W: A/T
N: A/C/G/T

Alphebetical List of Recognition Specificities of Vivantis Restriction Endonucleases

Recognition Sequence / Cleavage Site 5→3	Vivantis Restriction Endonucleases	Prototype	Product No	Page
AA↓CGTT	Ac/ I	Ac/I	RE1112	036
A↓AGCTT	Hin dIII	HindIII	RV1276	083
AAT ↓ ATT	Sspl	Ssp l	RE1344	105
↓AATT	Sse 91	Tsp El	RE1342	105
A↓CCGGT	Asi GI	Age l	RE1126	041
A↓CGCGT	Mlul	Mlul	RE1294	088
ACN↓GT	Bst 4CI	Tsp 4CI	RE1204	063
A [↓] CTAGT	Ahl l	Spel	RE1118	038
ACTGG (1/-1)	Bse I	Bsrl	RE1166	053
A↓GATCT	Bgl ll	Bg/ II	RV1144	046
AG↓CT	Alu l	Alu l	RE1120	038
AGG↓CCT	Pcel	Stu l	RE1308	093
AGT↓ACT	Bmc Al	Scal	RV1146	047
AT↓CGAT	Bsh VI	Clal	RV1180	056
ATGCA↓T	Zsp 21	AvaIII	RE1370	111
AT [↓] TAAT	Vspl	Vsp l	RE1362	108
ATTT [↓] AAAT	Smi I	Swal	RE1338	103
CAANNNNN [↓] NTGG	Bst XI	Bst XI	RE1240	072
CAC↓GTG	Psp CI	PmaCl	RE1314	095
CACNNN [↓] GTG	Dra III	Dra III	RE1256	076
CAG↓CTG	Pvu ll	Pvu II	RE1322	097
CA [↓] TATG	Fau NDI	Ndel	RE1266	080
CAYNN ↓ NNRTG	Smi MI	Ms/I	RE1378	104
CCANNNN [↓] NTGG	Acc B7I	Pfl MI	RE1108	035
C↓CATGG	Bsp 19 I	Ncol	RE1190	059
c↓ccggg	Xma l	Sma l*	RV1366	110
ccc↓ggg	Smal	Smal	RE1336	103
ccgc↓gg	Sfr 3031	SacII	RE1334	102
CCGCTC (-3/-3)	Acc BSI	Bsr BI	RE1110	035
c↓cgg	Hpa II	Hpa II	RE1282	085
c↓cgg	MsPI	Hpa ll	RE1302	090
CC [↓] NGG	Bmr FI	Scr FI	RV1154	049
CCNNNNN ↓ NNGG	Afi l	Bsi YI	RV1116	037
C↓CRYGG	Bst DSI	Dsa l	RE1214	066
cc↓sgg	Ври МІ	Caull	RV1162	051
C [‡] CTAGG	Asp A2I	AvrII	RE1128	041



^{*} Indicates neoschizomer

Alphebetical List of Recognition Specificities of Vivantis Restriction Endonucleases (cont'd)

Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonucleases	Prototype	Product No	Page
CCTC (7/6)	MnI	MnI	RE1296	089
CCTGCA [↓] GG	Sbfl	Sse 8387I	RE1328	100
CCTNAGC (-5/-2)	Bpu 10I	Bpu 10I	RE1158	050
CC↓TNAGG	Bse 21I	Saul	RE1170	054
CCTNN↓NNNAGG	Bst ENI	Eco NI	RE1216	066
cc↓wgg	Bst 2UI	EcoRII *	RE1202	063
c↓cwwgg	Bss T1I	Sty l	RE1198	062
cg↓cg	Bst FNI	Fnu DII	RE1220	067
c↓ggccg	Bse X3I	Xma lll	RE1178	056
cg↓gwccg	Rsr 21	Rsr II	RE1374	098
CGRY↓CG	Bst MCI	Mcrl	RE1230	070
CMG [↓] CKG	Msp A1I	Nsp BII	RE1306	091
C↓TCGAG	Sfr 274I	Xhol	RE1332	102
CTCTTC (1/4)	Bst 61	Ksp 6321	RE1206	064
CTGCA↓G	Pst l	Pst l	RE1320	096
C↓TNAG	Bst DEI	Ddel	RE1212	065
C↓TTAAG	Vha 464 I	Af/ II	RE1358	107
C↓YCGRG	Ama 871	Aval	RE1122	039
GAAGA (8/7)	Mbo ll	Mbo ll	RE1290	087
GAAGAC (2/6)	Bst V2I	Bbv ll	RE1238	072
GAANN [↓] NNTTC	Mro XI	Xmn l	RE1300	090
GAATGC (1/-1)	Pct l	Bsm l	RE1310	093
G↓AATTC	EcoRI	Eco RI	RE1260	078
GAC [↓] GTC	Zral	Aat II*	RE1368	111
GACGT↓C	Aat II	Aat II	RE1100	033
GACN↓NNGTC	Tth 111 I	Tth 111	RE1356	107
GACNN↓NNGTC	Bst PAI	Psh Al	RE1234	071
GACNNN↓NNGTC	Bme RI	Eam 1105 I	RV1150	048
GACNNNN ↓ NNGTC	Dse DI	Drd l	RE1372	077
GAG [↓] CTC	EcolCRI	Sac l*	RE1258	078
GAGCT↓C	Psp 124BI	Sacl	RE1312	094
G↓ANTC	Hinfl	Hinfl	RE1278	084
GAT [↓] ATC	Eco RV	Eco RV	RE1262	079
[↓] GATC	Bss MI	Mbol	RV1192	060
[↓] GATC	Bst MBI	Mbol	RE1228	069
GATNN↓NNATC	Bse 81	Bsa Bl	RE1174	055

Alphebetical List of Recognition Specificities of Vivantis Restriction Endonucleases (cont'd)

Recognition Sequence / Cleavage Site 5'→3'	Vivantis Restriction Endonucleases	Prototype	Product No	Page
GCAATG (2/0)	Bse 3DI	Bsr DI	RE1172	054
GCATC (5/9)	Sfa NI	Sfa NI	RE1376	101
GCATG↓C	Sph l	Sph l	RV1340	104
g↓ccggc	Mro NI	Nae l*	RE1289	089
GCCNNNN [↓] NGGC	Bg/ I	Bg/ I	RV1142	046
G↓CGC	Hsp Al	Hha l*	RE1284	085
gcg↓c	Asp LEI	Hhal	RE1130	042
gcg↓c	Bst HHI	Hha l	RE1224	068
G↓CGCGC	Bse PI	Bse PI	RE1176	055
gc↓ggccgc	CciNI	Not	RE1248	074
G↓CTAGC	Asu NHI	Nhel	RE1136	043
GCTAG↓C	Bmt l	Nhe l*	RE1156	049
GC↓TNAGC	Bsp 1720 I	Esp l	RE1188	059
GDGCH↓C	Mhl I	Sdu l	RE1292	088
G↓GATCC	Bam HI	Bam HI	RV1138	044
GGATG (2/0)	Bst F51	Fok l*	RE1218	067
GGATG (9/13)	Fok l	Fok l	RE1270	081
gg↓cc	Bsnl	HaeIII	RV1182	057
GGCCNNNN [↓] NGGCC	Sfi l	Sfil	RE1330	101
ggc↓gcc	Din l	Nar l*	RE1252	075
g↓ggccc	Psp OMI	Apa l*	RE1318	096
gggcc↓c	Apa l	Apa l	RE1124	040
G↓GNCC	Asp S9I	Asu l	RE1132	042
GGN↓NCC	Bmi l	Nla IV	RV1152	048
G [↓] GTACC	Acc 651	Kpn i*	RE1104	034
GGTAC↓C	Kpn l	Kpnl	RV1286	086
GGTCTC (1/5)	Bso 311	Eco 31I	RE1184	058
GGTGA (8/7)	Asu HPI	Hph l	RE1134	043
G [↓] GTNACC	Psp EI	Bst EII	RE1316	095
G [†] GWCC	Bme 18I	Ava ll	RE1148	047
G [↓] GYRCC	Acc B1I	Hgi CI	RE1106	034
GR↓CGYC	Bss NI	Acy l	RV1194	061
GRCGY↓C	Fri OI	Hgi JII	RE1272	082
GT↓AC	Rsal	Rsal	RE1324	097
GTA [↓] TAC	Bss NAI	Snal	RE1196	061
G↓TCGAC	Sall	Sall	RV1326	099

Alphabetical List of Recognition Specificities of Vivantis Restriction Endonucleases (cont'd)

GTCTC (1/5) Bst MAI Bsm AI RE1226 G¹TGCAC VneI ApaLI RE1360 GT¹MKAC FbII AccI RE1268 GTT¹AAC HpaI HpaI RE1280 GTY¹RAC HindII HindII RE1274 GWGCW¹C Bbv12I HgiAI RE1140 R¹AATTY AcsI ApoI RE1114 RCATG¹Y Bst NSI NspI RE1232 R¹CGGY Bse 118I Cfr10I RE1168 R¹GATCY Bst X2I XhoII RE1242 RGCGC¹Y Bst H2I HaeII RE1222 TAC¹GTA Bst SNI Sna BI RE1236 T¹CCGGA Bsp13I BspMII RE1186	069 108
GT $^{\downarrow}$ MKAC Fb/I AccI RE1268 GTT $^{\downarrow}$ AAC HpaI RE1280 GTY $^{\downarrow}$ RAC HindII HindII RE1274 GWGCW $^{\downarrow}$ C Bbv12I HgiAI RE1140 R\$\frac{1}{4}AATTY AcsI Apol RE1114 RCATG $^{\downarrow}$ Y Bst NSI NSpI RE1232 R\$\frac{1}{4}CCGGY Bse 118I Cfr 10I RE1168 R\$\frac{1}{4}CATCY Bst X2I XhoII RE1242 RGCGC $^{\downarrow}$ Y Bst H2I HaeII RE1226	108
GTT AAC Hpal Hpal RE1280 GTY RAC Hindll Hindll RE1274 GWGCW C Bbv12I HgiAl RE1140 RE1140 RATTY ACSI ACSI RCATG Y BSt NSI NSpl RE1232 RICGGY BSe 118I Cfr 10I RE1168 RIGGC Y BSt X2I XhoII RE1222 TAC GTA BSt SNI Sna BI RE1236	
GTY $^{\downarrow}$ RAC HindII HindII RE1274 GWGCW $^{\downarrow}$ C Bbv12I HgiAI RE1140 R $^{\downarrow}$ AATTY AcsI Apol RE1114 RCATG $^{\downarrow}$ Y Bst NSI NspI RE1232 R $^{\downarrow}$ CCGGY Bse 118I Cfr 10I RE1168 R $^{\downarrow}$ GATCY Bst X2I XhoII RE1242 RGCGC $^{\downarrow}$ Y Bst H2I HaeII RE1222 TAC $^{\downarrow}$ GTA Bst SNI Sna BI RE1236	081
GWGCW $^{\downarrow}$ C Bbv12I HgiAI RE1140 R $^{\downarrow}$ AATTY AcsI Apol RE1114 RCATG $^{\downarrow}$ Y Bst NSI Nspl RE1232 R $^{\downarrow}$ CCGGY Bse 118I Cfr10I RE1168 R $^{\downarrow}$ GATCY Bst X2I XhoII RE1242 RGCGC $^{\downarrow}$ Y Bst H2I HaeII RE1222 TAC $^{\downarrow}$ GTA Bst SNI Sna BI RE1236	084
R^{\downarrow} AATTY $AcsI$ $ApoI$ RE1114 $RCATG^{\downarrow}Y$ Bst NSI $NspI$ RE1232 R^{\downarrow} CCGGY Bse 118I Cfr 10IRE1168 R^{\downarrow} GATCY Bst X2I $XhoII$ RE1242 $RGCGC^{\downarrow}Y$ Bst H2I $HaeII$ RE1222 TAC^{\downarrow} GTA Bst SNI Sna BIRE1236	083
RCATG $^{\downarrow}$ YBst NSINspIRE1232R $^{\downarrow}$ CCGGYBse 118ICfr 10IRE1168R $^{\downarrow}$ GATCYBst X2IXhoIIRE1242RGCGC $^{\downarrow}$ YBst H2IHaeIIRE1222TAC $^{\downarrow}$ GTABst SNISna BIRE1236	045
R^{\downarrow} CCGGY $Bse118I$ $Cfr10I$ RE1168 R^{\downarrow} GATCY $BstX2I$ $XhoII$ RE1242RGCGC $^{\downarrow}$ Y $BstH2I$ $HaeII$ RE1222TAC $^{\downarrow}$ GTA $BstSNI$ $SnaBI$ RE1236	036
R \downarrow GATCYBst X2IXhoIIRE1242RGCGC \downarrow YBst H2IHaeIIRE1222TAC \downarrow GTABst SNISna BIRE1236	070
RGCGC	053
TAC ¹ GTA Bst SNI Sna BI RE1236	073
	068
T [↓] CCGGA Bsp13I BspMII RE1186	071
	058
T [↓] CGA TaqI TaqI RE1346	106
TCG CGA Btu MI Nru I RV1246	074
T [↓] CTAGA XbaI XbaI RV1364	109
T GATCA KSp 22I Bc/I RE1288	086
TGC GCA Acc 16I Mst I RE1102	033
TGG CCA Msp 201 Ball RE1304	091
T GTACA Bst AUI Bsp1407I RE1208	064
T TAA Tru 9I Msel RE1350	106
TT CGAA Bpu14I AsuII RE1160	051
TTT I AAA Dral Ahalii RE1254	076
YAC GTR Bst BAI Bsa AI RE1210	065

* Indicates neoschizomer

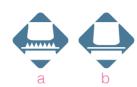
Base Nomenclature

B: C/G/T	M: A/C
D: A/G/T	Y : C/T
H: A/C/T	R: A/G
V : A/C/G	S: C/G
K: G/T	W: A/T
	N: A/C/G/T

Icons Description



Color-Tag Buffer System



Thermal Inactivation

- a indicates possible thermal inactivation of enzyme (65°C or 80°C for 20 minutes).
- b indicates enzyme cannot be thermally inactivated.



Star Activity

Indicates restriction endonuclease has star activity under non-standard conditions.



Recombinant Enzyme

Indicates enzyme is purified from a recombinant source.



Reaction Temperature

Indicates optimum incubation temperature for the enzyme.



Methylation Sensitivity

Cleavage is blocked when substrate DNA is methylated either by CG, dam or dcm methylase.

Aatll











Concentration

1-10u/µl

5'...GACGT¹C...3' 3'...C_↑TGCAG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.6), 50mM NaCl, 0.1mM EDTA, 200µg/ml BSA, 1mM DTT and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Aat*II, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Aat*II for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A (Diluent)

- * High enzyme concentration may result in Star Activity.
- * Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1100	200u



λDNA 0.7% Agarose

a -	Digestion	n aπer	I I	iour
b -	Digestion	n after	16	hour

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
50%	50%	25%	50%	100%		

Accl

Please refer to FbII (RE1268 - page 081)

Acc16| {Mst|}

Concentration

1-10u/µl

5'...TGC[↓]GCA...3' 3'...ACG_↑CGT...5'

Reaction Conditions

1X Buffer UB

25mM Tris-acetate (pH 7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol, and 50μg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA, and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Acc*16l, more than 80% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Acc*16l for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

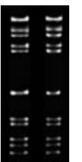
Ordering Information

Catalog No	Pack Size
RE1102	200u

UBBff







a b λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 10% 50% 50% 75% 50%

Acc65| {Kpn|*}

Concentration

10-30u/ul

5'...G¹GTACC...3' 3'...CCATG₁G...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCI, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assav

After 10-fold overdigestion with *Acc*65I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Acc*65I for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dcm-methylation (C^mCWGG): GGTACCWGG

Ordering Information

Catalog No	Pack Size
RE1104	500u

V4_{Bff}









λDNA (dam-&dcm-), 0.7% Agarose

a - Digestion after 1 hourb - Digestion after 16 hours

AccB1I {HgiCI}

Concentration

5-10u/µl

5'...G[↓]GYRCC...3' 3'...CCRYG_↑G...5'

Reaction Conditions

1X Buffer AccBII

10mM Tris-HCI (pH 7.5 at 25°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Acc*B1I, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of AccB1I for 16 hours at 37°C.

Supplied with 10X Buffer Acc B1I,10X Buffer UB and Viva Buffer A (Diluent)

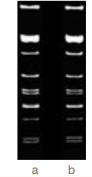
* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size	
RE1106	500u	







λDNA 1% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	75%	50%	50%	75%	



AccB7 | { PflM|}

Concentration

5u/µl

5'... CCANNNN¹NTGG...3' 3'...GGTN+NNNNACC...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH7.5 at 30°C). 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

AccBSI {BsrBI}

5'...CCG[‡]CTC...3'

3'...GGC+GAG...5'

Reaction Conditions

Storage Buffer

Thermal Inactivation 65°C for 20 minutes

2-mercaptoethanol, 200µg/ml BSA

and 50% glycerol. Store at -20°C.

Concentration

5-20u/ul

After 5-fold overdigestion with AccB7 I, 95% of the DNA fragments can be ligated and recut.









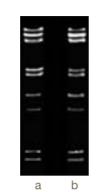
Overdigestion Assav

An unaltered banding pattern was observed after 1µg of DNA was digested with 2.5u of AccB7I for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

High enzyme concentration may result in Star Activity.

* Blocked by overlapping dcm-methylation (OMCWGG): CCANNNCCTGG or CAGGNNNTGG



λDNA (dam-&dcm-) 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1108	100u

Activity in Reaction Buffer V2 V3 V4 V5 V1 25% 100% 100% 100% 75%







Ligation / Recutting Assav After 5-fold overdigestion with

AccBSI, 90% of the DNA fragments can be ligated and of these 50% can be recut.

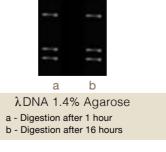
Overdigestion Assay

1X Buffer V5 An unaltered banding pattern was 30mM Tris-acetate (pH 7.9 at observed after 1µg of DNA was 30°C), 10mM Mg-acetate, 60mM digested with 10u of AccBSI for 16 K-acetate, and 100µg/ml BSA. hours at 37°C. Incubate at 37°C.

Supplied with 10X Buffer V5,10X 10mM Tris-HCl (pH 7.5), 100mM Buffer UB and Viva Buffer A. NaCl, 0.1mM EDTA, 7mM (Diluent)

Ordering Information

Catalog No	Pack Size
RE1110	500u



Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
100%	100%	75%	75%	100%	

AcII

Concentration

1-3u/ul

5'...AA[‡]CGTT...3' 3'...TTGC+AA...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 0.05% Triton X-100, 200 µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 3-fold overdigestion with Ac/I. 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 1.5u of Ac/I for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1112	100u











λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	75%	50%	75%	100%

Acsl {Apol}

Concentration

20-50u/µl

5'...R[‡]AATTY...3' 3'...YTTAA†R...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with Acsl, more than 95% of the DNA fragments can be ligated and recut.

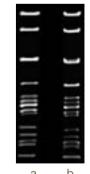
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Acsl for 16 hours at 50°C.

Supplied with 10X Buffer V4,10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1114	500u



λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
25%	75%	75%	100%	50%	



Please refer to BssNI (RV1194 – page 061)

Afil {BsiYI}

Concentration

2-10u/µl

5'...CCNNNNN¹NNGG...3' 3'...GGNN+NNNNNCC...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM KCI, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and 0.15% Triton X-100, and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assav

After 6-fold overdigestion with Afil, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Afil for 16 hours at 50°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

а

λDNA 1% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RV1116	500u

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
100%	75%	10%	100%	100%		

Please refer to Vha464I (RE1358 - page 107)

Aae

Please refer to AsiGI (RE1126 - page 041)

Ahdl

Please refer to Bme RI (RV1150 - page 048)

Ahll {Spel}

Concentration

10-30u/µl

5'...A[‡]CTAGT...3' 3'...TGATC₁A...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with Ahll, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of AhlI for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1118	300u









T7 DNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
100%	100%	50%	75%	75%	

Alul

Concentration

1-3u/µl

5'...AG[‡]CT...3' 3'...TC₁GA...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 3-fold overdigestion with Alul, 70% of the DNA fragments can be ligated and recut.

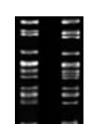
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 2u of Alul for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

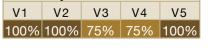
Ordering Information

Catalog No	Pack Size
RE1120	50u



λDNA 1% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5





Alw21I

Please refer to Bbv12I (RE1140 - page 045)

Alw261

Please refer to Bst MAI (RE1226 – page 069)

Alw441

Please refer to Vnel (RE1360 - page 108)

Ama87I {AvaI}

Concentration 10-30u/µl

5'...C¹YCGRG...3'

5'...C+YCGRG...3' 3'...GRGCY₁C...5'

Reaction Conditions

1X Buffer Ama87I 10mM Tris-HCI (pH8.5 at 30°C), 10mM MgCI₂, 100mM KCI, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM KH₂PO₄ (pH 7.2), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Ama*87I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of *Ama*87I for 16 hours at 37°C.

Supplied with 10X Buffer Ama87I, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1122	600u



λDNA 0.7% Agarose

a - Digestion after 1 hourb - Digestion after 16 hours

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
10%	25%	50%	75%	10%		

Aor13HI

Please refer to Bsp13I (RE1186 - page 058)

Apal

Concentration

10 - 30u/µl

5'...GGGCC¹C...3' 3'...C₁CCGGG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with *Apa*I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Apa*I for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dcm-methylation (CC^mWGG): GGGCCCWGG

Ordering Information

Catalog No	Pack Size
RE1124	600u











a b

λDNA (dam⁻&dcm⁻), (BamH I Digest) 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
75%	75%	75%	75%	100%		

ApaLl

Please refer to Vnel (RE1360 - page 108)

Apol

Please refer to Acsl (RE1114 - page 036)

Asel

Please refer to Vspl (RE1362 - page 108)

AsiGI {Agel}

Vivantis Technologies . Product Catalog Volume IV

Concentration

3-5u/µl

5'...A[‡]CCGGT...3' 3'...TGGCC₁A...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with AsiGI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

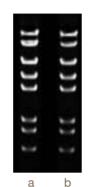
An unaltered banding pattern was observed after 1µg of DNA was digested with 3u of AsiGI for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1126	100u





λDNA 0.7% Agarose a - Digestion after 1 hour

Digestion		

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
50%	ó <u>1</u>	00%	100%	50%	75%

AspA2I {Avr II}

Concentration

10-20u/µl

5'...C[‡]CTAGG...3' 3'...GGATC₁C...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl_a, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with AspA2I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 15u of AspA2I for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

Catalog	No	Pack Size
RE1128		200u







V5

6 100%



λDNA (HindIII Digest) 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

talog No	Pack Size	Activity in Reaction Buffer				
	200u	V1	V2	V3	V4	V5
1128		25%	75%	75%	100%	1009

AspEl

Please refer to BmeRI (RV1150 - page 048)

AspLEI {Hhal}

Concentration

20-50u/ul

5'... GCG[‡]C...3' 3'...C_†GCG...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with AspLEI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of AspLEI for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1130	500u









λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
10%	50%	100%	25%	10%

AspS9I {AsuI}

Concentration

10-30u/µl

5'...G¹GNCC...3' 3'...CCNG₁G...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10 mM Tris-HCI (pH 7.5), 50 mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with AspS9I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of AspS9I for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dcm-methylation (OMCWGG): GGNCCWGG

Ordering Information

Catalog No	Pack Size
RE1132	600u







λDNA (dam-&dcm-) 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 50% 75% 100% 50%

AsuHPI {HphI}

Concentration 2-5u/µl

5'...GGTGA(N)₈¹...3' 3'...CCACT(N)₇†...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes







After 3-fold overdigestion with AsuHPI, 30% of the DNA fragments can be ligated and recut.

Ligation / Recutting Assay

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of AsuHPI for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (GMATC): GGTGAT

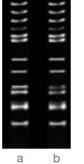
Ordering Information

Catalog No	Pack Size
RE1134	100u



λDNA (dam & dcm) 1.4% Agarose

alog No	Pack Size	4
		٧
1134	100u	25



a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
25%	75%	100%	75%	100%	



Concentration

10 - 20u/µl

5'...G¹CTAGC...3' 3'...CGATC₁G...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Condition

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with AsuNHI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of AsuNHI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1136	300u











λDNA (HindIII Digest) 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	75%	75%	75%	100%	

Concentration

40-150u/µl

5'...G¹GATCC...3' 3'...CCTAG_↑G...5'

1X Buffer UB

BSA. Incubate at 37°C.

NaCl, 0.1mM EDTA, 1mM DTT, and 50% glycerol. Store at -20°C.

65°C for 20 minutes

Please refer to Bpu14I (RE1160 - page 051)

Please refer to Ama87I (RE1122 - page 039)

Please refer to Bme18I (RE1148 - page 047)

Please refer to Acc16I (RE1102 - page 033)

Please refer to Asp A2I (RE1128 - page 041)

Please refer to Msp20l (RE1304 – page 091)







BamHI

Ball

Aval

Reaction Conditions

25mM Tris-acetate (pH7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol, and 50µg/ml

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM 0.15% Triton X-100, 200µg/ml BSA

Thermal Inactivation

Ligation / Recutting Assay

After 40-fold overdigestion with BamHI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BamHI for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RV1138	2500u



a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	75%	50%	75%	50%



Banl
Please refer to Acc B11 (RE1106 – page 034)

Ban | Please refer to Fri OI (RE1272 – page 082)

BbrPl Please refer to PspCI (RE1314 – page 95)

Bbs Please refer to Bst**V2I** (RE1238 – page 072)

Bbv12l {HgiAl}

Concentration

1-5u/µl

5'...GWGCW¹C...3' 3'...C₁WCGWG...5'

Reaction Conditions

1X Buffer Bbv12I 20mM Tris-HCI (pH8.5), 10mM MgCl₂, 20mM NaCl, and 1mM DTT.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Bcll

Ligation / Recutting Assay

After 5-fold overdigestion with *Bbv*12I, more than 90% of the DNA fragments can be ligated and recut.

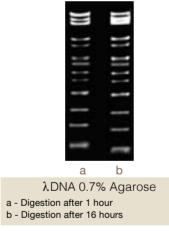
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of *Bbv*12l for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1140	100u



Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
10%	25%	75%	25%	25%		

Please refer to Ksp22 I (RE1288 - page 086)

Bgll

Concentration

5-30u/µl

5'...GCCNNNN¹NGGC...3' 3'...CGGN₁NNNNCCG...5'

Reaction Conditions

1X Buffer Bgll 20mM Tris-HCl (pH 8.5), 10mM MgCl₂, 200mM NaCl, and 1mM DTT. Incubate at 37°C.

Storage Condition

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *BgI*I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Bg/I* for 16 hours at 37°C.

Supplied with 10X Buffer *BgI*I, 10X Buffer UB and Viva Buffer A. (Diluent)

*High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RV1142	600u











λDNA 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
25%	10%	10%	10%	25%	

Bgl II

Concentration

10-30u/µl

5'...A[‡]GATCT...3' 3'...TCTAG_↑A...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCl (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with *BgI*II, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Bg/*II for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1144	300u





λDNA 0.7% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	75%	75%	100%	75%	



Blpl

Please refer to Bsp1720I (RE1188 – page 059)

BmcAl (Scal)

Concentration

1-10u/ul

5'...AGT[‡]ACT...3' 3'...TCA+TGA...5'

Reaction Conditions

0.5X Buffer UB

12.5mM Tris-acetate (pH 7.6 at 30°C), 5mM Mg-acetate, 50mM K-acetate, 3.5mM 2-mercaptoethanol, and 25µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with BmcAl. 70% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BmcAl for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

200u

λDNA 0.7% Agarose a - Digestion after 1 hour

75% 75% 10% 10%

b - Digestion after 16 hours

Ordering Informatio	n					
Catalog No Pack Size		Act	ivity in	React	ion Bu	ffer
	1 451 5125	V1	V2	V3	V4	V5

BmeRI (Eam1105I)

Concentration

5-10u/µl

5'...GACNNN¹NNGTC...3' 3'...CTGNN+NNNCAG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *Bme*R I. about 5% of the DNA fragments can be ligated.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of BmeRI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)



λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Catalog No	Pack Size
RV1150	200u

Ordering Information

Activity in Reaction Buffer

V1	V2	V3	V4	V5
75%	75%	10%	10%	100%

Bme18| {Ava||}

Concentration

5-20u/µl

5'...G¹GWCC...3' 3'...CCWG₁G...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with Bme181, more than 90% of the DNA fragments can be ligated.

Overdigestion Assay

RV1146

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bme18I for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1148	600u









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ı į a b

λDNA 1.0% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 50% 75% 100% 100% 100%

Bmil {Nla|V}

Concentration

10-30u/ul

5'...GGN¹NCC...3' 3'...CCN+NGG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with Bmil, 95% of the DNA fragments can be ligated and recut.

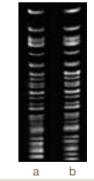
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Bmil for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1152	400u



λDNA 1.4% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	75%	75%	100%	100%



BmrFI {ScrFI}

Concentration 10-30u/µl

5'...CC¹TNAGC...3' 3'...GGANT₁CG...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCl (pH 7.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with *Bmr*FI, none of the DNA fragments can be ligated.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 30u of *Bmr*FI for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

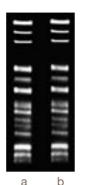
Bmr

* Blocked by overlapping dcm-methylation (OMCWGG): CCWGG

Ordering Information

Catalog No	Pack Size	
RV1154	400u	





λDNA (dam & dcm) 1% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	75%	100%	50%	75%	

*



Concentration 10-20u/ul

5'...GCTAG[↓]C...3' 3'...C_↑GATCG...5'

Reaction Conditions

10mM Tris-HCl (pH 7.5 at 30°C), 100mM MgCl $_2$, 50mM NaCl and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA. and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Bmt*l, about 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Bmt*116 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

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 $\lambda \text{DNA} \, (\textit{Hin} \text{dIII Digest}) \, 0.7\% \, \text{Agarose}$

a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RV1156	200u

Bmyl

Please refer to MhI (RE1292 - page 088)

Boxl

Please refer to Bst PA I (RE1234 – page 071)

Bpu101

Concentration 5u/µl

5'...CC¹TNAGC...3' 3'...GGANT₁CG...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCl (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *Bpu*10I, 80% of the DNA fragments can be ligated in the presence of 10% PEG and of these 90% can be recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 2u of *Bpu*10l for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1158	100u











λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 25% 25% 75% 100% 75%

Bpu14I {Asu II}

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Concentration 10u/µl

5'...TT[‡]CGAA...3' 3'...AAGC₁TT...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with Bpu14I, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Bpu14I for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

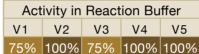




λDNA 0.7% Agarose

Ordering Information

Catalog No	Pack Size
RE1160	500u



BsaAl BsaBl BsaHl a - Digestion after 1 hour BsaMl b - Digestion after 16 hours

BpuMI {Cau II}

Concentration

20-50u/µl

5'...CC[‡]SGG...3' 3'...GGS₁CC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with BpuMI, less than 10% of the DNA fragments can be ligated.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of BpuMI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1162	500u











λDNA 1.0% Agarose

talog No Pack Size		Activity in		
		V1	V2	
V1162	500u	100%	100%	

a - Digestion after 1 hour b - Digestion after 16 hours

in Reaction Buffer V3 V4 V5 75% 100% 100%



Bse1 | {Bsr|}

Concentration

10-40u/µl

5'...ACTGGN¹...3' 3'...TGAC₁CN...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA.

Storage BufferThermal Inactivation

10mM Tris-HCl (pH 7.5), 100mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assav

After 10-fold overdigestion with Bse1 I. 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

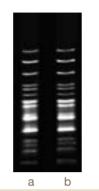
An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bse1 I for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1166	500u





λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
50%	75%	50%	25%	75%

Bse21 | {Saul}

Concentration

10-30u/ul

5'...CC[‡]TNAGG...3' 3'...GGANT_†CC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100ug/ml BSA. Incubate at 37°C.

Storage Condition

10mM KH₂PO₄ (pH 7.4), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assav

After 10-fold overdigestion with 211. 50% of the DNA fragments can be ligated by using high concentration of T4 DNA ligase and of these more than 90% can be recut.

Overdigestion Assay

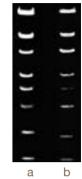
An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of 21l for 16 hours at 37°C.

Supplied with 10X Buffer V5. 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1170	400u





λDNA (Hind III Digest) 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer

V1 V2 V3 V4 V5 100% 100% 100% 75% 100%

Bse118| {Cfr10|}

Concentration

1-2u/µl

5'...R¹CCGGY...3' 3'...YGGCC_↑R...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM KH₂PO₄ (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

* Enzyme is stable for up to 6 months if properly stored. It is recommended that the enzyme is kept in small aliquots to avoid repeated freeze-thaw cycles.

Thermal Inactivation

None

Ligation / Recutting Assay

After 2-fold overdigestion with Bse118I, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 2u of Bse118l for 16 hours at 65°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1168	100u









λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
50%	100%	100%	50%	25%	

Bse3D | {BsrD|}

Concentration

1-10u/µl

5'...GCAATGNN¹...3' 3'...CGTTANC_↑NN...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with Bse3D I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 2u of Bse3D I. 16 hours at 60°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size	
RE1172	150u	



a b λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 50% 100% 75% 75% 100%



Bse8I {BsaBI}

Concentration 5-20u/µl

5'...GATNN¹NNATC...3' 3'...CTANN+NNTAG...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with Bse81, 80% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bse8l for 16 hours at 60°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1174	400u



λDNA 1.0% Agarose

a - Digestion after 1-fold b - Digestion after 5-fold

Activity in Poaction Buffor

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	100%	75%	75%	75%

BseX31

Concentration

1-5u/µl

5'...C¹GGCCG...3' 3'...GCCGG+C...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 8.2), 100mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C. Store at -70°C for period longer than 90 days.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with BseX3I, 90% of the DNA fragments can be ligated and of these 80% can be recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BseX3I for 16 hours at 50°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* Enzyme is stable for up to 6 months if properly stored. It is recommended that the enzyme is kept in small ali quots to avoid repeated freezethawed cycles.

Ordering Information

Catalog No	Pack Size	
RE1178	100u	





λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
50%	75%	100%	75%	50%

Please refer to Bst 2UI (RE1202 - page 063)

BsePl

Concentration

1-4u/µl

5'...G¹CGCGC...3' 3'...CGCGC₁G...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assav

After 4-fold overdigestion with BsePI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 8u of BsePI for 16 hours at 50°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)







λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1176	100u



BshVI {Clal}

Concentration 2-20u/µl

5'...AT¹CGAT...3' 3'...TAGC₁TA...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 2-fold overdigestion with BshVI, 90% of the DNA fragments can be ligated and recut.

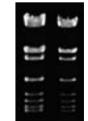
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BshVI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1180	400u



a b λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 25% 50% 75% 100% 100%



<i>Bsi</i> El	Please refer to Bst MCI (RE1230 – page 070)
<i>Bsi</i> HKAI	Please refer to Bbv12l (RE1140 – page 045)
Bsll	Please refer to Afi (RV1116 – page 037)
Bsml	Please refer to Pct I (RE1310 – page 093)

Please refer to BstMAI (RE1226 - page 069)

Bsnl {Haelll}

Concentration

BsmAl

2-10u/µl

5'...GG[‡]CC...3' 3'...CC_↑GG...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCl (pH 8.5 at 30°C), 10mM MgCl $_2$, 100mM KCl, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 0.15% Triton X-100, 500µg/ml BSA and 50% glycerol.Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 6-fold overdigestion with *Bsn*I, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *BsnI* for 16 hours at 37°C.

Diluent

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No		Pack Size	
	RV1182	1000u	

V4_{Bff}







λDNA 1.4% Agarose

a - Digestion after 1 hourb - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	75%	10%	100%	100%

Bso31| {Eco31|}

Concentration

2-5u/µl

5'...GGTCTC(N)₁¹...3' 3'...CCAGAG(N)₅†...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl $_{\rm 2}$, 100mM NaCl, and 100 μ g/ml BSA. Incubate at 55°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM KCI, 1mM EDTA, 7mM 2-mercaptoethanol, 100μg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 3-fold overdigestion with *Bso*31I, 90% of the DNA fragments can be ligated and of theses 80% can be recut.

Overdigestion Assay

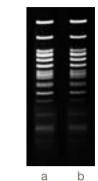
An unaltered banding pattern was observed after 1µg of DNA was digested with 2u of *Bso*31l for 16 hours at 55°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1184	100u





T7 DNA 1.0% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Activity in Reaction Buffer					
V1	V1 V2 V3 V4				
75%	100%	100%	100%	75%	

Bsp13| {BspMII}

Concentration

10-30u/ul

5'...T¹CCGGA...3' 3'...AGGCC₁T...5'

Reaction Conditions

1X Buffer Bsp13I

10mM Tris-HCl (pH7.6), 10mM MgCl2, 200mM KCl and 100μg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with *Bsp*13I, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Bsp*13l for 16 hours at 50°C.

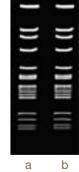
Supplied with 10X Buffer *Bsp*13I, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (G^mATC): TCCGGATC and GATCCGGA

Ordering Information

Catalog No	Pack Size
RE1186	500u





λDNA (dam & dcm) 0.7% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
25%	50%	75%	75%	75%		



Please refer to BstH2I (RE1222 - page 068)

Bsp17201 {Espl}

Concentration

1-10u/µl

5'...GC¹TNAGC...3' 3'...CGANT_†CG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with Bsp1720I, 80% of the DNA fragments can be ligated and of these 95% can be recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bsp1720I for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1188	150u





λDNA 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer						
V1	V1 V2		V4	V5		
50%	75%	50%	75%	100%		

Bsp19| {Ncol}

Concentration

10-30u/µl

5'...C¹CATGG...3' 3'...GGTAC₁C...5'

Reaction Conditions

1X Buffer Bsp191 20mM Tris-HCI (pH8.5), 10mM MgCl₂, 200mM NaCl and 1mM DTT. Incubate at 37°C.

Storage Condition

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with Bsp19I, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bsp19I for for 16 hours at 37°C.

Supplied with 10X Buffer Bsp19I, 10X Buffer UB and Viva Buffer A. (Diluent)

Catalog No Pack Si RE1190 300u









λDNA 0.7% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Ordering Information

ize	Activity in Reaction Buffer					
	V1	V2	V3	V4	٧	
	25%	50%	75%	75%	50	

BspEl

Please refer to Bsp13I (RE1186 - page 058)

Bsrl

Please refer to Bse11 (RE1166 - page 053)

BsrBl

Please refer to AccBSI (RE1110 - page 035)

BsrDI

Please refer to Bse3DI (RE1172 – page 054)

BsGl

Please refer to BstAUI (RE1208 - page 064)

BssHI

Please refer to BsePI (RE1176 – page 055)

BssMI (Mbol)

Concentration

1- 10u/µl

5'...¹GATC...3' 3'...CTAG_↑...5'

Reaction Conditions

2X Buffer UB

50mM Tris-acetate (pH7.6 at 30°C), 20mM Mg-acetate, 14mM 2-mercaptoethanol, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH7.5), 200mM KCI, 0.1mM EDTA, 1mM DTT, 14mM 2-mercaptoethanol, and 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 2-fold overdigestion with BssMI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

Ordering Information

Catalog No

RV1192

An unaltered banding pattern was observed after 1µg of DNA was digested with 4u of BssMI for for 16 hours at 37°C.

Supplied with 10X Buffer UB and VIva Buffer A. (Diluent)

a b

λDNA 1.0% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Pack Size	Activity in Reaction Buffe				
100u	V1	V2	V3	V4	V5
	25%	75%	75%	50%	75%



BssNI {Acyl}

Concentration

20-40u/µl

5'...GR¹CGYC...3' 3'...CYGC₁RG...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCl (pH8.5 at 30°C), 10mM MgCl₂, 100mM KCl and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with BssNI, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BssNI for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)



а λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RV1194	400u

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
50%	75%	50%	100%	100%	

BssT1I {Styl}

Concentration 20-40u/µl

5'...C¹CWWGG...3' 3'...GGWWC₁C...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH 7.5 at 30°C),

10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with BssT1 I, more than 95% of the DNA fragments can be ligated and

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BssT1I for 16 hours at 60°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1198	1000u









λDNA 1% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5

25% 75% 100% 100% 75%

Bst 11071

Please refer to BssNAI (RE1196 – page 061)

BssNAI {Snal}

Concentration

1-10u/µl

5'...GTA¹TAC...3' 3'...CAT₁ATG...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with BssNAI, more than 90% of the DNA pBR322 can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BssNAI for for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1196	200u



λDNA 0.7% Agarose a - Digestion after 1 hour

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
75%	75%	100%	100%	100%	



Bst2UI {EcoRII*}

Vivantis Technologies . Product Catalog Volume IV

Concentration

10-20u/ul

5'...CC[‡]WGG...3' 3'...GGW₁CC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH7.9 at 30°C, 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assav

After 2-fold overdigestion with Bst2UI, none of the DNA fragments can be

Overdigestion Assay

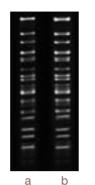
An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bst2UI for 16 hours at 60°C.

Supplied with 10X Buffer V5. 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1202	1000u





λDNA 1.0% Agarose

O
a - Digestion after 1 hour
b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	٧3	V4	V5
10%	50%	50%	100%	100%

Bst61 {Ksp6321}

Concentration

1-5u/µl

5'...CTCTTC(N)₁¹...3' 3'...GAGAAG(N)41...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C. Store at -70°C for period longer than 30 days.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 2-fold overdigestion with Bst61, 90% of the DNA fragments can be ligated and of these 80% can be recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 1u of Bst6l for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1206	100u











λDNA 1% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
100%	75%	75%	75%	100%

Bst981

Please refer to Vha464I (RE1358 - page 107)

Bst4Cl {Tsp4Cl}

Concentration 10 u/µl

5'...ACN[‡]GT...3' 3'...TG₁NCA...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100ug/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assav

After 5-fold overdigestion with Bst4Cl, less than 10% of the DNA fragments can be ligated.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Bst4Cl for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1204	150u



λDNA 1.4% Agarose a - Digestion after 1 hour

Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
00%	100%	50%	75%	100%		

b - Digestion after 16 hours

BstAUI {Bsp1407I}

Concentration

10-20u/µl

5'...T¹GTACA...3' 3'...ACATG_↑T...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 10mM NaCl, and 100µg/ml BSA. Store at -37°C.

Storage Buffer

50mM Tris-HCI (pH 7.5), 100mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with Bst AUI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 1u of BstAUI for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1208	200u



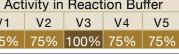






λDNA 0.7% Agarose a - Digestion after 1 hour

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	75%	100%	75%	75%	





BstBAI {BsaAI}

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Concentration

5-20u/µl

5'...YAC¹GTR...3' 3'...RTG_↑CAY...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCI, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with BstBAI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of *Bst*BAI for 16 hours at 65°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)



a b λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size	
RE1210	400u	

Activity in Reaction Buffer V1 V2 V3 V4 V5 50% 50% 100% 100% 50%

BstDSI {Dsal}

Concentration

1-10 u/µl

5'...C¹CRYGG...3' 3'...GGYRC_↑C...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with BstDSI, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

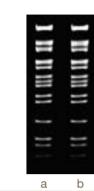
An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Bst*DSI for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1214	500u





λDNA 1.2% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
25%	75%	100%	75%	100%		

BstEll

Please refer to PspEI (RE1316 - page 095)

Bst DEI {Ddel}

Concentration

10-50u/μl

5'...C¹TNAG...3' 3'...GANT₁C...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH 7.5 at 30°C).

10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

10mM KH₂PO₄ (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200μg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with *Bst*DEI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of *Bst*DEI for 16 hours at 60°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

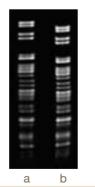
Ordering Information

Catalog No	Pack Size
RE1212	500u









 $\lambda \, a \text{ - Digestion after 1 hour} \\ \text{ b - Digestion after 16 hours}$

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 100% 50% 75% 50%

BstENI {EcoNI}

Concentration

2-5u/µl

5'...CCTNN¹NNNAGG...3' 3'...GGANNN₁NNTCC...5'

Reaction Conditions

1X Buffer UB

25mM Tris-acetate (pH 7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol, and 50μg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with BstENI, 60% of the DNA fragments can be ligated and of these 90% can be recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 3u of *Bst*ENI for 16 hours at 65°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1216	100u



a b λDNA 1.4% Agarose

a - Digestion after 1 hour

Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
75%	75%	75%	75%	75%		



BstF5 | {Fok|*}

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Concentration

1-10u/µl

5'...GGATGNN¹...3' 3'...CCTAC₁NN...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with BstF5I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

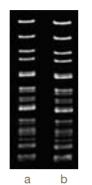
An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of BstF5I for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1218	200u





λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	75%	100%	100%	100%	

BstH2| {Hae||}

Concentration

10-30u/ul

5'...RGCGC¹Y...3' 3'...Y_↑CGCGR...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM KH₂PO₄ (pH 7.2), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with BstH2I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BstH2I for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1222	600u











λDNA 1.0% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
25%	50%	50%	75%	100%		

BstFNI {FnuDII}

Concentration 2-10 u/µl

5'...CG¹CG...3' 3'...GC₁GC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

20mM Tris-HCI (pH 7.5), 300mM NaCl, 10mM MgCl₂ 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with BstFNI, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BstFNI for 16 hours at 60°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1220	100u









λDNA 1.4% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer						
V1	V2	٧3	V4	V5		
75%	75%	50%	75%	100%		

BstHHI {Hhal}

Concentration

50u/µl

5'...GCG¹C...3' 3'...C_†GCG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 50-fold overdigestion with BstHHI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 100u of BstHHI for 16 hours at 50°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

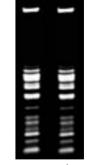
Ordering Information

Catalog No	Pack Size
RE1224	1000u









λDNA 1.2% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
100%	75%	50%	75%	100%



BstMAI {BsmAI}

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Concentration

30-100u/µl

5'...GTCTC(N)1[↓]...3' 3'...CAGAG(N)5_↑...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCI, and 100µg/ml BSA. Incubate at 55°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 30-fold overdigestion with BstMAI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 30u of *Bst*MAI for 16 hours at 55°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1226	1000u





λDNA 1% Agarose

a - Digestion after 1 hourb - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	75%	75%	100%	100%

BstMCI {Mcrl}

Concentration

5u/µl

5'...CGRY¹CG...3' 3'...GC₁YRGC...5'

Reaction Conditions

1X Buffer V1 10mM Tris-HCl (pH 7.5 at 30°C), 10mM MgCl₂, and 100μg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with BstMCI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 2.5u of *Bst*MCI for 16 hours at 50°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size	
RE1230	200u	









a b

λDNA 1% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
100%	100%	75%	50%	75%	

BstMBI {Mbol}

Concentration

5-20u/µl

5'...¹GATC...3' 3'...CTAG₁...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCI, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with BstMBI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Bst*MBI for 16 hours at 65°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (GMATC): GATC

Ordering Information

Catalog No	Pack Size
RE1228	200u

V4_{Bff}









a b λDNA (dam-&dcm-) 1.4% Agarose

Activity in Reaction Buffer
V1 V2 V3 V4 V5
50% 100% 75% 100% 100%

a - Digestion after 1 hour

b - Digestion after 16 hours

BstNl

Please refer to Bst2UI (RE1202 - page 063)

BstNSI {Nspl}

Concentration

1-10u/ul

5'...RCATG¹Y...3' 3'...Y_↑GTACR...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with BstNSI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of *Bst*NSI for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

T7 DNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Catalog No	Pack Size
RE1232	200u

BstPAI {PshAI}

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Concentration

20-50u/µl

5'...GACNN¹NNGTC...3' 3'...CTGNN+NNCAG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C). 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with BstPA I. more than 5% of the DNA fragments can be ligated.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of Bsp PAI for 16 hours at 65°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

* Prolong Incubation will result in Star Activity.

Ordering Information

Catalog No	Pack Size	
RE1234	500u	





λDNA 1% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Act	ivity in	Reaction Buffer		
V1	V2	V3	V4	V5
750/	1000/	750/	750/	1000/



BstSNI {SnaBI}

Concentration

5-15u/µl

5'...TAC¹GTA...3' 3'...ATG_↑CAT...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 5-fold overdigestion with BstXI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of BstSNI for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1236	200u









T7 DNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
100%	100%	50%	75%	100%	

BstUI

Please refer to BstFNI (RE1220 - page 067)

BstV2I {Bbvll}

Concentration

5-15u/ul

5'...GAAGAC(N)₂¹...3' 3'...CTTCTG(N)6+...5'

Reaction Conditions

1 X Buffer UB

25mM Tris-acetate (pH7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol and 25µg/ml BSA. Incubate at 55°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with BstV21, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of BstV2I for 16 hours at 55°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in

Ordering Information

Catalog No	Pack Size
RE1238	200u



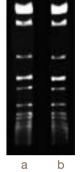












λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Act	ivity in	React	ion Bu	ffer
V1	V2	V3	V4	V5
50%	75%	50%	75%	75%

BstXI

Concentration

5-15u/µl

5'...CCANNNNN¹NTGG...3' 3'...GGTN+NNNNNACC...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100μg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with BstXI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of BstXI for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1240	200u







λDNA 0.7% Agarose a - Digestion after 1 hour

	Activity in Reaction Buffer				
V1 V2 V3 V4					
	50%	50%	100%	50%	25%



BstX2I {Xholl}

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Concentration

3-5u/µl

5'...R¹GATCY...3' 3'...YCTAG₁R...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 60°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Thermal Inactivation

Bsu Rl

None

Ligation / Recutting Assay

After 5-fold overdigestion with BstX21, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 6u of BstX2I for 16 hours at 60°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

а

λ DNA 0.7% Agarose

100% 100% 75% 75% 100%

- a Digestion after 1 hour
- b Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1242	100u

Please refer to BsnI (RV1182 - page 057)

Activity in Reaction Buffer V1 V2 V3 V4 V5

BtuMI {Nrul}

Concentration

5-10u/ul

5'...TCG¹CGA...3' 3'...AGC₁GCT...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM observed after 1µg of DNA was MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, (Diluent) and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with BtuM I, about 50% of the DNA fragments can be ligated and of these 90% can be recut.

Overdigestion Assay

An unaltered banding pattern was digested with 5u of BtuMI for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A.

* Blocked by overlapping dam-methylation (G^mATC): GATCGCGA, (TOGCGATC).

Ordering Information

Catalog No	Pack Size
RV1246	200u





Restriction Endonucleases





λDNA (dam-&dcm-) 0.7% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Activity in Reaction Buffer

7 totivity iii i toaotion banoi				
V1	V2	V3	V4	V5
50%	100%	100%	100%	75%

CciNI {Notl}

Concentration

2-5u/ul

5'...GC[†]GGCCGC...3' 3'...CGCCGG_↑CG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with NI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of NI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

- * High enzyme concentration may result in Star Activity.
- * Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1248	100u



Ad2 DNA 0.7% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V					
50%	75%	75%	75%	100%	



Please refer to Bsp1720 I (RE1188 – page 059)

Cfr101 Please refer to Bse 118 I (RE1168 – page 053)

Cfr131 Please refer to AspS9 I (RE1132 - page 042)

Cfr421 Please refer to Sfr**303 I** (RE1334 – page 102)

Please refer to BshVI (RV1180 - page 056)

Ddel Please refer to Bst DEI (RE1212 - page 065)

Dinl {Narl}

Concentration

1-20u/ul

5'...GGC[†]GCC...3'

3'...CCG₁CGG...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM KCI, 0.1mM EDTA, 1mM DTT, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 6-fold overdigestion with Dinl, 70% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Dinl for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No Pack Size RV1252 200u



a b λDNA (Hind III Digest) 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
75%	75%	50%	100%	100%	

Dral {Ahalll}

Concentration

10-40u/µl

5'...TTT¹AAA...3' 3'...AAA_↑TTT...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with Dral, 50% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 80u of *Dral* for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1254	600u





a b

λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
75%	100%	75%	50%	75%	

Dra III

Concentration

3-10u/µl

5'...CACNNN¹GTG...3' 3'...GTG_↑NNNCAC...5'

Reaction Conditions

1X Buffer Dralll

10mM Tris-HCI (pH 7.6), 10mM MgCl₂, 200mM KCl, 1mM DTT and 100μg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 200mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with Dra III. 70% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of Dralll for 16 hours at 37°C.

Supplied with 5X Buffer Dralll, 10X Buffer UB and Viva Buffer A

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1256	200u



λDNA 1.0% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 50% 50% 75% 75%





Please refer to BstDSI (RE1214 - page 066)

DseD | {Drd|}

Concentration

10-30u/µl

5'...GACNNNN¹NNGTC...3' 3'...CTGNN₁NNNNCAG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Condition

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with DseDI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of DseDI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1372	300u





λDNA 0.7% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V					
75%	50%	25%	50%	100%	

Eagl Please refer to BseX3I (RE1178 – page 056) Eam11051 Please refer to BmeRI (RV1150 - page 048) Please refer to Bst 61 (RE1206 - page 064) Eco311 Please refer to Bso31I (RE1184 - page 058)

EcolCRI {Sacl}

Concentration

2-10u/µl

5'...GAG[‡]CTC...3' 3'...CTC₁GAG...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl_a, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assav

After 5-fold overdigestion with Eco1CRI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assav

An unaltered banding pattern was observed after 1µg of DNA was digested with 1.5u of EcolCRI for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

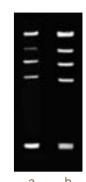
Ordering Information

Catalog No	Pack Size
RE1258	200u









λDNA (Hind III Digest) 1.0% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Catalog No	Pack Size
RE1258	200u

Activity in Reaction Buffer V1 V2 V3 V4 V5 100% 100% 50% 75% 75%

EcoN

Please refer to Bst ENI (RE1216 - page 066)

FcoRI

Concentration

20-100u/µl

5'...G[‡]AATTC...3' 3'...CTTAA₁G...5'

Reaction Conditions

1X Buffer V2

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl_a, 100mM NaCl, 0.02% Triton X-100, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM NaCl, 0.1mM EDTA, 200µg/ml BSA 1m/m DTT and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 50-fold overdigestion with EcoRI, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 50u of EcoRI for 16 hours at 37°C.

Supplied with 10X Buffer EcoRI, 10X Buffer UB and Viva Buffer A. (Diluent)

High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1260	2500u



λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 50% 50% 100% 100% 50%



EcoRII

Please refer to Bst2UI (RE1202 - page 063)

EcoRV



Concentration

 $20-40u/\mu I$

5'...GAT¹ATC...3' 3'...CTA₁TAG...5'

Reaction Conditions

1X Buffer EcoR V 10mM Tris-HCl (pH 8.5 at 30°C), 10mM MgCl $_2$, 100mM NaCl, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 50mM NaCl, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with R V, 80% of the DNA fragments can be ligated and recut.

Overdigestion Assay

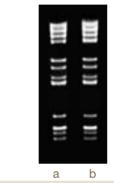
An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of RV for 16 hours at 37°C.

Supplied with 10X Buffer RV, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1262	1000u



λDNA 1.0% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1 V2 V3 V4 V5					
0%	0%	100%	75%	0%	

Ege

Please refer to Din1 (RV1252 - page 075)

FauNDI {Ndel}

Concentration

5-20 u/µl

5'...CA[‡]TATG...3' 3'...GTAT_†AC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *Fau*NDI, 60% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Fau*NDI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Sensitive to impurities present in some DNA preparation.

Ordering Information

Catalog No		Pack Size
	RE1266	500u





λDNA 0.7% Agarose

a - Digestion after 1 hour

	Activity in Reaction Buffer				
V1 V2 V3 V4 V5					
	50%	75%	50%	75%	100%

FbII {Accl}

Concentration

2-10u/ul

5'...GT \ MKAC ...3' 3'...CAKM+TG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 55°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None









а λDNA 1% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1268	150u

Catalog No	Pack Size
RE1268	150u

Ligation / Recutting Assav

ligated and recut.

hours at 55°C.

(Diluent)

Overdigestion Assay

After 3-fold overdigestion with Fb/I.

90% of the DNA fragments can be

An unaltered banding pattern was

observed after 1µg of DNA was

Supplied with 10X Buffer V5, 10X

Buffer UB and Viva Buffer A.

digested with 4u of Fb/I for 16

Activity in Reaction Buffer				
V1	V2	٧3	V4	V5
100%	100%	50%	75%	100%









Fokl

Concentration

1-3u/µl

5'...GGATG(N)9¹...3' 3'...CCTAC(N)_{13↑}...5'

Reaction Conditions

1X Buffer V2 50mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, 100mM NaCl, and 100μg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 2-fold overdigestion with Fokl. more than 95% of the DNA fragments can be ligated and recut.

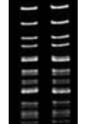
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Fok I for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1270	150u



a b λDNA 1.0% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Act	ivity in	React	ion Bu	ffer
V1	V2	V3	V4	V5
50%	100%	25%	50%	50%

FriOI {HgiJII}

Concentration

10-40u/µl

5'...GRGCY¹C...3' 3'...C_†YCGRG...5

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

*Fs*PI

Ligation / Recutting Assay

After 10-fold overdigestion with FriOI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of FriOI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Enzyme is stable for up to 6 months if properly stored. It is recommended that the enzyme is kept in small aliquots to avoid repeated freeze-thaw cycles.

Ordering Information

Catalog No	Pack Size
RE1272	800u







λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 50% 100% 50% 50% 100%

Please refer to Acc16 I (RE1102 - page 033)

Haell Please refer to Bst H2I (RE1222 - page 068)

Haelll Please refer to Bsn I (RV1182 – page 057)

Hhal Please refer to AspLE I (RE1130 – page 042)

Hhal Please refer to BstHH I (RE1224 – page 068)

Hin11 Please refer to BssN I (RV1194 - page 061)

Hincl Please refer to Hind II (RE1274 – page 083)

Hind II

Concentration

5-20u/µl

5'...GTY¹RAC...3' 3'...CAR_†YTG...5'

Reaction Conditions

0.5X Buffer UB

12.5mM Tris-acetate (pH 7.6 at 30°C), 5mM Mg-acetate, 50mM k-acetate, 3.5mM 2-mercaptoethanol, 25µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with HindII, more than 60% of the DNA fragments can be ligated and recut. In the presence of 10% PEG, ligation is better.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Hindll for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

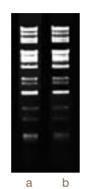
Catalog No	Pack Size
RE1274	400u











λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
100%	100%	75%	100%	100%

Hinfl {Ndel}

Concentration

10-30u/µl

5'...G[‡]ANTC...3' 3'...CTNA₁G...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM NaCl, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and 50% glycerol.Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with Hinfl, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assav

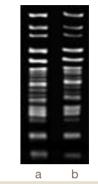
An unaltered banding pattern was observed after 1µg of DNA was digested with 30u of Hinfl for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

a - Digestion after 1 hour

Ordering Information

Catalog No	Pack Size
RE1278	1000u



λDNA 1.4% Agarose

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5

50% 100% 100% 100% 50%

Hind III

Concentration

30-100u/µl

5'...A[‡]AGCTT...3' 3'...TTCGA₁A...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 300mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assav

After 40-fold overdigestion with HindIII. 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of HindIII for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1276	2500u









а λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	100%	75%	75%	75%

Hpal

Concentration

5-10u/µl

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM

Thermal Inactivation

65°C for 20 minutes









5'...GTT¹AAC...3' 3'...CAA₁TTG...5'

Reaction Conditions

1X Buffer V5

NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Ligation / Recutting Assay

After 5-fold overdigestion with Hpal, more than 60% of the DNA fragments can be ligated and recut.

Overdigestion Assay

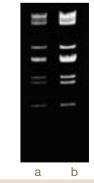
Star Activity is observed at greater than 5-fold over digestion of 1µg substrate with Hpal.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1280	200u



λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
75%	75%	50%	75%	100%		



Hpall

Concentration 5-10u/µl

5'...C[†]CGG...3' 3'...GGC tC...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂ and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM NaCl, 0.1mM EDTA, 1mM DTT, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assav

After 5-fold overdigestion with Hpall, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assav

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Hpall for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation.

Ordering Information

Catalog No	Pack Size
RE1282	500u



λDNA 1.4% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
100%	100%	75%	75%	100%		

Hphl

Please refer to AsuHPI (RE1134 – page 043)

HspA | {Hhal*}

Concentration 10-30u/µl

5'...G¹CGC...3' 3'...CGC₁G...5'

Reaction Conditions

0.5X Buffer UB

12.5mM Tris-acetate (pH 7.6 at 30°C), 5mM Mg-acetate, 50mM K-acetate, 3.5mM 2-mercaptoethanol, and 25µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with HspAI, more than 90% of the DNA fragments can be ligated recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Hsp AI for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation.

Ordering Information

Catalog No	Pack Size
RE1284	500u







λDNA 1.0% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% | 75% | 75% | 75% | 75%

Kpnl

Concentration

10-20u/µl

5'...GGTAC¹C...3' 3'...C₁CATGG...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C). 10mM MgCl₂, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with KpnI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Kpnl for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1286	1000u











λDNA 1% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 100% 25% 25% 25% 75%

Ksp22I {BcII}

Concentration 10-30u/µl

5'...T¹GATCA...3' 3'...ACTAG_↑T...5'

Reaction Conditions

1X Buffer V5

30mM Tris-HCI (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM KH₂PO₄ (pH 7.2), 200mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with Ksp22I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Ksp22l for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (GMATC): TGATCA

Ordering Information

Catalog No	Pack Size
RE1288	1000u









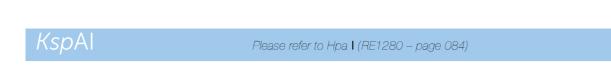


λDNA (dam⁻& dcm⁻) 1.0% Agarose

a - Digestion after 1 hour

Activity in Reaction Buffer				
V1 V2		V3	V4	V5
75%	100%	75%	75%	100%





Kzo91 Please refer to BssM I (RV1192 - page 060)

Kzo91 Please refer to BstMBI (RE1228 - page 069)

Maml Please refer to Bse 8 I (RE1174 – page 055)

Mbil Please refer to Acc B SI (RE1110 - page 035)

Mbol Please refer to BssM I (RV1192 - page 060)

Mhll (Sdul)

Concentration

5-10u/µl

5'...GDGCH¹C...3' 3'...C+HCGDG...5'

Reaction Conditions

1.5X Buffer UB

37.5mM Tris-acetate (pH 7.6 at 30°C). 15mM Mg-acetate. 150mM K-acetate, 10.5 mM 2-mercaptoethanol and 75µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with Mhll, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of Mh/I for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A

* High enzyme concentration may result in Star Activity.

Ordering Information

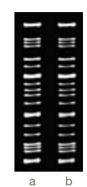
Catalog No	Pack Size
RE1292	200u











λDNA 1.0% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						
25%	50%	50%	25%	75%		

Please refer to Msp20l (RE1304 – page 091)

Mlul

Concentration

10-20u/µl

5'...A[‡]CGCGT...3' 3'...TGCGC₁A...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation 65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with Mlul, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Mlul for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size	
RE1294	500u	









a b

λDNA 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5

25% 75% 100% 75% 50%

Mholl

Concentration $0.5 - 3u/\mu I$

5'...GAAGA(N)₈¹...3' 3'...CTTCT (N)71...5'

Reaction Conditions

1X Buffer Mboll

33mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 66mM K-acetate, and 1mM DTT. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation 65°C for 20 minutes



After 3-fold overdigestion with Mbo II,

about 60% of the DNA fragments can

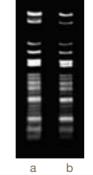
Ligation / Recutting Assay

observed after 1µg of DNA was digested with 3u of Mbo II for 16 hours at 37°C.

Supplied with 10X Buffer Mbo II, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (G^MATC): GAAGATC

Orderi



λDNA (dam-&dcm-) 1.4% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

ring Informa	ation		
		٥.	Δctivit

Catalog No	Pack Size
RE1290	100u

Mnll

Concentration

2-10u/µl

5'...CCTC(N)₆¹...3' 3'...GGAG(N)71...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with Mn/I. 50% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1ug of DNA was digested with 20u of Mn/I for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CG methylation: CCTmCG

λDNA 1% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Catalog No	Pack Size	
RE1296	150u	

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 100% 50% 50% 75%

Mrol

Please refer to Bsp13I (RE1186 - page 058)

MroNI {Nael*}

Concentration

2-10u/µl

5'...G¹CCGGC...3' 3'...CGGCC₁G...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with NI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1ug of DNA was digested with 10u of NI for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

λDNA (Hind III Digest) 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size	
RE1298	150u	

Activity in Reaction Buffer V2 V3 V4 100% 50% 10% 10% 10%

MroXI {Xmn|}

Concentration

5-15u/ul

5'...GAANN¹NNTTC...3' 3'...CTTNN+NNAAG...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with MroXI, 50% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of MroXI for 16 hours at 37°C.

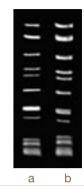
Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size	
RE1300	300u	







λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Catalog No	Pack Size	
RE1300	300u	

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 100% 100% 100% 75%

Msel

Please refer to Tru9I (RE1350 - page 106)

Mspl {Hpall}

Concentration

10-40u/µl

5'...C¹CGG...3' 3'...GGC₁C...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of I for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size	
RE1302	1000u	







λDNA 1.2% Agarose a - Digestion after 1 hour

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	100%	75%	75%	75%



Msp201 {Ball}

Concentration

1-3u/µl

5'...TGG CCA...3' 3'...ACCGGT...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCI. 0.1mM EDTA. 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes









λDNA (dam-&dcm-) 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

* Blocked by overlapping dcm⁻methylation (O^mCWGG): TGGCCAGG

After 3-fold overdigestion with 201, 80% of the DNA fragments can be

An unaltered banding pattern was

digested with 2u of 20l for 16 hours

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A.

observed after 1µg of DNA was

Ordering Information

ligated and recut.

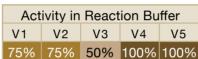
at 37°C.

(Diluent)

Overdigestion Assay

Catalog No	Pack Size		
RE1304	50u		

Pack Size	
50u	













λDNA 0.7% Agarose a - Digestion after 1 hour

Reaction Conditions 1X Buffer V5

5'...CMG[‡]CKG...3'

3'...GKC₁GMC...5'

Concentration

2-10u/µl

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

MspA1I {NspBII}

Storage Buffer

20mM Tris-HCI (pH 7.6), 300mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 10mM MgCl₃ 200µg/ml BSA and 50% glycerol. Store at -20°C. Store at -70°C for period longer than 30 days.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with MspA1I, 60% of the DNA fragments can be ligated and recut.

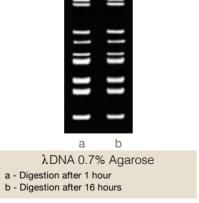
Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of MspA1I for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size	
RE1306	200u	



Activity in Reaction Buffer				
V1	V2	٧3	V4	V5
100%	100%	75%	100%	100%

Please refer to MroNI (RE1298 - page 089) NaRI* Please refer to Din1 (RV1252 - page 075) Ncil Please refer to BpuMI (RV1162 – page 051) Ncol Please refer to Bsp19I (RE1190 - page 059) Ndel Please refer to FauNDI (RE1266 - page 080) Please refer to AsuNHI (RE1136 - page 043) Please refer to Bmil (RV1152 - page 048) Notl Please refer to CciNI (RE1248 - page 074) Nrul Please refer to BtuMI (RV1246 – page 074) Nsil Please refer to Zsp2I (RE1370 - page 111) Nsp Please refer to Bst NSI (RE1232 - page 070) PaeR71 Please refer to Sfr274I (RE1332 - page 102)

Pce | {Stul}

Concentration 10-20u/µl

5'...AGG[‡]CCT...3' 3'...TCC_↑GGA...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with I, 70% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of I for 16 hours at 50°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)





λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1308	500u

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 75% 75% 100% 100%

PflMI Please refer to Acc B7I (RE1108 – page 035) PinAI Please refer to Asi GI (RE1126 – page 041) PmII Please refer to PspCI (RE1314 – page 095) PshAI Please refer to Bst PAI (RE1234 – page 071)

Please refer to Tth 1111 (RE1356 - page 107)

Psp124BI {SacI}

Concentration

20-50u/µl

5'...GAGCT¹C...3' 3'...C_↑TCGAG...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100μg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation 65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Psp*124BI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 80u of *Psp*124BI for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1312	500u



a b

λDNA (Hind III Digest) 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Act	ivity in	React	ion Bu	ffer
V1	V2	V3	V4	V5
100%	100%	75%	100%	100%

Pct l {Bsml}

Concentration 10-40 u/µl

5'...GAATGCN¹...3' 3'...CTTAC₁GN...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCl (pH 7.5 at 30°C), 10mM MgCl $_2$, 100mM NaCl, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with *Pct*I, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 30u of *Pct*I for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size	
RE1310	400u	



λDNA 1.4% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

50% 75% 100% 75% 75%

talog No Pack Size Activity in Reaction Buffer V1 V2 V3 V4 V5



PspCl {PmaCl}

Concentration

10-30u/µl

5'...CAC[‡]GTG...3' 3'...GTG+CAC...5'

Reaction Conditions

1X Buffer V1 10mM Tris-HCI (pH 7.5 at 30°C). 10mM MaCl₂, and 100ua/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C. Store at -70°C for period longer than 30 days.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with PspCI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assav

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of PspCl for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1314	500u





λDNA 0.7% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer							
V1	V2	V3	V4	V5			

100% 75% 25% 50% 100%

PspOMI {Apal*}

Concentration

10-30u/µl

Reaction Conditions

5'...G¹GGCCC...3' 3'...CCCGG+G...5'

Storage Buffer

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, and 100µg/ml BSA. Incubate at 37°C.

Thermal Inactivation

10mM Tris-HCI (pH 7.5), 100mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C. Ligation / Recutting Assay

Overdigestion Assay

After 10-fold overdigestion with PspCI, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of PspOMI for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1318	800u







λDNA (BamHI Digest) 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Catalog No	Pack Size
RE1318	800u

Activity in Reaction Buffer V2 V3 V4 V5 100% 25% 10% 25% 10%

PspEl {BstEll}

Concentration

5-10u/µl

5'...G¹GTNACC...3' 3'...CCANTG₁G...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C). 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5). 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with PspEI, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of PspEI for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No		Pack Size
RE131	6	500u









λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer						
V1	V2	V3	V4	V5		
75%	75%	75%	75%	100%		

Pstl

Concentration 20-100u/ul

5'...CTGCA[‡]G...3' 3'...G_↑ACGTC...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 30-fold overdigestion with Pstl, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Pstl for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1320	2000u



λa - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
25%	50%	100%	50%	50%	



Pvull

Concentration 20-50u/µl

5'...CAG[‡]CTG...3' 3'...GTC_†GAC...5'

Reaction Conditions

1X Buffer V2 10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, and 50% glycerol. Store at -20°C.

Thermal Inactivation

None











Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of Pvull for 16 hours at 37°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1322	800u



λDNA 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
75%	100%	50%	25%	50%	

Rsr2l {Rsrll}

Concentration

10-30u/µl

5'...CG[‡]GWCCG...3' 3'...GCCWG_†GC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl(pH 7.5), 50mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation 65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with Rsr2I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 60u of *Rsr*2l for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1374	300u









λDNA 0.7% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
75%	75%	25%	25%	100%

Rsal

Concentration

10-30u/µl

5'...GT[‡]AC...3' 3'...CA_↑TG...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCl (pH 7.5 at 30°C), 10mM MgCl₂, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

20mM Tris-HCl (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 20-fold overdigestion with Rsal, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of *Rsa*l for 16 hours at 37°C

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

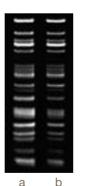
Catalog No Pack Size

RE1324 500u









λDNA 1.4% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

g information

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
100%	100%	50%	50%	100%

Sacl

Please refer to Psp124BI (RE1312 - page 094)

SacII

Please refer to Sfr303I (RE1334 - page 102)



Sall

Vivantis Technologies . Product Catalog Volume IV

Concentration 10-20 u/µl

5'...G[‡]TCGAC...3' 3'...CAGCT₁G...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes











Ligation / Recutting Assay

After 10-fold overdigestion with Sall, 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 15u of I for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

- * High enzyme concentration may result in
- * Blocked by CpG methylation



λDNA (Hind III Digest) 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RV1326	1000u

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
0%	75%	100%	100%	100%	

Sau3Al

Please refer to BssMI (RV1192 - page 060)

Sau3Al

Please refer to Bst MBI (RE1228 – page 069)

Sau961

Please refer to AspS9I (RE1132 - page 042)

Sbfl {Sse8387l}

Concentration

2-5u/µl

5'...CCTGCA[‡]GG...3' 3'...GG_↑ACGTCC...5'

Reaction Conditions

1X Buffer V5 30mM Tris-HCI (pH 7.9 at 30°C),

10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5). 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with Sbfl, more than 90% of the DNA fragments can be ligated and of these 90% can be recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of Sbfl for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1328	100u



a b

λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Scal

Please refer to BmcA I (RV1146 - page 047)

ScrFI

Please refer to BmrF I (RV1154 - page 049)

Sdal

Please refer to Sbf I (RE1328 - page 100)



SfaNI

Concentration

5-15u/µl

5'...GCATC(N)₅¹...3' 3'...CGTAG(N)₉₁...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 300mM NaCl, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assav

After 10-fold overdigestion with *SfaNI*, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 15u of *Sfa*NI for 16 hours at 37°C.

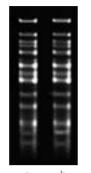
Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1376	200u

V3BH 37°C CG



λDNA 1.4% Agarose

a - Digestion after 1 hourb - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
50%	75%	100%	100%	75%	

Sfil

Concentration

10-40u/µl

5'...GGCCNNNN¹NGGCC...3' 3'...CCGGN₁NNNNCCGG...5'

Reaction Conditions

1X Buffer V2

10mM Tris-HCl (pH 7.5 at 30°C), 10mM MgCl₂, 50mM NaCl, and 100µg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.6), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Sfol

Ligation / Recutting Assav

After 10-fold overdigestion with *Sfil*, 70% of the DNA fragments can be ligated and recut.

Overdigestion Assay

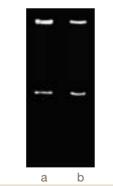
An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of *Sfil* for 16 hours at 50°C.

Supplied with 10X Buffer V2, 10X Buffer UB and Viva Buffer A. (Diluent)

- * Blocked by overlapping dcm-methylation (C^mCWGG): GGCCWGGNNGGCC
- * Not blocked by overlapping dam-methylation (C^mCWGG): GGCCNNNNNGGCCWGG.

Ordering Information

Catalog No	Pack Size
RE1330	500u



T7 DNA 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer					
V1	V2	V3	V4	V5	
100%	100%	10%	50%	75%	

Please refer to Din I (RV1252 - page 075)

*Sfr*274I {*XhoI*}

Concentration

10-40u/µl

5'...C¹TCGAG...3' 3'...GAGCT_↑C...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, and 100μg/ml BSA. Incubate at 50°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with *Sfr*274 I, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of *Sfr*274l for 16 hours at 50°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

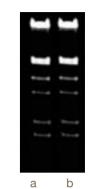
- * Blocked by overlapping dcm-methylation CTOG^mAG:
- * Not blocked by overlapping dam-methylation CT^mCGAG

V1_{Bff}









λDNA (Hind III Digest) 0.7% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1332	500u

Sfr303| {Sac||}

Concentration

5-20u/µl

5'...CCGC¹GG...3' 3'...GG₁CGCC...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCl (pH 7.5 at 30°C), 10mM MgCl $_2$, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *Sfr*303I, more than 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Sfr*303l for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1334	1000u







λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours



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Please refer to Bpu14I (RE1160 - page 051)

Smal

Concentration 10-20u/µl

5'...CCC[†]GGG...3' 3'...GGG+CCC...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 25°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes









recut.

Ligation / Recutting Assav

After 5-fold overdigestion with

fragments can be ligated and

Smal, more than 90% of the DNA

Overdigestion Assav An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Smal for 16 hours at 25°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by CpG methylation

Ordering Information

Catalog No	Pack Size
RE1336	1000u



λDNA (Hind III Digest) 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Act	ivity in	React	ion Bu	ffer
V1	V2	V3	V4	V5
25%	10%	10%	75%	100%



Smil (Swal)

Concentration 10-30u/µl

5'...ATTT[‡]AAAT...3' 3'...TAAA₁TTTA...5'

Reaction Conditions

1X Buffer V3 50mM Tris-HCI (pH 7.5 at 30°C).

10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl. 0.1mM EDTA. 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with Smil, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Smil for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1338	500u









T7 DNA (Sspl Digest) 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
50%	50%	100%	75%	25%

SmiMI {Msll}

Concentration

5-10u/µl

5'...CAYNN¹NNRTG...3' 3'...GTRNN+NNYAC...5'

Reaction Conditions

1X Buffer V4 10mM Tris-HCI (pH 8.5 at 30°C), 10mM MgCl₂, 100mM KCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 250mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with SmiMI, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of SmiMI for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)



λDNA 1.2% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1378	100u

Activity in Reaction Buffer

V1	V2	V3	V4	V5
50%	75%	100%	100%	100%

Spel

Please refer to Ahl (RE1118 – page 038)

Sphl

Concentration 3-10u/µl

5'...GCATG¹C...3' 3'...C_†GTACG...5'

Reaction Conditions

1X Buffer UB

25mM Tris-acetate (pH 7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol, and 50µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 3-fold overdigestion with Sphl, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assav

An unaltered banding pattern was observed after 1ug of DNA was digested with 6u of Sphl for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RV1340	200u









λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 75% 75% 50% 75% 75%



Sse9I {TspEI}

Concentration

2-10u/µl

5'...¹AATT...3' 3'...TTAA₁...5'

Reaction Conditions

1x Buffer V1

10mM Tris-HCl (pH7.5 at 30°C), 10mM MgCl $_{2}$, and 100µg/ml BSA. Incubate at 55°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 100µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *Sse*9I, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Sse*9I for 16 hours at 55°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1342	200u





pBR322 2% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

Act	ivity in	React	ion Bu	ffer
V1	V2	V3	V4	V5
100%	75%	25%	50%	75%

Concentration

3-30u/µl

Tagl

5'...T¹CGA...3' 3'...AGC_↑T...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

None

Ligation / Recutting Assay

After 10-fold overdigestion with *Taq*l, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

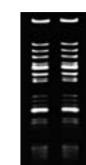
An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Taq*I for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (G^mATC): TCGATC.

Ordering Information

Catalog No	Pack Size
RE1346	2000u



a k

DNA (dam⁻& dcm⁻) 1.0% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Tasl

Concentration

1X Buffer V1

5'...T¹TAA...3'

3'...AAT₁T...5'

Incubate at 65°C.

Storage Buffer

Reaction Conditions

10mM Tris-HCI (pH 7.5 at 30%),

10mM MgCl₂, and 100µg/ml BSA.

10mM Tris-HCI (pH 7.5), 50mM KCI

0.1mM EDTA, 10mM DTT, 200µg/

ml BSA and 50% glycerol. Store at

Tru9| {Msel}

Please refer to Sse9 I (RE1342 - page 105)

Sspl

Concentration

5-20u/µl

5'...AAT[‡]ATT...3' 3'...TTA_†AAT...5'

Reaction Conditions

1X Buffer Sspl 10mM Tris-HCI (pH 7.6 at 30°C), 10mM MgCl₂, 100mM KCI, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCl (pH 7.5), 200mM KCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with Sspl, 90% of the DNA fragments can be ligated and recut

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of I for 16 hours at 37°C.

Supplied with 10X Buffer I, 10X Buffer UB and Viva Buffer A. (Diluent)

- * High enzyme concentration may result in Star Activity.
- * Blocked by A^mATATT methylation

Ordering Information

Catalog No	Pack Size
RE1344	200u



λDNA 1% Agarose

Activity in Reaction Buffer

V1 V2 V3 V4 V5

75% 75% 50% 75% 75%

- a Digestion after 1 hour
- b Digestion after 16 hours

Thermal Inactivation None

-20°C

Ligation / Recutting Assay

After 10-fold overdigestion with *Tru*9I, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u a Trust for 16 hours at 65°C.

Supplied with 10X Buffer I, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by TTA^mA methylation

Ordering Information

Catalo	g No	Pack Size
RE135	60	200u



a b λDNA 1.0% Agarose

-

a - Digestion after 1 hourb - Digestion after 16 hours

Activity in Reaction Buffer

V1 V2 V3 V4 V5

100% 75% 10% 50% 25%



Tth1111

Concentration

5-10u/µl

5'...GACN¹NNGTC...3' 3'...CTGNN₁NCAG...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 65°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, and 50% glycerol. Store at -20°C.

Thermal Inactivation

None



Ligation / Recutting Assay

After 2-fold overdigestion with *Tth*1111, 10% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 5u of *Tth*111I for 16 hours at 65°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.

Ordering Information

Catalog No	Pack Size
RE1356	200u



λDNA (d III Digest) 0.7% Agarose a - Digestion after 1-fold

b - Digestion after 3-fold

Activity in Reaction Buffer						
V1 V2 V3 V4 V5						

50% 75% 25% 50% 100%

Van 911

Please refer to AccB7I (RE1108 - page 035

Vha464| {Aft||}

Concentration

4-20u/µl

5'...C[↓]TTAAG...3' 3'...GAATT_↑C...5'

Reaction Conditions

1X Buffer V3

50mM Tris-HCI (pH7.5 at 30°C), 10mM MgCI2, 100mM NaCI, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 5-fold overdigestion with *Vha*464I, 40% of the DNA fragments can be ligated and recut.

Overdigestion Assay

Ordering Information

An unaltered banding pattern was observed after 1µg of DNA was digested with 4u of *Vha*464l for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

a b

λDNA 0.7% Agarose a - Digestion after 1 hour

b - Digestion after 16 hours

Catalog No	Pack Size	
RE1358	600u	



Vnel {ApaLl}

Concentration

1-10u/µl

5'...G¹TGCAC...3' 3'...CACGT₁G...5'

Reaction Conditions

1X Buffer V3

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂, 10mM NaCl, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *Vnel*, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of *Vne*l for 16 hours at 37°C.

Supplied with 10X Buffer V3, 10X Buffer UB and Viva Buffer A. (Diluent)

Ordering Information

Catalog No	Pack Size
RE1360	500u

V3_{Bff}







λDNA 0.7% Agarose

a - Digestion after 1 hour b - Digestion after 16 hours

Vspl

Concentration

 $10-30u/\mu l$

5'...AT¹TAAT...3' 3'...TAAT₁TA...5'

Reaction Conditions

1X Buffer V4

10mM Tris-HCl (pH 8.5 at 30°C), 10mM MgCl $_{\rm 2}$, 100mM KCl, and 100 μ g/ml BSA. Incubate at 37°C.

Storage Buffer

10mM $\rm KH_2PO_4$ (pH 7.2), 50mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200 $\mu g/ml$ BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 10-fold overdigestion with *VspI*, 70% of the DNA fragments can be ligated and recut.

Overdigestion Assay

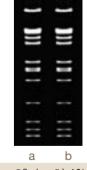
An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of *Vsp*I for 16 hours at 37°C.

Supplied with 10X Buffer V4, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam- methylation ATTA^mAT

Ordering Information

Catalog No	Pack Size
RE1362	600u



DNA (dam⁻& dcm⁻) 1% Agarose a - Digestion after 1 hour b - Digestion after 16 hours

 Activity in Reaction Buffer

 V1
 V2
 V3
 V4
 V5

 75%
 75%
 50%
 100%
 10%

Xagl

Please refer to Bst EN I (RE1216 - page 066)



Xapl

Please refer to Acl I (RE1112 - page 036)

Xbal

Concentration

10-50u/µl

5'...T¹CTAGA...3' 3'...AGATC₁T...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM NaCl. 0.1mM EDTA, 7mM 2-mercaptoethanol, and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 30-fold overdigestion with Xbal. 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 30u of Xbal for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation (G^mATC): TCTAGATC

Ordering Information

Catalog No	Pack Size
RV1364	800u



λDNA (dam-&dcm-) (Hind III Digest) 1.0% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
10%	75%	75%	10%	100%

Xmal {Smal*}

Concentration

1-3u/µl

5'...C¹CCGGG...3' 3'...GGGCC₁C...5'

Reaction Conditions

1X Buffer V5

30mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 60mM K-acetate, and 100ug/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 100mM NaCl, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 3-fold overdigestion with I, more than 95% of the DNA fragments can be ligated and recut.

Overdigestion Assav

An unaltered banding pattern was observed after 1µg of DNA was digested with 2u of I for 16 hours at 37°C.

Supplied with 10X Buffer V5, 10X Buffer UB and Viva Buffer A. (Diluent)

* Blocked by overlapping dam-methylation ATTA^mAT

Ordering Information

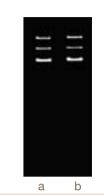
Catalog No	Pack Size
RV1366	100u











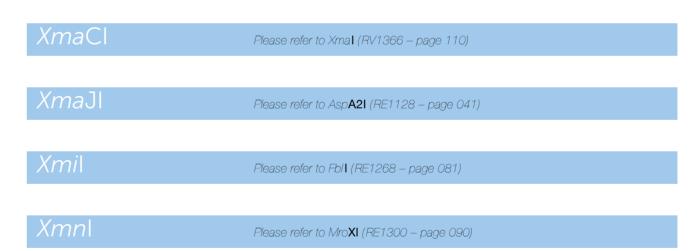
λDNA 0.7% Agarose

a - Digestion after 1 hour

b - Digestion after 16 hours

Activity in Reaction Buffer V1 V2 V3 V4 V5 100% 100% 75% 75% 100%

Xcel Please refer to Bst NSI (RE1232 - page 070) Xhol Please refer to Sfr274I (RE1332 - page 102) Xholl Please refer to Bst X2I (RE1242 - page 073)



Restriction Endonucleases

Vivantis Technologies . Product Catalog Volume IV

Zral {Aatll*}

Concentration

3-10u/µl

5'...GAC[‡]GTC...3' 3'...CTG_↑CAG...5'

Reaction Conditions

1X Buffer UB

25mM Tris-acetate (pH 7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol and 50µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 50mM KCI, 0.1mM EDTA, 10mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

80°C for 20 minutes



After 5-fold overdigestion with Zral, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 4u of Zral for 16 hours at 37°C.

Supplied with 10X Buffer UB and Viva Buffer A. (Diluent)

* High enzyme concentration may result in Star Activity.



λDNA 0.7% Agarose

- a Digestion after 1 hour
- b Digestion after 16 hours

Ordering Information

Catalog No	Pack Size
RE1368	100u

Activity in Reaction Buffer				
V1	V2	V3	V4	V5
50%	75%	25%	50%	50%

Zrml

Please refer to BmcAl (RV1146 - page 047)

Zsp2l {Avalll}

Concentration

5-20u/µl

5'...ATGCA[‡]T...3' 3'...T₁ACGTA...5'

Reaction Conditions

1X Buffer V1

10mM Tris-HCI (pH 7.5 at 30°C), 10mM MgCl₂ and 100µg/ml BSA. Incubate at 37°C.

Storage Buffer

10mM Tris-HCI (pH 7.5), 200mM KCI, 0.1mM EDTA, 7mM 2-mercaptoethanol, 200µg/ml BSA and 50% glycerol. Store at -20°C.

Thermal Inactivation

65°C for 20 minutes

Ligation / Recutting Assay

After 20-fold overdigestion with Zsp2I, 90% of the DNA fragments can be ligated and recut.

Overdigestion Assay

An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Zsp2l for 16 hours at 37°C.

Supplied with 10X Buffer V1, 10X Buffer UB and Viva Buffer A. (Diluent)

Enzyme is stable for up to 6 months if properly stored. It is recommended that the enzyme is kept in small aliquot to avoid repeated freeze-thaw cycles.

Catalog No	Pack Size
RE1370	600u











λDNA 0.7% Agarose

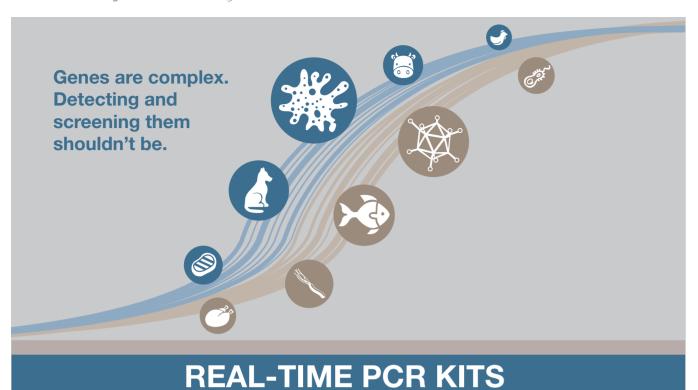
- a Digestion after 1 hour b - Digestion after 16 hours
- Activity in Reaction Buffer V1 V2 V3 V4 V5 100% 50% 25% 50% 100%





Real-time PCR Products

Real-time PCR Products



ViPrimePLUS qPCR/RT-qPCR Kits



Real-time PCR has become one of the gold standard approaches in gene quantification for detection of infectious diseases owing to its sensitivity, specificity, speed and quantitative features. ViPrimePLUS PCR/RT-qPCR products from Vivantis offer rapid and reliable qPCR solution with more than 200 kits for detection and screening of pathogens in human and veterinary as well as meat identification. The primers and probes are specifically designed to have 100% specificity against a broad range of reference samples.

Features

Advanced : E

: Equipped with Taqman probe technology

Broad selections : Wide selection of kits for more than 200 subtypes of pathogens

Rapid : Fast and Easy protoc

Reliable : Reliable quantification & confirmation of results by inclusion of internal extraction control

Sensitive : Sensitive to <100 copies of target genome of pathogens

Specific : Highly specific genome detection

ViPrimePLUS qPCR/RT-qPCR Products

Human Infection qPCR/RT-qPCR Kits

Description

ViPrimePLUS Human Infection qPCR/RT-qPCR Kits provide wide selection of kits for rapid and reliable screening and detection of clinically important diseases such as sexually transmitted infections, TORCH infections, respiratory tract infections, intestinal infections as well as viral infections which include human herpes virus, human papillomavirus, hepatitis virus, influenza virus and etc.

No	Catalog No	Description	Pack Size
Ame	eoba		
1	QM1001	ViPrimePLUS Acanthamoeba qPCR Kit	150 reactions
2	QM1002	ViPrimePLUS Balamuthia mandrillaris qPCR Kit	150 reactions
3	QM1003	ViPrimePLUS Entamoeba qPCR Kit	150 reactions
4	QM1004	ViPrimePLUS Entamoeba histolytica qPCR Kit	150 reactions
5	QM1005	ViPrimePLUS Naegleria qPCR Kit	150 reactions
Bac	terial		
1	QM2001	ViPrimePLUS Acinetobacter baumannii qPCR kit	150 reactions
2	QM2002	ViPrimePLUS Aggregatibacter actinomycetemcomitans qPCR Kit	150 reactions
3	QM2003	ViPrimePLUS Bacillus anthracis qPCR Kit	150 reactions
4	QM2004	ViPrimePLUS Bacillus cereus E33 qPCR Kit	150 reactions
5	QM2005	ViPrimePLUS Bordetella pertussis qPCR Kit	150 reactions
6	QM2006	ViPrimePLUS Brucella abortus qPCR Kit	150 reactions
7	QM2007	ViPrimePLUS Brucella qPCR Kit	150 reactions
8	QM2008	ViPrimePLUS Burkholderia cepacia complex qPCR Kit	150 reactions
9	QM2009	ViPrimePLUS Burkholderia pseudomallei qPCR Kit	150 reactions
10	QM2010	ViPrimePLUS Campylobacter coli qPCR Kit	150 reactions
11	QM2011	ViPrimePLUS Campylobacter jejuni qPCR Kit	150 reactions
12	QM2012	ViPrimePLUS Chlamydia trachomatis qPCR Kit	150 reactions
13	QM2013	ViPrimePLUS Chlamydophila pneumoniae qPCR Kit	150 reactions
14	QM2014	ViPrimePLUS Clostridium difficile (toxin A) qPCR Kit	150 reactions
15	QM2015	ViPrimePLUS Clostridium difficile (toxin B) qPCR Kit	150 reactions
16	QM2016	ViPrimePLUS Clostridium estertheticum qPCR Kit	150 reactions
17	QM2017	ViPrimePLUS Clostridium perfringens qPCR Kit	150 reactions
18	QM2018	ViPrimePLUS Clostridium perfringens A & B qPCR Kit	150 reactions
19	QM2019	ViPrimePLUS Clostridium tetani qPCR Kit	150 reactions
20	QM2020	ViPrimePLUS Corynebacterium diphtheriae A & B qPCR Kit	150 reactions
21	QM2021	ViPrimePLUS Cryptococcus neoformans qPCR Kit	150 reactions
22	QM2022	ViPrimePLUS Enterococcus casseliflavus qPCR Kit	150 reactions
23	QM2023	ViPrimePLUS Enterococcus faecalis qPCR Kit	150 reactions
24	QM2024	ViPrimePLUS Enterococcus faecium qPCR Kit	150 reactions
25	QM2025	ViPrimePLUS Escherichia coli qPCR Kit	150 reactions

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Bac	terial		1
26	QM2026	ViPrimePLUS Escherichia coli O157:H7 qPCR Kit	150 reactions
27	QM2027	ViPrimePLUS Escherichia coli O104:H4 qPCR Kit	150 reactions
28	QM2028	ViPrimePLUS Francisella tularensis qPCR Kit	150 reactions
29	QM2029	ViPrimePLUS Haemophilus ducreyi qPCR Kit	150 reactions
30	QM2030	ViPrimePLUS Haemophilus influenzae qPCR Kit	150 reactions
31	QM2031	ViPrimePLUS Helicobacter pylori qPCR Kit	150 reactions
32	QM2032	ViPrimePLUS Klebsiella pneumoniae qPCR Kit	150 reactions
33	QM2033	ViPrimePLUS Legionella qPCR Kit	150 reactions
34	QM2034	ViPrimePLUS Legionella pneumophila qPCR Kit	150 reactions
35	QM2035	ViPrimePLUS Leptospirosis qPCR Kit	150 reactions
36	QM2036	ViPrimePLUS Listeria monocytogenes qPCR Kit	150 reactions
37	QM2037	ViPrimePLUS Lyme Disease qPCR Kit	150 reactions
38	QM2038	ViPrimePLUS MRSA qPCR Kit	150 reactions
39	QM2039	ViPrimePLUS Moraxella catarrhalis qPCR Kit	150 reactions
40	QM2040	ViPrimePLUS Mycobacterium avium complex qPCR Kit	150 reactions
41	QM2041	ViPrimePLUS Mycobacterium tuberculosis complex qPCR Kit	150 reactions
42	QM2042	ViPrimePLUS Mycobacterium tuberculosis qPCR Kit	150 reactions
43	QM2043	ViPrimePLUS Mycoplasma hominis qPCR Kit	150 reactions
44	QM2044	ViPrimePLUS Mycoplasma pneumoniae qPCR Kit	150 reactions
45	QM2045	ViPrimePLUS Neisseria gonorrhoeae qPCR Kit	150 reactions
46	QM2046	ViPrimePLUS Neisseria meningitidis qPCR Kit	150 reactions
47	QM2047	ViPrimePLUS Porphyromonas gingivalis qPCR Kit	150 reactions
48	QM2048	ViPrimePLUS Pseudomonas aeruginosa qPCR Kit	150 reactions
49	QM2049	ViPrimePLUS Rickettsia qPCR Kit	150 reactions
50	QM2050	ViPrimePLUS Salmonella enterica qPCR Kit	150 reactions
51	QM2051	ViPrimePLUS Salmonella qPCR Kit	150 reactions
52	QM2052	ViPrimePLUS Shewanella putrefaciens qPCR Kit	150 reactions
53	QM2053	ViPrimePLUS Shigella qPCR Kit	150 reactions
54	QM2054	ViPrimePLUS Simkania negevensis qPCR Kit	150 reactions
55	QM2055	ViPrimePLUS Staphylococcus aureus qPCR Kit	150 reactions
56	QM2056	ViPrimePLUS Staphylococcus epidermidis qPCR Kit	150 reactions
57	QM2057	ViPrimePLUS Staphylococcus haemolyticus qPCR Kit	150 reactions
58	QM2058	ViPrimePLUS Streptococcus agalactiae qPCR Kit	150 reactions
59	QM2059	ViPrimePLUS Streptococcus mitis qPCR Kit	150 reactions
60	QM2060	ViPrimePLUS Streptococcus mutans qPCR Kit	150 reactions
61	QM2061	ViPrimePLUS Streptococcus oralis qPCR Kit	150 reactions
62	QM2062	ViPrimePLUS Streptococcus pneumoniae qPCR Kit	150 reactions
63	QM2063	ViPrimePLUS Streptococcus pyogenes qPCR Kit	150 reactions
64	QM2064	ViPrimePLUS Streptococcus sanguinis qPCR Kit	150 reactions
65	QM2065	ViPrimePLUS Treponema pallidum qPCR Kit	150 reactions
66	QM2066	ViPrimePLUS Ureaplasma urealyticum qPCR Kit	150 reactions
67	QM2067	ViPrimePLUS Vibrio qPCR Kit	150 reactions
68	QM2068	ViPrimePLUS Vibrio cholerae qPCR Kit	150 reactions
69	QM2069	ViPrimePLUS Yersinia enterolitica qPCR Kit	150 reactions

Bac	terial		I
63	QM2063	ViPrimePLUS Streptococcus pyogenes qPCR Kit	150 reactions
64	QM2064	ViPrimePLUS Streptococcus sanguinis qPCR Kit	150 reactions
65	QM2065	ViPrimePLUS Treponema pallidum qPCR Kit	150 reactions
66	QM2066	ViPrimePLUS <i>Ureaplasma urealyticum</i> qPCR Kit	150 reactions
67	QM2067	ViPrimePLUS Vibrio qPCR Kit	150 reactions
68	QM2068	ViPrimePLUS Vibrio cholerae qPCR Kit	150 reactions
69	QM2069	ViPrimePLUS Yersinia enterolitica qPCR Kit	150 reactions
Funç	gal		
1	QM3001	ViPrimePLUS Ajellomyces capsulatus qPCR Kit	150 reactions
2	QM3002	ViPrimePLUS Aspergillus qPCR Kit	150 reactions
3	QM3003	ViPrimePLUS Aspergillus fumigatus qPCR Kit	150 reactions
4	QM3004	ViPrimePLUS Botrytis cinerea qPCR Kit	150 reactions
5	QM3005	ViPrimePLUS Candida albicans qPCR Kit	150 reactions
6	QM3006	ViPrimePLUS Encephalitozoon qPCR Kit	150 reactions
7	QM3007	ViPrimePLUS Enterocytozoon bieneusi qPCR Kit	150 reactions
8	QM3008	ViPrimePLUS Fungi qPCR Kit	150 reactions
9	QM3009	ViPrimePLUS <i>Pneumocystis jirovecii</i> qPCR Kit	150 reactions
Para	asitic		
1	QM4001	ViPrimePLUS African trypanosomiasis qPCR Kit	150 reactions
2	QM4002	ViPrimePLUS Ancylostoma duodenale qPCR Kit	150 reactions
3	QM4003	ViPrimePLUS Cryptosporidium qPCR Kit	150 reactions
4	QM4004	ViPrimePLUS <i>Ehrlichia</i> qPCR Kit	150 reactions
5	QM4005	ViPrimePLUS Giardia intestinalis qPCR Kit	150 reactions
6	QM4006	ViPrimePLUS Leishmania qPCR Kit	150 reactions
7	QM4007	ViPrimePLUS Leishmania infantum qPCR Kit	150 reactions
8	QM4008	ViPrimePLUS Leishmania major qPCR Kit	150 reactions
9	QM4009	ViPrimePLUS Leishmania tropica qPCR Kit	150 reactions
10	QM4010	ViPrimePLUS <i>Plasmodium</i> qPCR Kit	150 reactions
11	QM4011	ViPrimePLUS Plasmodium falciparum qPCR Kit	150 reactions
12	QM4012	ViPrimePLUS Plasmodium knowlesi qPCR Kit	150 reactions
13	QM4013	ViPrimePLUS Plasmodium malariae qPCR Kit	150 reactions
14	QM4014	ViPrimePLUS Plasmodium ovale qPCR Kit	150 reactions
15	QM4015	ViPrimePLUS <i>Plasmodium vivax</i> qPCR Kit	150 reactions
16	QM4016	ViPrimePLUS Toxoplasma gondii qPCR Kit	150 reactions
17	QM4017	ViPrimePLUS Trichomonas vaginalis qPCR Kit	150 reactions
18	QM4018	ViPrimePLUS Trypanosoma cruzi qPCR Kit	150 reactions
Viral			
1	QM5001	ViPrimePLUS Adenovirus Type B qPCR Kit	150 reactions
2	QM5002	ViPrimePLUS Adenovirus Type C qPCR Kit	150 reactions
3	QM5003	ViPrimePLUS Adenovirus Type F and G qPCR Kit	150 reactions
4	QM5004	ViPrimePLUS Chikungunya Virus RT-qPCR Kit	150 reactions
5	QM5005	ViPrimePLUS Crimean-Congo Haemorrhagic Fever Virus RT-qPCR Kit	150 reactions
6	QM5006	ViPrimePLUS Dengue Virus subtypes 1, 2, 3, 4 RT-qPCR Kit	150 reactions
7	QM5007	ViPrimePLUS Dobrava-Belgrade Virus RT-qPCR Kit	150 reactions

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Vira			
8	QM5008	ViPrimePLUS Ebola Virus (2014 Outbreak) RT-qPCR Kit	150 reactions
9	QM5009	ViPrimePLUS Enterovirus RT-qPCR Kit	150 reactions
10	QM5010	ViPrimePLUS Hepatitis A Virus RT-qPCR Kit	150 reactions
11	QM5011	ViPrimePLUS Hepatitis B Virus qPCR Kit	150 reactions
12	QM5012	ViPrimePLUS Hepatitis C Virus RT-qPCR Kit	150 reactions
13	QM5013	ViPrimePLUS Hepatitis Delta Virus RT-qPCR Kit	150 reactions
14	QM5014	ViPrimePLUS Hepatitis E Virus RT-qPCR Kit	150 reactions
15	QM5015	ViPrimePLUS Human Bocavirus qPCR Kit	150 reactions
16	QM5016	ViPrimePLUS Human Coronavirus Group 1b RT-qPCR Kit	150 reactions
17	QM5017	ViPrimePLUS Human Coronavirus Group 2a RT-qPCR Kit	150 reactions
18	QM5018	ViPrimePLUS Herpes Simplex Virus 1 and 2 qPCR Kit	150 reactions
19	QM5019	ViPrimePLUS Herpes Simplex Virus 1 qPCR Kit	150 reactions
20	QM5020	ViPrimePLUS Herpes Simplex Virus 2 qPCR Kit	150 reactions
21	QM5021	ViPrimePLUS Human Herpes Virus 3 qPCR Kit	150 reactions
22	QM5022	ViPrimePLUS Human Herpes Virus 4 qPCR Kit	150 reactions
23	QM5023	ViPrimePLUS Human Herpes Virus 5 qPCR Kit	150 reactions
24	QM5024	ViPrimePLUS Human Herpes Virus 6 qPCR Kit	150 reactions
25	QM5025	ViPrimePLUS Human Herpes Virus 7 qPCR Kit	150 reactions
26	QM5026	ViPrimePLUS Human Herpes Virus 8 qPCR Kit	150 reactions
27	QM5027	ViPrimePLUS Human Immunodeficiency Virus type 1 RT-qPCR Kit	150 reactions
28	QM5028	ViPrimePLUS Human Immunodeficiency Virus type 2 RT-qPCR Kit	150 reactions
29	QM5029	ViPrimePLUS Human Influenza A Virus (M1) RT-qPCR Kit	150 reactions
30	QM5030	ViPrimePLUS Human Influenza A Virus (M2) RT-qPCR Kit	150 reactions
31	QM5031	ViPrimePLUS Human Influenza A Virus subtype H1 RT-qPCR Kit	150 reactions
32	QM5032	ViPrimePLUS Human Influenza A Virus subtype H3 RT-qPCR Kit	150 reactions
33	QM5033	ViPrimePLUS Human Influenza B Virus RT-qPCR Kit	150 reactions
34	QM5034	ViPrimePLUS Human Measles Virus RT-qPCR Kit	150 reactions
35	QM5035	ViPrimePLUS Human Metapneumovirus RT-qPCR Kit	150 reactions
36	QM5036	ViPrimePLUS Human Papillomavirus 11 qPCR Kit	150 reactions
37	QM5037	ViPrimePLUS Human Papillomavirus 16 qPCR Kit	150 reactions
38	QM5038	ViPrimePLUS Human Papillomavirus 18 qPCR Kit	150 reactions
39	QM5039	ViPrimePLUS Human Papillomavirus 33 qPCR Kit	150 reactions
40	QM5040	ViPrimePLUS Human Papillomavirus 52 & 52b qPCR Kit	150 reactions
41	QM5041	ViPrimePLUS Human Papillomavirus 58 qPCR Kit	150 reactions
42	QM5042	ViPrimePLUS Human Papillomavirus 6 qPCR Kit	150 reactions
43	QM5043	ViPrimePLUS Human Parainfluenza Virus Type 1 RT-qPCR Kit	150 reactions
44	QM5044	ViPrimePLUS Human Parainfluenza Virus Type 2 RT-qPCR Kit	150 reactions
45	QM5045	ViPrimePLUS Human Parainfluenza Virus Type 3 RT-qPCR Kit	150 reactions
46	QM5046	ViPrimePLUS Human Parainfluenza Virus Type 4a RT-qPCR Kit	150 reactions
47	QM5047	ViPrimePLUS Human Parainfluenza Virus Type 4b RT-qPCR Kit	150 reactions
48	QM5048	ViPrimePLUS Human Parvovirus B19 qPCR Kit	150 reactions
49	QM5049	ViPrimePLUS Human Rhinovirus 14 RT-qPCR Kit	150 reactions
50	QM5050	ViPrimePLUS Human Rhinovirus 16 RT-qPCR Kit	150 reactions

52 C 53 C 54 C 55 C 56 C 57 C	QM5051 QM5052 QM5053 QM5054 QM5055 QM5056 QM5057 QM5058 QM5059	ViPrimePLUS Human Rhinovirus 1B RT-qPCR Kit ViPrimePLUS Human Rhinovirus 29 RT-qPCR Kit ViPrimePLUS Human Rhinovirus 9 RT-qPCR Kit ViPrimePLUS Human Rhinovirus RT-qPCR Kit ViPrimePLUS Human Rotavirus B RT-qPCR Kit ViPrimePLUS Human Rotavirus C RT-qPCR Kit ViPrimePLUS Human T-lymphotropic Virus 1 RT-qPCR Kit ViPrimePLUS Human T-lymphotropic Virus 2 RT-qPCR Kit	150 reactions
53 C 54 C 55 C 56 C 57 C	QM5053 QM5054 QM5055 QM5056 QM5057 QM5058	ViPrimePLUS Human Rhinovirus 9 RT-qPCR Kit ViPrimePLUS Human Rhinovirus RT-qPCR Kit ViPrimePLUS Human Rotavirus B RT-qPCR Kit ViPrimePLUS Human Rotavirus C RT-qPCR Kit ViPrimePLUS Human T-lymphotropic Virus 1 RT-qPCR Kit	150 reactions 150 reactions 150 reactions 150 reactions
54 C 55 C 56 C 57 C	QM5054 QM5055 QM5056 QM5057 QM5058	ViPrimePLUS Human Rhinovirus RT-qPCR Kit ViPrimePLUS Human Rotavirus B RT-qPCR Kit ViPrimePLUS Human Rotavirus C RT-qPCR Kit ViPrimePLUS Human T-lymphotropic Virus 1 RT-qPCR Kit	150 reactions 150 reactions 150 reactions
55 C 56 C 57 C	QM5055 QM5056 QM5057 QM5058	ViPrimePLUS Human Rotavirus B RT-qPCR Kit ViPrimePLUS Human Rotavirus C RT-qPCR Kit ViPrimePLUS Human T-lymphotropic Virus 1 RT-qPCR Kit	150 reactions 150 reactions
56 C	QM5056 QM5057 QM5058	ViPrimePLUS Human Rotavirus C RT-qPCR Kit ViPrimePLUS Human T-lymphotropic Virus 1 RT-qPCR Kit	150 reactions
57 C	QM5057 QM5058	ViPrimePLUS Human T-lymphotropic Virus 1 RT-qPCR Kit	
	QM5058		150 reactions
		ViPrimePLUS Human T-lymphotropic Virus 2 RT-qPCR Kit	
58 C	M5059		150 reactions
59 C		ViPrimePLUS Japanese Encephalitis Virus RT-qPCR Kit	150 reactions
60 C	QM5060	ViPrimePLUS Merkel Cell Polyomavirus qPCR Kit	150 reactions
61 G	QM5061	ViPrimePLUS Mumps Virus RT-qPCR Kit	150 reactions
62 G	QM5062	ViPrimePLUS Norovirus Genotype 1 RT-qPCR Kit	150 reactions
63 C	QM5063	ViPrimePLUS Norovirus Genotype 2 RT-qPCR Kit	150 reactions
64 C	QM5064	ViPrimePLUS Novel Coronavirus hCoV-EMC / MERS RT-qPCR Kit	150 reactions
65 C	QM5065	ViPrimePLUS Polyomavirus BK qPCR Kit	150 reactions
66 C	QM5066	ViPrimePLUS Polyomavirus JC qPCR Kit	150 reactions
67 C	QM5067	ViPrimePLUS Rabies Virus RT-qPCR Kit	150 reactions
68 C	QM5068	ViPrimePLUS Respiratory Syncytial Virus RT-qPCR Kit	150 reactions
69 C	QM5069	ViPrimePLUS Respiratory Syncytial Virus Type A RT-qPCR Kit	150 reactions
70 C	QM5070	ViPrimePLUS Respiratory Syncytial Virus Type B RT-qPCR Kit	150 reactions
71 C	QM5071	ViPrimePLUS Rotavirus A RT-qPCR Kit	150 reactions
72 C	QM5072	ViPrimePLUS Rubella Virus RT-qPCR Kit	150 reactions
73 C	QM5073	ViPrimePLUS Severe Acute Respiratory Syndrome RT-qPCR Kit	150 reactions
74 C	QM5074	ViPrimePLUS Simian Virus 40 qPCR Kit	150 reactions
75 C	QM5075	ViPrimePLUS Sin Nombre Hantavirus RT-qPCR Kit	150 reactions
76 C	QM5076	ViPrimePLUS H1N1 Influenza Human Pandemic Strain RT-qPCR Kit	150 reactions
77 C	QM5077	ViPrimePLUS West Nile Virus RT-qPCR Kit	150 reactions
78 C	QM5078	ViPrimePLUS Yellow Fever Virus RT-qPCR Kit	150 reactions
79 C	QM5079	ViPrimePLUS Zika Virus RT-qPCR Kit	150 reactions

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Veterinary Infection qPCR/RT-qPCR Kits

Description

ViPrimePLUS Veterinary Infection qPCR/RT-qPCR Kits are designed for rapid and reliable screening and detection of pathogen genomes which infect animals.

No	Catalog No	Description	Pack Size
Avi	an		
1	QV1001	ViPrimePLUS Anaplasma phagocytophilum qPCR Kit	150 reactions
2	QV1002	ViPrimePLUS Avian Adenovirus (Egg Drop Syndrome) qPCR Kit	150 reactions
3	QV1003	ViPrimePLUS Avian Infectious Bronchitis Virus RT-qPCR Kit	150 reactions
4	QV1004	ViPrimePLUS Avian Influenza A Virus subtype (H5) RT-qPCR Kit	150 reactions
5	QV1005	ViPrimePLUS Avian Influenza A Virus subtype (H6) RT-qPCR Kit	150 reactions
6	QV1006	ViPrimePLUS Avian Influenza A Virus subtype (H7) RT-qPCR Kit	150 reactions
7	QV1007	ViPrimePLUS Avian Influenza A Virus subtype (H9) RT-qPCR Kit	150 reactions
8	QV1008	ViPrimePLUS Avian Influenza A Virus subtype H5N1 RT-qPCR Kit	150 reactions
9	QV1009	ViPrimePLUS Beak and Feather Disease Virus qPCR Kit	150 reactions
10	QV1010	ViPrimePLUS Budgerigar Fledgling Disease Virus qPCR Kit	150 reactions
11	QV1011	ViPrimePLUS Burkholderia mallei qPCR Kit	150 reactions
12	QV1012	ViPrimePLUS Chicken Anemia Virus qPCR Kit	150 reactions
13	QV1013	ViPrimePLUS Chlamydophila psittaci qPCR Kit	150 reactions
14	QV1014	ViPrimePLUS Duck Hepatitis B Virus qPCR Kit	150 reactions
15	QV1015	ViPrimePLUS Fowlpox Virus qPCR Kit	150 reactions
16	QV1016	ViPrimePLUS Gallid herpesvirus 1 qPCR Kit	150 reactions
17	QV1017	ViPrimePLUS Gallid herpesvirus 2 qPCR Kit	150 reactions
18	QV1018	ViPrimePLUS Infectious Bursal Disease Virus RT-qPCR Kit	150 reactions
19	QV1019	ViPrimePLUS Influenza Virus H7N9 RT-qPCR Kit	150 reactions
20	QV1020	ViPrimePLUS Mycoplasma gallisepticum qPCR Kit	150 reactions
21	QV1021	ViPrimePLUS Newcastle Disease Virus RT-qPCR Kit	150 reactions
Ca	nine & Feline		
1	QV2001	ViPrimePLUS Ancylostoma duodenale qPCR Kit	150 reactions
2	QV2002	ViPrimePLUS Canine babesiosis qPCR Kit	150 reactions
3	QV2003	ViPrimePLUS Canine Distemper Virus RT-qPCR Kit	150 reactions
4	QV2004	ViPrimePLUS Canine Herpesvirus qPCR Kit	150 reactions
5	QV2005	ViPrimePLUS Chlamydophila felis qPCR Kit	150 reactions
6	QV2006	ViPrimePLUS Feline Calicivirus RT-qPCR Kit	150 reactions
7	QV2007	ViPrimePLUS Feline Coronavirus RT-qPCR Kit	150 reactions
8	QV2008	ViPrimePLUS Feline Herpesvirus qPCR Kit	150 reactions
9	QV2009	ViPrimePLUS Feline Immunodeficiency Virus RT-qPCR Kit	150 reactions
10	QV2010	ViPrimePLUS Feline Leukaemia Virus RT-qPCR Kit	150 reactions
11	QV2011	ViPrimePLUS Microsporum canis qPCR Kit	150 reactions
12	QV2012	ViPrimePLUS Mycoplasma felis qPCR Kit	150 reactions
13	QV2013	ViPrimePLUS Mycoplasma haemofelis qPCR Kit	150 reactions
50	QM5050	ViPrimePLUS Human Rhinovirus 16 RT-qPCR Kit	150 reactions

Equ	uine		I
1	QV3001	ViPrimePLUS African Horse Sickness Virus RT-qPCR Kit	150 reactions
2	QV3002	ViPrimePLUS <i>Babesia caballi</i> qPCR Kit	150 reactions
3	QV3003	ViPrimePLUS Equid Herpesvirus 1 qPCR Kit	150 reactions
4	QV3004	ViPrimePLUS Equid Herpesvirus 4 qPCR Kit	150 reactions
5	QV3005	ViPrimePLUS Equine Infectious Anaemia Virus RT-qPCR Kit	150 reactions
6	QV3006	ViPrimePLUS Theileria equi qPCR Kit	150 reactions
7	QV3007	ViPrimePLUS Trypanosoma equiperdum qPCR Kit	150 reactions
Piso	cean (Fish)		
1	QV4001	ViPrimePLUS Cyprinid Herpesvirus 3 qPCR Kit	150 reactions
2	QV4002	ViPrimePLUS Grass Carp Reovirus RT-qPCR Kit	150 reactions
3	QV4003	ViPrimePLUS Infectious Hematopoietic Necrosis Virus RT-qPCR Kit	150 reactions
4 5	QV4004 QV4005	ViPrimePLUS Infectious Pancreatic Necrosis Virus RT-qPCR Kit ViPrimePLUS Spring Viremia of Carp Virus RT-qPCR Kit	150 reactions 150 reactions
6	QV4006	ViPrimePLUS Viral Hemorrhagic Septicemia Virus RT-qPCR Kit	150 reactions
	cine	The same and the s	.001000.01.0
1	QV5001	ViPrimePLUS <i>Chlamydiaceae</i> (all species) qPCR Kit	150 reactions
2	QV5001	ViPrimePLUS <i>Porcine Circovirus</i> 1 qPCR Kit	150 reactions
3	QV5003	ViPrimePLUS <i>Porcine Circovirus 2</i> qPCR Kit	150 reactions
4	QV5004	ViPrimePLUS Porcine Reproductive and Respiratory Syndrome Virus-	150 reactions
		European Genotype RT-qPCR Kit	
5	QV5005	ViPrimePLUS Porcine Reproductive and Respiratory Syndrome Virus-	150 reactions
		US Genotype RT-qPCR Kit	
Rur	minants		
1	QV6001	ViPrimePLUS Anaplasma centrale qPCR Kit	150 reactions
2	QV6002	ViPrimePLUS Anaplasma marginale qPCR Kit	150 reactions
3	QV6003	ViPrimePLUS <i>Babesia bigemina</i> qPCR Kit	150 reactions
4	QV6004	ViPrimePLUS Babesia bovis qPCR Kit	150 reactions
5	QV6005	ViPrimePLUS Bluetongue Virus RT-qPCR Kit	150 reactions
6	QV6006	ViPrimePLUS Bluetongue Virus 1 RT-qPCR Kit	150 reactions
7	QV6007	ViPrimePLUS Bluetongue Virus 8 RT-qPCR Kit	150 reactions
8	QV6008	ViPrimePLUS Bovine Herpesvirus 1 qPCR Kit	150 reactions
9	QV6009	ViPrimePLUS Bovine Leukemia Virus RT-qPCR Kit	150 reactions
10	QV6010	ViPrimePLUS Campylobacter fetus-venerialis qPCR Kit	150 reactions
11	QV6011	ViPrimePLUS Campylobacter fetus qPCR Kit	150 reactions
12	QV6012	ViPrimePLUS Capripoxvirus qPCR Kit	150 reactions
13	QV6013	ViPrimePLUS Chlamydophila abortus qPCR Kit	150 reactions
14	QV6014	ViPrimePLUS Foot & Mouth Disease Virus RT-qPCR Kit	150 reactions
15	QV6015	ViPrimePLUS Mycobacterium avium subsp. paratuberculosis qPCR Kit	150 reactions

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Rur	ninants		I
16	QV6016	ViPrimePLUS Mycoplasma mycoides cluster qPCR Kit	150 reactions
17	QV6017	ViPrimePLUS Neospora caninum qPCR Kit	150 reactions
18	QV6018	ViPrimePLUS Pasteurella multocida qPCR Kit	150 reactions
19	QV6019	ViPrimePLUS Peste-des-petits-ruminants Virus RT-qPCR Kit	150 reactions
20	QV6020	ViPrimePLUS Sheep Pox Virus qPCR Kit	150 reactions
21	QV6021	ViPrimePLUS Theileria annulata qPCR Kit	150 reactions
22	QV6022	ViPrimePLUS Theileria parva qPCR Kit	150 reactions
23	QV6023	ViPrimePLUS Tritrichomonas foetus qPCR Kit	150 reactions
24	QV6024	ViPrimePLUS <i>Trypanosoma evansi</i> qPCR Kit	150 reactions
25	QV6025	ViPrimePLUS Wesselsbron Virus RT-qPCR Kit	150 reactions
Oth	er Veterinary		
1	QV7001	ViPrimePLUS Aleutian Disease Virus qPCR Kit	150 reactions
2	QV7002	ViPrimePLUS Epizootic Hemorrhagic Disease Virus RT-qPCR Kit	150 reactions
3	QV7003	ViPrimePLUS Israeli Acute Paralysis Virus RT-qPCR Kit	150 reactions
4	QV7004	ViPrimePLUS Vesivirus_2117 RT-qPCR Kit	150 reactions

Meat Identification qPCR Kits

Description

ViPrime PLUS Meat Identification qPCR Kits are designed for rapid and reliable screening tool for detection of species genome in various food samples especially in raw or cooked meats and meat products.

No	Catalog No	Description	Pack Size
Mea	at Identification		
1	Q10001	ViPrimePLUS Bos taurus (Beef) qPCR Kit	100 reactions
2	Q10002	ViPrimePLUS Gallus gallus (Chicken) qPCR Kit	100 reactions
3	Q10003	ViPrimePLUS Sus scrofa (Pig) qPCR Kit	100 reactions
4	Q10004	ViPrimePLUS Ovies aries (Sheep) qPCR Kit	100 reactions
5	Q10005	ViPrimePLUS Equus caballus (Horse) qPCR Kit	100 reactions
6	Q10006	ViPrimePLUS Felis catus (Cat) qPCR Kit	100 reactions
7	Q10007	ViPrimePLUS Canis familiaris (Dog) qPCR Kit	100 reactions
8	Q10008	ViPrimePLUS <i>Meleagris gallopavo</i> (Turkey) qPCR Kit	100 reactions
9	Q10009	ViPrimePLUS Equus asinus (Donkey) qPCR Kit	100 reactions
10	QI0010	ViPrimePLUS Capra hircus (Goat) qPCR Kit	100 reactions
11	QI0011	ViPrimePLUS Phacochoerus africanus (Warthog) qPCR Kit	100 reactions
12	QI0012	ViPrimePLUS Bubalus bubalis (Buffalo) qPCR Kit	100 reactions



DNA Amplification Products

High Fidelity DNA Polymerases

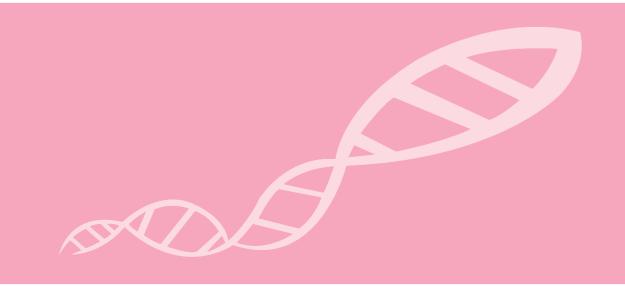
Pfu DNA Polymerase
Chromo Pfu DNA Polymerase
Max Taq DNA Polymerase
Chromo Max Taq DNA Polymerase
AtMax Taq DNA Polymerase

DNA Polymerases

Taq DNA PolymeraseChromo Taq DNA PolymeraseAt Taq DNA PolymeraseChromo At Taq DNA Polymerase

2X Ampli-Optimization Kit
2X At *Taq* Master Mix
2X *Taq* Master Mix

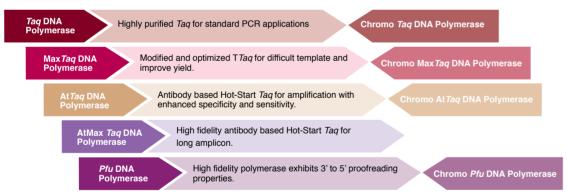
DNA Amplification Kits
My PCR Kits
Buffers





PCR is an invaluable tool in molecular biology research, and at the heart of this application is the DNA polymerase. At Vivantis Technologies, we believe that a successful PCR starts with quality Polymerases. You

polymerase. At Vivantis Technologies, we believe that a successful PCR starts with quality Polymerases. You can choose from a premium selection of our polymerases, for standard PCR or Multiplex PCR, to Hot-Start PCR applications. It is our goal to make PCR a simple and easy process for researchers around the world. With Vivantis Technologies, PCR will be a walk in the park.



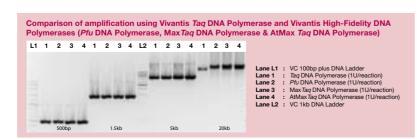
The Chromo DNA Polymerase series is a blend of polymerase with inert colour tracer dyes for easy visualization of the addition of polymerase to the reaction and serve as tracking dye during PCR.

High Fidelity DNA Polymerases

High-Fidelity DNA Polymerases are DNA polymerases which have thermostable properties with 5' to 3' polymerase activity as well as 3' to 5' exonuclease activity which are important for proofreading amplification (the DNA sequence needs to be correct after amplification).

High Fidelity DNA Polymeases:

- Pfu DNA Polymerase
- MaxTaq DNA Polymerase
- AtMax Taq DNA Polymerase



Pfu DNA Polymerase (Pvrococcus furiosus)

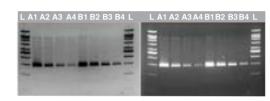
Description

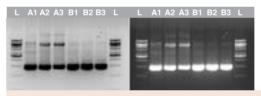
Pfu DNA Polymerase is an extremely thermostable proofreading DNA polymerase, suitable for applications requiring high temperatures synthesis of DNA. *Pfu* DNA Polymerase catalyzes the polymerization of nucleotides into duplex DNA in the 5' to 3' direction with the presence of Mg²⁺. It exhibits the 3' to 5' proofreading activity.

Features

- Ultra pure recombinant protein allows amplification up to 8kb.
- 5X ViBuffer *Pfu* provided for both short and long amplification.

Amplification Using Vivantis Pfu DNA Polymerase





Lane M1: VC 1kb DNA Ladder Lane 1: 0.5kb amplicon

Lane 2: 1.5kb amplicon

Lane 3: 5.0kb amplicon
Lane 4: 8.0kb amplicon

Lane M2: VC Lambda / Hind III Marker

Unit Definition

1u is defined as the amount of enzymes that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCl (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200 μ M each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [³H]dTTP), 10 μ g activated calf thymus DNA and 100 μ g/ml BSA in a final volume of 50 μ l.

Supplied With

- 50X ViBuffer Pfu
- 50mM MgCl₂

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

50mM Tris-HCl (pH 8.0 at 22°C), 100mM KCl, 0.1% TweenTM 20, 0.1% Nonidet-P40, 0.1mM EDTA, 1mM DTT, and 50% glycerol. Store at -20°C.

Catalog No	Description	Pack Size
PL5201 PL5202	Pfu DNA Polymerase Pfu DNA Polymerase	100ս, 5ս/µl 500ս, 5ս/µl

^{*} Please refer to Appendix for the amplification protocol.





Chromo Pfu DNA Polymerase

(Pyrococcus furiosus)



Chromo *Pfu* DNA Polymerase is an extremely thermostable proofreading DNA polymerase, suitable for applications requiring high temperatures synthesis of DNA. *Pfu* DNA Polymerase catalyzes the polymerization of nucleotides into duplex DNA in the 5' to 3' direction with the presence of Mg²⁺. It exhibits the 3' to 5' proofreading activity. The enzyme is supplemented with inert color tracer dyes.

Features

- Color tracer dyes for ease of visualization of the addition of polymerase to the reaction.
- Use of 1u in every 25µl final reaction allows direct loading and serves as tracking dyes during gel electrophoresis. The blue and pink color dyes migrate approximately at 4kb and 0.3kb respectively on 1% TAE agarose gel.
- Ultra pure recombinant protein allows amplification up to 8kb.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon.
- Recommended for use in high-fidelity amplification and cloning of bluntended amplification products.

Amplification Using Vivantis Pfu DNA Polymerase



0.7% TAE agarose gel

Lane M1: VC 1kb DNA Ladder

Lane 0.5 & 1.5kb: 0.5kb PCR amplification product generated using 0.2mM dNTPs and 2.0u Vivantis Pfu DNA Polymerasa

Lane 5 & 8kb: 5kb and 8kb amplification products generated using 0.25mM dNTPs, 2.5u Vivantis Pfu DNA Polymerase and 3% of formamide.

Lane M2: VC Lambda/HindIII Marker

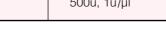
Ordering Information

Catalog No	Description	Pack Size
PL5205	Chromo <i>Pfu</i> DNA Polymerase	100u, 1u/µl
PL5206	Chromo <i>Pfu</i> DNA Polymerase	500u, 1u/µl

^{*} Please refer to Appendix for the amplification protocol.







indicators

Supplemented with

romo

Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCI (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [³H]dTTP), 10µg activated calf thymus DNA and 100µg/ml BSA in a final volume of 50µl.

Supplied With

- 50mM MgCl₂
- 5X ViBuffer Pfu

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

50mM Tris-HCl (pH 8.0 at 22°C), 100mM KCl, 0.1% Tween™ 20, 0.1% Nonidet-P40, 0.1mM EDTA, 1mM DTT, color dyes and 50% glycerol. Store at -20°C.

MaxTaq DNA Polymerase

Description

Max Taq DNA Polymerase is a modified and optimized thermostable enzyme blend containing Taq DNA Polymerase, Pfu DNA Polymerase and enhancing factors. It exhibits the 3' to 5' proofreading activity, resulting in considerably higher amplification fidelity than possible with unmodified Taq DNA Polymerase. Recommended for use in amplification to obtain DNA products up to 20kb.

Features

- Ultra pure recombinant protein allows amplification up to 20kb.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon.
- Excellent for multiplex amplification as it exhibits wider tolerance for Mg²⁺ and salt concentrations.
- Improves amplification results with critical templates, such as those containing GC-rich regions, palindromes or multiple repeats.
- · Increased amplification product yields and purity.
- Generates a mixture of blunt end and 3' dA overhang amplification products, majority of the products are blunt ended.

Amplification Using Vivantis MaxTag DNA Polymerase



0.7% TAE agarose gel

Lane M1: VC Lambda / Hind III Marker

Lane 1: 8kb amplicon
Lane 2: 10kb amplicon

Lane 3: 12kb amplicon

Lane 4: 15kb amplicon

Lane 5: 20kb amplicon Lane 6: 30kb amplicon

Lane 7: 40kb amplicon

Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCl (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [³H]dTTP), 10µg activated calf thymus DNA and 100µg/ml BSA in a final volume of 50µl.

Supplied With

- 10X ViBuffer A (without MgCl₂)
 500mM KCl, 100mM Tris-HCl (pH
 9.1 at 20°C) and 0.1% Triton™
 X-100. The buffer is optimized for
 use with 0.1-0.2mM of each dNTP.
- 10X ViBuffer S 160mM (NH₄)₂SO₄, 500mM Tris-HCl (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% Triton™ X-100. The buffer is optimized for use with 0.35mM of each dNTP.
- 50mM MgCl_a

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

20mM Tris-HCl (pH 8.0 at 22°C), 100mM KCl, 0.5% Tween™ 20, 0.5% Nonidet-P40, 0.1mM EDTA, 1mM DTT and 50% glycerol. Store at -20°C.

Catalog No	Description	Pack Size
PL2201	Max <i>Taq</i> DNA Polymerase	200u, 5u/µl
PL2202	Max <i>Taq</i> DNA Polymerase	500u, 5u/µl

^{*} Please refer to Appendix for the amplification protocol.





Chromo MaxTaq DNA Polymerase

Description

Chromo Max Tag DNA Polymerase is a modified and optimized thermostable enzyme blend containing *Tag* DNA Polymerase. Pfu DNA Polymerase and enhancing factors. It exhibits the 3' to 5' proofreading activity, resulting in considerably higher amplification fidelity than possible with unmodified Tag DNA Polymerase. The enzyme is supplemented with inert color tracer dyes. Recommended for use in amplification to obtain DNA

Features

- Color tracer dyes for ease of visualization of the addition of polymerase to the reaction.
- Use of 1u in every 25µl final reaction allows direct loading and serves as tracking dyes during gel electrophoresis. The blue and pink color dyes migrate approximately at 4kb and 0.3kb respectively on 1% TAE agarose gel.
- Ultra pure recombinant protein allows amplification up to 20kb.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon.
- Excellent for multiplex amplification as it exhibits wider tolerance for Mg²⁺ and salt concentrations.
- Improves amplification results with critical templates, such as those containing GC-rich regions, palindromes or multiple repeats.
- Increased amplification product yields and purity.
- Generates a mixture of blunt end and 3' dA overhang amplification products, majority of the products are blunt ended.

Lane M2: Lambda DNA

Amplification Using Vivantis MaxTag DNA Polymerase



0.7% TAE agarose gel

Lane M1: VC Lamba / Hind III Marker Lane 1: 8kb amplicon Lane 2: 10kb amplicon Lane 3: 12kb amplicon Lane 4: 15kb amplicon Lane 5: 20kb amplicon : 30kb amplicon Lane 7: 40kb amplicon





Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCI (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [3H]dTTP), 10µg activated calf thymus DNA and 100 µg/ml BSA in a final volume of 50µl.

Supplied With

- 10X ViBuffer A (without MaCl_a) 500mM KCl, 100mM Tris-HCl (pH 9.1 at 20°C) and 0.1% Triton™ X-100. The buffer is optimized for use with 0.1-0.2mM of each dNTP.
- · 10X ViBuffer S 160mM (NH₄)₂SO₄, 500mM Tris-HCI (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% Triton™X-100. The buffer is optimized for use with 0.35mM of each dNTP.
- 50mM MgCl_a

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

20mM Tris-HCI (pH 8.0 at 22°C), 100mM KCl, 0.5% Tween™ 20, 0.5% Nonidet-P40, 0.1mM EDTA, 1mM DTT, color dyes and 50% glycerol. Store at -20°C.

Ordering Information

Catalog No	Description	Pack Size
PL2205	Chromo Max <i>Taq</i> DNA Polymerase	200u, 1u/µl
PL2206	Chromo Max <i>Taq</i> DNA Polymerase	500u, 1u/µl

^{*} Please refer to Appendix for the amplification protocol.







AtMax Tag DNA Polymerase (Hot Start Long Amplification)

AtMax Tag DNA Polymerase is a mixture of thermostable Tag DNA Polymerase, proofreading *Pfu* DNA Polymerase, anti-*Taq* DNA Polymerase antibodies, reversible inhibitors and enhancers for automatic "Hot Start" amplification. It exhibits the 3' to 5' proofreading activity, resulting in considerably higher amplification fidelity than possible with unmodified Tag DNA Polymerase.

Features

- Ultra pure recombinant protein is reversibly complexed with an anti-Tag monoclonal antibody that blocks replication activity of the enzyme at moderate temperatures.
- Excellent for multiplex amplification as it exhibits wider tolerance for Mg²⁺ and salt concentrations.
- Improves amplification results with critical templates, such as those containing GC-rich regions, palindromes or multiple repeats.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon

Amplification Using Vivantis AtMax Tag DNA Polymerase



0.5% TAE agarose gel Lane M1: VC 1kb DNA Ladder

Lane 1: 5kb amplicon Lane 2: 8kb amplicon

Lane 3: 10kb amplicon

Lane 4: 15kb amplicon Lane 5: 20kb amplicon

Lane M2: VC Lambda / Hind III Marker

Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCI (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [3H]dTTP), 10µg activated calf thymus DNA and 100µg/ml BSA in a final volume of 50µl.

Supplied With

- 10X ViBuffer A (without MgCl₂) 500mM KCI, 100mM Tris-HCI (pH 9.1 at 20°C) and 0.1% Triton™ X-100. The buffer is optimized for use with 0.1-0.2mM of each dNTP.
- 10X ViBuffer S 160mM (NH₄)₂SO₄, 500mM Tris-HCI (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% Triton™ X-100. The buffer is optimized for use with 0.35mM of each dNTP.
- 50mM MgCl₂

Quality Control

All preparations are assayed for contaminating endonuclease, 3'-exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

20mM Tris-HCI (pH 8.0 at 22°C). 100mM KCl. 0.5% Tween™ 20. 0.5% Nonidet-P40, 0.1mM EDTA, 1mM DTT, and 50% glycerol. Store at -20°C.

Ordering Information

Catalog No	Description	Pack Size
PL4201	AtMax <i>Taq</i> DNA Polymerase	200u, 2.5u/µl
PL4202	AtMax <i>Taq</i> DNA Polymerase	500u, 2.5u/µl

^{*} Please refer to Appendix for the amplification protocol.





Non-Thermal Inactivation

Tag DNA Polymerase

Vivantis Technologies . Product Catalog Volume IV

(Thermus aquaticus)

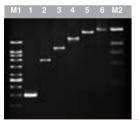


Tag DNA Polymerase is a thermostable DNA polymerase. It is suitable for applications requiring high temperature synthesis of DNA. Tag DNA Polymerase catalyzes the polymerization of nucleotides into duplex DNA in the 5' to 3' direction with the presence of Mg²⁺ and has the 5' to 3'exonuclease activity.

Features

- Ultra pure recombinant protein allows amplification up to 8kb.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon
- Generates mostly 3' dA overhang amplification products which are suitable for TA cloning.

Amplification Using Vivantis *Taq* DNA Polymerase



0.5% TAE agarose gel

Lane M1: VC 1kb DNA Ladder Lane 1: 1.5kb amplicon

Lane 2: 5.0kb amplicon Lane 3: 8.0kb amplicon Lane 4: 10kb amplicon Lane 5: 15kb amplicon

Lane 6: 20kb amplicon

Lane M2: VC Lambda / Hind III Marker

Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCI (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [3H]dTTP), 10µg activated calf thymus DNA and 100µg/ ml BSA in a final volume of 50µl.

Supplied With

- 10X ViBuffer A (without MgCl_a) 500mM KCI, 100mM Tris-HCI (pH 9.1 at 20°C) and 0.1% Triton™ X-100. The buffer is optimized for use with 0.1-0.2mM of each dNTP.
- 10X ViBuffer S 160mM (NH₄)₂SO₄, 500mM Tris-HCI (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% Triton™ X-100. The buffer is optimized for use with 0.35mM of each dNTP.
- 50mM MgCl₂

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

20mM Tris-HCI (pH 8.0 at 22°C). 100mM KCl, 0.5% Tween™ 20, 0.5% Nonidet-P40, 0.1mM EDTA, 1mM DTT and 50% glycerol. Store at -20°C.

Ordering Information

Catalog No	Description	Pack Size
PL1202 PL1204	Taq DNA Polymerase Taq DNA Polymerase	500u, 5u/µl 2 X 1000u, 5u/µl

Please refer to Appendix for the amplification protocol.

Recombinant Enzyme



Chromo Tag DNA Polymerase

(Thermus aquaticus)

Description

Chromo Tag DNA Polymerase is a thermostable DNA polymerase. It is suitable for applications requiring high temperature synthesis of DNA. Tag DNA Polymerase catalyzed the polymerization of nucleotides into duplex DNA in the 5' to 3' direction with the presence of MgCl₂ and has the 5' to 3' exonuclease activity. The enzyme is supplemented with inert color tracer dyes.

Features

- Color tracer dyes for ease of visualization of the addition of polymerase to the reaction.
- Use of 1u in every 25µl final reaction allows direct loading and serves as tracking dyes during gel electrophoresis. The blue and pink color dyes migrate approximately at 4kb and 0.3kb respectively on 1% TAE agarose
- Ultra pure recombinant protein allows amplification up to 20kb.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon.
- Generates mostly 3' dA overhang amplification products which are suitable for TA cloning.

Amplification Using Vivantis Chromo Tag DNA Polymerase



0.5 % TAE agarosa gel

Lane M1: VC 1kb DNA Ladder Lane 1: 1.5kb amplicon Lane 2: 5.0kb amplicon

Lane 3: 8.0kb amplicon

Lane 4: 10kb amplicon

Lane 5: 15kb amplicon Lane 6: 20kb amplicon

Lane M2: VC Lambda / Hind III Marker





Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCI (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [3H]dTTP), 10µg activated calf thymus DNA and 100 µg/ml BSA in a final volume of 50µl.

Supplied With

- 10X ViBuffer A (without MgCl₂) 500mM KCI, 100mM Tris-HCI (pH 9.1 at 20°C) and 0.1% Triton™ X-100. The buffer is optimized for use with 0.1-0.2mM of each dNTP
- 10X ViBuffer S 160mM (NH₄)₂SO₄, 500mM Tris-HCI (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% Triton™ X-100. The buffer is optimized for use with 0.35mM of each dNTP.
- 50mM MgCl₂

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

20mM Tris-HCI (pH 8.0 at 22°C), 100mM KCI, 0.5% Tween™ 20, 0.5% Nonidet-P40, 0.1mM EDTA, 1mM DTT. color dves and 50% glycerol. Store at -20°C.

Catalog No	Catalog No Description	
PL1205	Chromo <i>Taq</i> DNA Polymerase	200u, 1u/µl
PL1206	Chromo <i>Taq</i> DNA Polymerase	500u, 1u/µl

^{*} Please refer to Appendix for the amplification protocol.







AtTaq DNA Polymerase (Hot Start)

Vivantis Technologies . Product Catalog Volume IV

Description

At *Taq* DNA Polymerase is a complex of specific anti-*Taq* monoclonal antibody with top quality thermostable *Taq* DNA Polymerase for automatic "Hot Start" amplification, resulting in greatly enhanced amplification specificity, sensitivity and yield. At *Taq* DNA Polymerase catalyzes the polymerization of nucleotides into duplex DNA in the 5' to 3' direction with the presence of Mg²⁺ and has the 5' to 3' exonuclease activity.

Features

- Ultra pure recombinant protein which is reversibly complex with anti-Taq monoclonal antibody that blocks replication activity of the enzyme at moderate temperatures.
- Carefully selected anti-Taq antibodies have high thermal stability, providing protection against non-specific primer extension from room temperature to 70°C.
- Formation of complexes between *Taq* DNA Polymerase and an anti-*Taq* antibody forms a basis for automatic "Hot Start" amplification, which allows for the assembly of amplification reactions at room temperature.
- High stability of the complexes allows enormous increase in amplification specificity, sensitivity and yield in comparison to the conventional amplification assembly method.
- Increased specificity as a result of reduced amplification artefacts such as primer-dimer formation and mispriming in multiplex amplification.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon.

Amplification Using Vivantis AtTaq DNA Polymerase



VC 1kb DNA Ladder 1.5kb amplicon 5kb amplicon 8kb amplicon 15kb amplicon VC Lamba / *Hin*d III Marker

0.7% TAE agarose gel

Unit Definition

1u is defined as the amount of enzyme that is required to catalyze the incorporation of 10nmol of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50mM Tris-HCl (pH 9.0 at 25°C), 50mM NaCl, 5mM MgCl₂, 200µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [³H]dTTP), 10µg activated calf thymus DNA and 100µg/ml BSA in a final volume of 50µl.

Supplied With

- 10X ViBuffer A (without MgCl₂)
 500mM KCl, 100mM Tris-HCl (pH
 9.1 at 20°C) and 0.1% Triton™ X 100. The buffer is optimized for use
 with 0.1-0.2mM of each dNTP.
- 10X ViBuffer S
 160mM (NH₄)₂SO₄ , 500mM Tris-HCI (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% Triton™ X-100. The buffer is optimized for use with 0.35mM of each dNTP.
- 50mM MgCl₂

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage Buffer

20mM Tris-HCI (pH 8.0 at 22°C), 100mM KCI, 0.5% Tween™ 20, 0.5% Nonidet-P40, 0.1mM EDTA, 1mM DTT, and 50% glycerol. Store at -20°C.

Ordering Information

Cata	alog No	Description	Pack Size
PL32		At <i>Taq</i> DNA Polymerase At <i>Taq</i> DNA Polymerase	200u, 5u/µl 500u, 5u/µl

^{*} Please refer to Appendix for the amplification protocol.

Recombinant Enzyme



Chromo AtTaq DNA Polymerase (Hot Start)

Chromo

1u is defined as the amount of

the incorporation of 10nmol of

enzyme that is required to catalyze

dNTP into acid-insoluble material

in 30 minutes at 74°C. The reaction

conditions are: 50mM Tris-HCI (pH

MgCl₂, 200µM each of dATP, dCTP,

and [3H]dTTP), 10µg activated calf

thymus DNA and 100µg/ml BSA in a

• 10X ViBuffer A (without MgCl₂)

500mM KCl, 100mM Tris-HCl (pH

X-100. The buffer is optimized for

use with 0.1-0.2mM of each dNTP.

160mM (NH₄)₂ SO₄, 500mM Tris-

HCI (pH 9.2 at 22°C), 17.5mM

MgCl₂ and 0.1% Triton™ X-100.

The buffer is optimized for use

with 0.35mM of each dNTP.

All preparations are assayed for

contaminating endonuclease,

exonuclease, and non-specific

DNase activities. Functionally tested

20mM Tris-HCI (pH 8.0 at 22°C),

Nonidet-P40, 0.1mM EDTA, 1mM

DTT, color dves and 50% glycerol.

100mM KCl, 0.5% Tween™ 20, 0.5%

9.1 at 20°C) and 0.1% Triton™

9.0 at 25°C), 50mM NaCl, 5mM

dGTP, dTTP (a mix of unlabeled

final volume of 50µl.

Supplied With

10X ViBuffer S

50mM MgCl₂

in DNA amplification.

Quality Control

Storage Buffer

Store at -20°C.

Unit Definition





Description

Chromo At *Taq* DNA Polymerase is a complex of specific anti-*Taq* monoclonal antibody with top quality thermostable *Taq* DNA Polymerase for automatic "Hot Start" amplification, resulting in greatly enhanced amplification specificity, sensitivity and yield. At *Taq* DNA Polymerase catalyzes the polymerization of nucleotides into duplex DNA in the 5' to 3' direction with the presence of Mg²+ and has the 5' to 3' exonuclease activity. The enzyme is supplemented with inert color tracer dyes.

Features

- Color tracer dyes for ease of visualization of the addition of polymerase to the reaction.
- Use of 1u in every 25µl final reaction allows direct loading and serves as tracking dyes during gel electrophoresis. The blue and pink color dyes migrate approximately at 4kb and 0.3kb respectively on 1% TAE agarose gel.
- Ultra pure recombinant protein which is reversibly complex with anti-Taq monoclonal antibody that blocks replication activity of the enzyme at moderate temperatures.
- Carefully selected anti-Taq antibodies have high thermal stability, providing protection against non-specific primer extension from room temperature to 70°C.
- Formation of complexes between Taq DNA Polymerase and an anti-Taq antibody forms a basis for automatic "Hot Start" amplification, which allows for the assembly of amplification reactions at room temperature.
- High stability of the complexes allows for the enormous increase in amplification specificity, sensitivity and yield in comparison to the conventional amplification assembly method.
- Increased specificity as a result of reduced amplification artefacts such as primer-dimer formation and mispriming in multiplex amplification.
- 10X ViBuffer S provided for amplification of more than 5kb amplicon.

Amplification Using Vivantis Chromo AtTaq DNA Polymerase



Lane M1: VC 1kb DNA Ladder

Lane 1: 1.5kb amplicon
Lane 2: 5kb amplicon

Lane 3: 8kb amplicon
Lane 4: 15kb amplicon

Lane 4: 15kb amplicon

Lane 5: VC Lamba / Hind III Marker

0.7% TAE agarose gel

Ordering Information

Catalog No	Description	Pack Size
PL3205	Chromo At <i>Taq</i> DNA Polymerase	200u, 1u/µl
PL3206	Chromo At <i>Taq</i> DNA Polymerase	500u, 1u/µl

^{*} Please refer to Appendix for the amplification protocol.







Supplemented with indicators

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2X Ampli - Optimization Kit



Description

The 2X Ampli-Optimization Kit is designed to provide an easy and convenient way to optimize reaction conditions specific for your amplicon. This kit contains the 2X Tag Master Mix with a range of MgCl₂ concentrations (2.0-7.0mM). Other reaction components included in 2X Tag Master Mix such as dNTPs, reaction buffers and *Tag* DNA Polymerase are provided at the optimized concentration. This kit enables the user to determine the specific MgCl₂ concentration for their amplification reaction.

Features

- Easy and convenient for optimization of MgCl₂ concentration.
- Saves time and reduces contamination due to reduced number of pipetting steps.
- Color caps are provided for convenient recognition of MgCl₂ concentration within the 2X Tag Master Mix.
- Stable at 4°C for 6 months, allowing immediate reaction setup without the time-consuming thawing of reagents.
- Reaction buffer, dNTPs and Taq DNA Polymerase are provided at an optimized concentration.
- Suitable for all routine DNA amplification applications.

Composition

Taq DNA Polymerase (0.05u/μl), 2X ViBuffer A (100mM KCl, 20mM Tris-HCI (pH9.1 at 20°C) and 0.02% Triton™ X-100), 0.4mM dNTPs and 2.0-7.0 mM MgCl₂.

Quality Control

All preparations are assayed for contamination endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage & Stability

- Stable at -20°C for one year or at 4°C for 6 months if properly stored
- Stable for 20 freeze-thaw cycles. To avoid frequent freeze-thaw, keeping small aliquots at -20°C is recommended.
- For daily use, keeping an aliquot at 4°C is recommended.

Ordering Information

Catalog No	Description	Pack Size
PLAO01	2X Ampli-Optimization Kit	150 applications

^{*} Please refer to Appendix for the amplification protocol.

Recombinant Enzyme



2X AtTag Master Mix (Hot Start)

Description

2X At Tag Master Mix is an optimized ready-to-use 2X concentrated DNA ampli cation mixture containing AtTag DNA Polymerase, reaction buffer, dNTPs and MgCl2. It contains all the components required for routine DNA ampli cation, except template and primers. AtTag DNA polymerase is a complex of specific anti-Tag monoclonal antibody with top quality thermostable Tag DNA Polymerase for automatic "Hot Start" amplification, resulting in greatly improved amplication specificity, sensitivity and yield.

Features

- Saves time and reduces contamination due to reduced number of pipetting steps.
- Stable at 4°C for 6 months, allowing immediate reaction setup without the time consuming thawing of reagents.
- Suitable for all routine DNA ampli cation applications

Amplification of 1.5kb DNA fragment from pTZ region using Vivantis 2X AtTag Master Mix



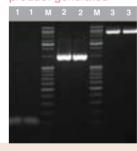
Lane M: VC 1kb DNA Ladder

Lane 1: DNA amplification product generated with 1.25u of At Taq DNA Polymerase

Lane 2: DNA amplification product generated

Lane 3: DNA amplification product generated thaw cycles)

Efficiency analysis of Vivantis 2X AtTag Master Mix - minimum and maximum base pair size of PCR product generated



Lane M: VC DNA Ladder Mix

Lane 10:DNA amplification 100bp product generated with 2X At Tag Master Mix

Lane 15: DNA amplification 1.5kb product generated with 2X At Tag Master Mix

Lane 20:DNA amplification 5kb product generated with 2X At Tag Master Mix

Composition

At Tag DNA Polymerase (0.05u/ μI), 2X ViBuffer A (100mM KCl, 20mM Tris-HCI (pH9.1 at 20°C) and (0.02% Triton™ X-100), 0.4mM dNTPs and 3.0mM MgCl₂.

Supplied With

- 50mM MqCl₂
- Nuclease-free Water

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage & Stability

- Stable at -20°C for one year or at 4°C for 6 months if properly stored.
- Stable for 20 freeze-thaw cycles. To avoid frequent freeze-thaw, keeping small aliquots at -20°C is recommended,
- For daily use, keeping aliquots at 4°C is recommended.

with 2X Taq Master Mix (stored at -20°C)

with 2X At Taq Master Mix (after 20 freeze-

Catalog No	Description	Pack Size
PLMM02	2X At Taq Master Mix	100 applications

^{*} Please refer to Appendix for the amplification protocol.





2X Tag Master Mix



Description

2X *Tag* Master Mix is an optimized ready-to-use 2X concentrated DNA amplification mixture containing *Tag* DNA polymerase, reaction buffer, dNTPs and MgCl₂. It contains all the components HCl(pH9.1 at 20°C) and (0.02% required for routine DNA amplification, except template and primers.

Features

- Saves time and reduces contamination due to reduced number of pipetting
- Stable at 4°C for 6 months, allowing immediate reaction setup without the time consuming thawing of reagents.
- Suitable for all routine DNA amplification applications.

Amplification of 5kb DNA fragment from lambda DNA using VIVANTIS 2X Tag Master Mix



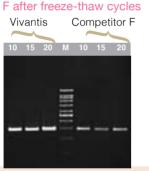
Lane M: VC 1kb DNA Ladder

Lane 1: DNA amplification product generated with 1.25u of Tag DNA

Lane 2: DNA amplification product generated with 2X Taq Master Mix (stored at -20°C)

Lane 3: DNA amplification product generated with 2X Taq Master Mix (after 20 freeze-thaw cycles)

Efficiency analysis between Vivantis 2X Tag Master Mix with competitor



Lane M: VC 1kb DNA Ladder

Lane 10: DNA amplification product generated with 2X Tag Master Mix (10 freeze-thaw cycles)

Lane 15: DNA amplification product generated with 2X Taq Master Mix (15 freeze-thaw cycles)

Lane 20: DNA amplification product generated with 2X Taq Master Mix (20 freeze-thaw cycles)

Tag DNA Polymerase (0.05u/µl), 2X ViBuffer A (100mM KCl, 20mM Tris-Triton™ X-100), 0.4mM dNTPs and 3.0mM MgCl₂.

Supplied With

- 50mM MaCl₂
- Nuclease-free Water

Quality Control

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

Storage & Stability

- Stable at -20°C for one year or at 4°C for 6 months if properly stored.
- Stable for 20 freeze-thaw cycles. To avoid frequent freeze-thaw, keeping small aliquots at -20°C is recommended,
- · For daily use, keeping aliquots at 4°C is recommended.

Ordering Information

Catalog No	Description	Pack Size
PLMM01	2X Taq Master Mix	100 applications

^{*} Please refer to Appendix for the amplification protocol.

Recombinant Enzyme



Selection Chart

Catalog No / Pack size

Catalog No / Pack size

DNA Polymerases Selection Chart

Divit of includes colorion chart					
Properties	<i>Taq</i> DNA Polymerase	Max <i>Taq</i> DNA Polymerase	At <i>Taq</i> DNA Polymerase	AtMax <i>Taq</i> DNA Polymerase	<i>Pfu</i> DNA Polymerase
Half Life	50 cycles	> 50 cycles	50 cycles	> 50 cycles	50 cycles
Target Length	Up to 8kb	Up to 40kb	Up to 15kb	Up to 20kb	Up to 8kb
Error Rate	1-2 x 10 ⁻⁵	1 x 10 ⁻⁶	1-2 x 10 ⁻⁵	1 x 10 ⁻⁶	5 x 10 ⁻⁶
Units / 50µl Reaction	2.0U	0.5-2.0U	2.0U	0.5-2.0U	0.5-1.0U
Hot Start			Yes	Yes	
Proofreading Activity		Yes		Yes	Yes
Fidelity vs Taq	1X	8-10X	1X	8-10X	2-3X
PCR Product End	3'A	Blunt / 3'A	3'A	Blunt / 3'A	Blunt
High Yield		Yes		Yes	
High Fidelity		Yes		Yes	Yes
High Throughput				Yes	
		Applications			
Routine PCR	Yes	Yes	Yes	Yes	Yes
Long PCR		Yes		Yes	
Colony PCR	Yes		Yes		
TA Cloning	Yes		Yes		
GC-rich Targets	Yes	Yes		Yes	
Long Amplicon		Yes		Yes	
DNA-labeling			Yes		
Palindrome / Multiple Repea	ts	Yes		Yes	
Multiplex Amplification		Yes	Yes	Yes	
		O	nformation		

PL1202 – 500u	PL2201 – 200u	PL3201 – 200u	PL4201 – 200u	PL5201- 100u
PL1204 – 2 x 1000u	PL2202 – 500u	PL3202 – 500u	PL4202 – 500u	PL5202 – 500u

Chromo DNA Polymerases Selection Chart

Ordering information					
Catalog No / Pack size	PL1205 – 200u	PL2205 – 200u	PL3205 – 200u	PL5205 – 100u	
Catalog No / Pack size	PL1206 – 500u	PL2206 – 500u	PL3206 – 500u	PL5206 – 500u	

My PCR Kits

Description

My PCR Kits contain all necessary components for PCR. It comes in three packaging with different combinations of reagents to meet various needs while providing convenience in many ways. My PCR Kits are an ideal solution to minimize risk of contamination when sharing reagents among different users. All My PCR Kits come with Vivantis ready-to-use VC 100bp Plus DNA Ladder pre-mixed with loading dye for 50 applications.

Kit Components

Mv PCR Kit 1

- 100 app of 2X *Taq* Master Mix (with Taq DNA Polymerase [0.05µl/ul], 0.4mM, dNTPs, 3.0mM MgCl2, 2X ViBuffer A)
- 1ml of 50mM MqCl2
- 1ml of 6X Loading Dye
- 3ml of Nuclease-Free Water
- 25µg of 100bp Plus DNA Ladder

My PCR Kit 2

- 200u of Chromo Taq DNA Plymerase
- 2ml of 10X ViBuffer A & 1ml of 10X ViBuffer S
- 1ml of 50mM MgCl2
- 1ml of 2mM dNTPs Mix
- 1ml of 6X Loading Dye
- 25µg of 100bp Plus DNA Ladder

My PCR Kit 3

- 200u of Taq DNA Plymerase
- 2ml of 10X ViBuffer A &
- 1ml of 10X ViBuffer S
- 1ml of 50mM MgCl2
- 1ml of 2mM dNTPs Mix
- 1ml of 6X Loading Dye
- 25µg of 100bp Plus DNA Ladder

Ordering Information

Catalog No	Description	Pack Size
PL8881 PL8882 PL8883	My PCR Kit 1 My PCR Kit 2 My PCR Kit 3	100 applications 100 applications 100 applications

^{*} Please refer to Appendix for the amplification protocol.

DNA Amplification Kits

Description

DNA Amplification Kits contain high quality Taq / Chromo Taq / Max Taq/ Chromo Max Taq DNA Polymerases, nucleotides, reaction buffers, MgCl₂ and nuclease free water for DNA amplification experiments. This kit allows user to carry out DNA amplification experiments by simply providing the sample DNA and primers as the other components are supplied. Positive control Plus is also included with all DNA amplification kits. Both VC 100bp DNA Ladder and VC 1kb DNA Ladder are included.

Kit Components

- 500u Taq DNA Polymerase / 500u Chromo Taq DNA Polymerase / 500u MaxTaq DNA Polymerase / 500u Chromo MaxTaq DNA Polymerase
- 2ml of 10X ViBuffer A
- 1ml of 10X ViBuffer S
- 1ml of 50mM MgCl₂
- 1ml of 2mM dNTP mix
- 100ng of control DNA
- 25µl of each 10µM forward and reverse primers for control DNA
- 100 applications of ready-to-use VC 100bp Plus DNA Ladder
- 100 applications of ready-to-use VC 1kb DNA Ladder
- 1ml of 6X loading dye
- Store at -20°C

Catalog No	Description	Pack Size
PL1202-K PL1206-K PL2202-K PL2206-K	DNA Amplification Kit (with <i>Taq</i> DNA Polymerase) DNA Amplification Kit (with Chromo <i>Taq</i> DNA Polymerase) DNA Amplification Kit (with Max <i>Taq</i> DNA Polymerase) DNA Amplification Kit (with Chromo Max <i>Taq</i> DNA Polymerase)	200 applications 200 applications 200 applications 200 applications

^{*} Please refer to Appendix for the amplification protocol.

DNA Amplification Products

Vivantis Technologies . Product Catalog Volume IV

10X ViBuffer A

Description

General reagent used for various PCR reactions.

Composition

500mM KCI, 100mM Tris-HCI (pH 9.1 at 20°C) and 0.1% Triton™ X-100. The buffer is optimized for use with 0.1-0.2mM of each dNTP.

Quality Assurance

Functionally tested in PCR with *Taq* and *Pfu* DNA Polymerases. Store at -20°C.

Ordering Information

Catalog No	Description	Pack Size
RB0201	10X ViBuffer A	5 x 1ml

10X ViBuffer S

Description

General reagent used for long PCR amplification.

Composition

160mM (NH₄)₂SO₄, 500mM Tris-HCl (pH 9.2 at 22°C), 17.5mM MgCl₂ and 0.1% TritonTM X-100. The buffer is optimized for use with 0.35mM of each dNTP.

Quality Assurance

Functionally tested in PCR with *Taq* and *Pfu* DNA Polymerases. Store at -20°C.

Ordering Information

Catalog No	Description	Pack Size
RB0203	10X ViBuffer S	5 x 1ml

50mM MgCl₂

Description

General reagent used for various PCR reactions.

Composition

 $50 \mathrm{mM} \ \mathrm{MgCl_2}$ and $6 \mathrm{H_2O}$. The buffer is optimized for use with 0.35 mM of each dNTP.

Quality Assurance

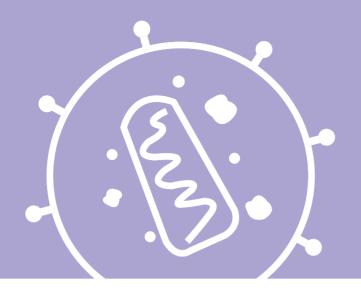
Functionally tested in PCR with *Taq* and *Pfu* DNA Polymerases. Store at -20°C.

Catalog No	Description	Pack Size
RB0204	50mM MgCl₂	5 x 1ml





RNA Amplification Products



RNA Amplification Products

2X OneStep *Taq* ReverseTrans PCR Master Mix
2X ViRed OneStep *Taq* ReverseTrans PCR Master Mix
Viva cDNA Synthesis Kit
Viva 2-step RT-PCR Kit

RNA Amplification Products

2X OneStep *Taq* ReverseTrans PCR Master Mix

Description

2X OneStep *Taq* ReverseTrans PCR Master Mix offers rapid and sensitive end-point detection of RNA templates in a single step. 2X OneStep *Taq* ReverseTrans PCR Master Mix is an optimized ready-to-use 2X concentrated RNA amplification mixture containing M-MuLV Reverse Transcriptase, RNase Inhibitors, *Taq* DNA Polymerase, reaction buffer and dNTPs. It contains all the components required for routine RNA amplification except template and primers. M-MuLV Reverse Transcriptase has the absence of RNase H activities that enhance the synthesis of long cDNAs and amplification of long transcripts. 2X OneStep *Taq* ReverseTrans PCR Master Mix allows cDNA synthesis and PCR to be performed using only gene-specific primers.

Features

- Saves time and reduces contamination due to reduced number of tests and pipetting steps
- Stable at 4°C for 6 months, allowing immediate reaction setup without the time-consuming thawing of reagent
- Suitable for all routine RNA amplification applications

Kit Components

- 1ml 2X OneStep Taq ReverseTrans PCR Master Mix
- · 2ml of Nuclease-free Water

2X ViRed OneStep *Taq* ReverseTrans PCR Master Mix

Description

2X ViRed OneStep Tag ReverseTrans PCR Master Mix offers rapid and sensitive end-point detection of RNA templates in a single step. 2X ViRed OneStep Tag ReverseTrans PCR Master Mix is an optimized ready-to-use 2X concentrated RNA amplification mixture containing M-MuLV Reverse Transcriptase, RNase Inhibitors, Tag DNA Polymerase, reaction buffer and dNTPs. It contains all the components required for routine RNA amplification except template and primers. M-MuLV Reverse Transcriptase has the absence of RNase H activities that enhances the synthesis of long cDNAs and amplification of long transcripts. 2X ViRed OneStep Tag ReverseTrans PCR Master Mix allows one-step RT-PCR using only gene-specific primers. 2X ViRed OneStep Tag ReverseTrans PCR Master Mix contains the inert red dye and stabilizers that allow direct loading of final PCR products onto gels for electrophoresis. The red color dve migrates at approximately 400bp on 1% agarose gel in 1X TBE Buffer.

Features

- Suitable for all routine RNA amplification applications
- Reduces set-up time and buffer-dye mixing
- Minimizes potential contamination due to reduced number of tests and pipetting steps
- Easy confirmation of complete mixing
- No additional loading dye needed direct loading of final products onto gels

Kit Components

- 1ml 2X ViRed OneStep *Taq* ReverseTrans PCR Master Mix
- 2ml of Nuclease-free Water

Viva 2-Steps RT-PCR Kits

Description

Viva 2-Steps RT-PCR Kits are specially designed to provide reliable synthesis of full-length cDNA and convenient application of cDNA in PCR. M-MuLV Reverse Transcriptase synthesize complementary DNA strand initiating from a specific primer, oligo d(T) or random hexamer. The absence of RNase H enhances the synthesis of long cDNA as RNA strand does not degraded in DNA-RNA hybrid during first strand cDNA synthesis. With variety of kit options for standard PCR and long PCR, Viva 2-Steps RT-PCR Kits provide flexibility in an easy use format.

Features

- Absence of RNase H activity allows high amount of full length cDNA synthesis with RNA templates up to 10kb.
- Wide selection of primers, oligo d(T) or random hexamer. *Taq* DNA Polymerase and Max*Taq* DNA Polymerase for ampli cation for short and long DNA fragments.

Kit Components

Reverse Transcription

- 10X M-MuLV RT Buffer, 250µl
- 10,000u M-MuLV Reverse Transcriptase
- 10mM dNTPs mix, 0.25ml
- RNase-free water 3ml
- 40μM Oligo d(T), 100μl
- 50ng/µl Random hexamer, 100µl
- Store at -20°C

PCR

- 500u Taq DNA Polymerase / 500u Chromo Taq DNA Polymerase / 500u MaxTaq DNA Polymerase / 500u Chromo MaxTaq DNA Polymerase
- 10X PCR Reaction Buffers
- 50mM MqCl₂
- 10mM dNTPs Mix, 0.25ml
- Store at -20°C

Viva cDNA Synthesis Kit

Description

Viva cDNA Synthesis Kit is specially designed to provide reliable synthesis of full-length cDNA. M-MuLV RNase H-synthesizes complementary DNA strand initiating from a specific primer, oligo d(T) or random hexamer. The absence of RNase H enhances the synthesis of long cDNA as the RNA strand does not degrade in DNA-RNA hybrid during first strand cDNA synthesis. This cDNA synthesis kit is readily compatible with various cDNA-dependent downstream applications.

Features

- Absence of RNase H activity allows high yield of full length cDNA synthesis with RNA templates up to 10kb
- Wide selection of primers, oligo d(T) or random hexamer
- Highly compatible with various downstream applications
- Allows synthesis of full length cDNA from various RNA templates
- High capacity and able to copy up to 2µg of purified mRNA

Kit Components

Reverse Transcription

- M-MuLV Reverse Transcriptase
- 10X Buffer M-MuLV
- 10mM dNTPs mix
- Oligo d(T)18 (40μm)
- Random hexamer (50ng/µl)
 - Traindom moxamor (don
- Nuclease-free water

Catalog No	Description	Pack Size
RTPL12 RTPL16 RTPL22 RTPL26	Viva 2-Steps RT-PCR kit with M-MuLV RT/ Taq DNA Polymerase Viva 2-Steps RT-PCR Kit with M-MuLV RT/Chromo Taq DNA Polymerase Viva 2-Steps RT-PCR Kit with M-MuLV RT/MaxTaq DNA Polymerase Viva 2-Steps RT-PCR Kit with M-MuLV RT/Chromo MaxTaq DNA Polymerase	100 applications 100 applications 100 applications 100 applications
RTMM01 RTMM02	2X OneStep <i>Taq</i> ReverseTrans PCR Master Mix 2X ViRed OneStep <i>Taq</i> ReverseTrans PCR Master Mix	100 applications
cDSK01-050 cDSK01-100	Viva cDNA Synthesis Kit Viva cDNA Synthesis Kit	50 reactions 100 reactions

^{*} Please refer to Appendix for the amplification protocol.



GF-1 Nucleic Acid Extraction Kits

Bacterial DNA Extraction Kit
Blood DNA Extraction Kit
Tissue DNA Extraction Kit
Tissue Blood Combi DNA Extraction Kit
Plant DNA Extraction Kit
Plasmid DNA Extraction Kit

PCR Clean-up Kit Gel DNA Recovery Kit AmbiClean Kit

Forensic DNA Extraction Kit
Soil Sample DNA Extraction Kit
Food DNA Extraction Kit

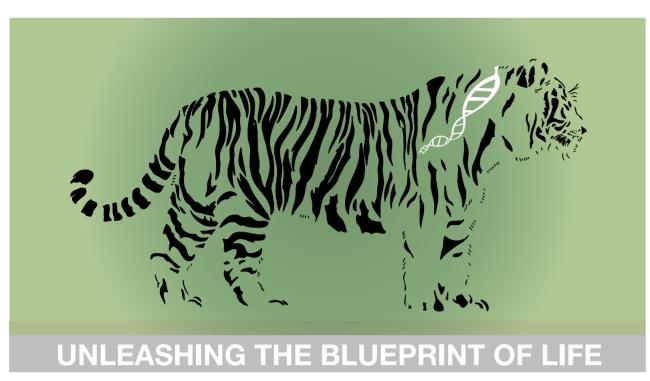
MicroTotal RNA Extraction Kit
Total RNA Extraction Kit
Blood Total RNA Extraction Kit

Viral Nucleic Acid Extraction Kit

GF-1 Starter Kits

GF-1 96-well Nucleic Acid Extraction Kits

Bacterial Genomic DNA Extraction Kit
Tissue DNA Extraction Kit
PCR Clean-up Kit
Plasmid DNA Extraction Kit
Total RNA DNA Extraction Kit



GF-1 Nucleic Acid Extraction Kits provide a rapid and efficient method for purification of nucleic acid from various samples. The purification columns in the kits are fixed with a specially treated glass filter membrane that is uniquely designed to efficiently bind nucleic acid in the presence of high salts. The kits applies the principle of a spin mini-column technology and the use of optimized buffers ensure that only DNA and / or RNA is isolated while cellular proteins, metabolites, salts and other impurities are removed during subsequent washing steps. Water or low salt buffers with the appropriated pH are then used to elute highly pure nucleic acid, ready-to-use in many routine molecular biology applications.





Features:

 Efficient High purity as intracellular proteins and nucleases are completely removed

• User-friendly Easy to use, reliable & reproducible Safe As no toxic or organic-based

extraction required

Convenient

Wide application for many routine molecular biology manipulations

Universal

Various kits for different samples Spin mini-column technology

 Base High yield

up to 20µg of DNA/RNA

Quality control:

All components of the kits are tested for the purification of DNA and RNA from different sources. The DNA and RNA obtained has been proven suitable for different downstream applications.

GF-1 Nucleic Acid Extraction Kits (Mini-Prep Kit) Unleashing the Blueprint of Life

Principle

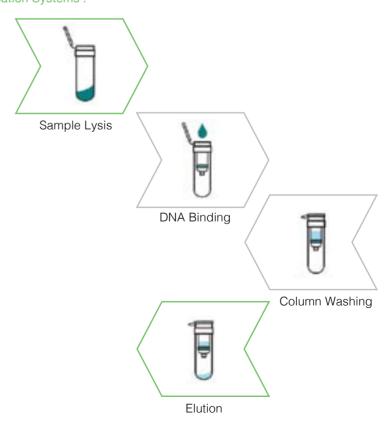
The purification columns in the kits are fixed with a specially-treated glass filter membrane that is uniquely designed to efficiently bind DNA or RNA in the presence of high salts. The kit applies the principle of a mini column spin technology and the use of optimized buffers ensure that only DNA and/or RNA is isolated while cellular proteins, metabolites. salts and other impurities are removed during subsequent washing steps. Water or low salt buffers with the appropriate pH is then used to elute highly pure DNA and/or RNA, ready to use in many routine molecular biology applications.

Quality Control

All components of the kits are tested for the purification of DNA and RNA from different sources The DNA and RNA obtained have been proven suitable for different downstream applications.

- High yield, reliable and reproducible
- High purity as intracellular proteins and nucleases are completely removed
- Fast and easy purification
- Wide application for many routine molecular biology manipulation
- Convenient for all standard operating laboratories
- No toxic or organic-based extraction required

Nucleic Acid Purification Systems:



GF-1 Bacterial DNA Extraction Kit

Description

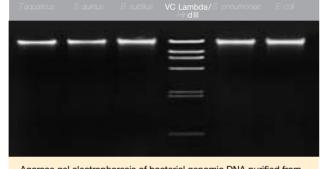
The GF-1 Bacterial DNA Extraction Kit provides a rapid and efficient method for purification of high molecular weight genomic DNA from either Gram-negative or Gram-positive bacteria. The purification is based on the usage of denaturing agents to provide lysis of cells, denaturation of proteins and subsequently release of genomic DNA. Special buffers provided in the kit are optimized to enhance the binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure genomic DNA.

Features

- Suitable for both Gram negative or Gram positive bacteria
- Yields up to 20µg of DNA
- No organic-based extraction required
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting and DNA fingerprinting.

Kit Components

- Buffer R1
- Buffer R2
- Buffer BG
- Wash Buffer (concentrate)
- Elution Buffer
- Proteinase K



Agarose gel electrophoresis of bacterial genomic DNA purified from various strains using the GF-1 Bacterial DNA Extraction Kit.

GF-1 Blood DNA Extraction Kit

Description

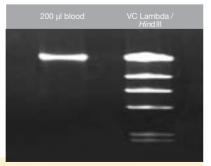
The GF-1 Blood DNA Extraction Kit is designed for rapid and efficient purification of genomic DNA from fresh and frozen anti-coagulated whole blood. The purification is based on the usage of denaturing agents to provide lysis of cells, denaturation of proteins and subsequently release of genomic DNA. Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure genomic DNA.

Features

- Yields up to 20µg of DNA
- No organic-based extraction required
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting and DNA fingerprinting.

Kit Components

- Buffer BB
- Wash Buffer 1 (concentrate)
- Wash Buffer 2 (concentrate)
- Elution Buffer
- Proteinase K



Agarose gel electrophoresis of blood genomic DNA purified from human whole blood using the GF-1 Blood DNA Extraction Kit.

GF-1 Tissue DNA Extraction Kit

Description

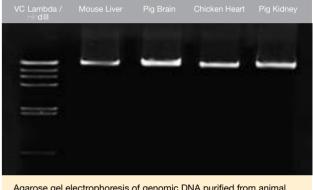
The GF-1 Tissue DNA Extraction Kit is designed for rapid and efficient purification of genomic DNA from various tissue samples such as kidney, heart, lungs, brain, muscles, liver, spleen, and animal cultured cells. The purification is based on the usage of denaturing agents to provide lysis of cells, denaturation of proteins and subsequently release of genomic DNA. Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure genomic DNA.

Kit Components

- Buffer TL
- Lysis Enhancer
- Buffer TB
- Wash Buffer (concentrate)
- Elution Buffer
- Proteinase K

Features

- Yields up to 20µg of DNA
- No organic-based extraction required
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting and DNA fingerprinting.



Agarose gel electrophoresis of genomic DNA purified from animal tissue using the GF-1 Tissue DNA Extraction Kit.

GF-1 Tissue Blood Combi DNA Extraction Kit

Description

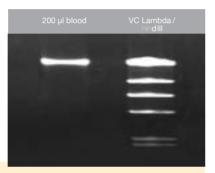
The GF-1 Tissue Blood DNA Extraction Kit is designed for rapid and efficient purification of genomic DNA from blood and various fresh or frozen tissue samples such as kidney, heart, lungs, brains, muscles, liver, spleen, and animal cultured cells. The purification is based on the usage of denaturing agents to provide lysis of cells, denaturation of proteins and subsequently release of genomic DNA.

Features

- Yields up to 20µg of DNA
- No organic-based extraction required
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting and DNA fingerprinting.

Kit Components

- Buffer TL
- Lysis Enhancer
- Buffer CB
- Wash Buffer 1 (concentrate)
- Wash Buffer 2 (concentrate)
- Elution Buffer
- Proteinase K



Agarose gel electrophoresis of blood genomic DNA purified from human whole blood using the GF-1 Blood DNA Extraction Kit.

GF-1 Plasmid DNA Extraction Kit

Description

The GF-1 Plasmid DNA Extraction Kit is designed for rapid and efficient purification of high copy and low copy plasmid DNA from bacterial lysates. The kit uses the alkaline lysis-SDS method to lyse cells and release plasmid DNA. Special buffers provided in the kit are optimized to enhance binding DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure plasmid DNA.

Kit Components

- Solution 1
- Solution 2
- Buffer NB
- Wash Buffer (concentrate)
- Elution Buffer
- RNase A

Features

- Yields up to 20µg of DNA
- Multiple samples can be processed rapidly in less than 30 minutes
- No organic-based extraction required
- Highly pure plasmid DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, ligation, DNA sequencing, transformation, etc.



Agarose gel electrophoresis of different types of plasmid DNA purified using the GF-1 Plasmid DNA Extraction Kit.

- pCambia (Cambia)
 pUC18
- pGEM-T (Promega)
- 4. pBR322

GF-1 Plant DNA Extraction Kit

Description

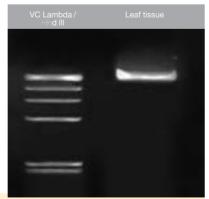
The GF-1 Plant DNA Extraction Kit is designed for rapid and efficient purification of genomic DNA from a wide variety of plant tissues. The purification is based on the usage of denaturing agents to provide lysis of tissue cells, denaturation of proteins and subsequently release of genomic DNA. Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure genomic DNA.

Features

- Yields up to 20µg of DNA
- No organic-based extraction required
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting and DNA fingerprinting.

Kit Components

- Buffer PL
- Buffer PB
- Wash Buffer (concentrate)
- Elution Buffer
- Proteinase K



Agarose gel electrophoresis of genomic DNA purified from plant tissue using the GF-1 Plant DNA Extraction Kit.

Catalog No	Description	Pack Size
GF-BA-050	GF-1 Bacterial DNA Extraction Kit	50 preps
GF-BA-100	GF-1 Bacterial DNA Extraction Kit	100 preps
GF-BD-050	GF-1 Blood DNA Extraction Kit	50 preps
GF-BD-100	GF-1 Blood DNA Extraction Kit	100 preps
GF-TD-050	GF-1 Tissue DNA Extraction Kit	50 preps
GF-TD-100	GF-1 Tissue DNA Extraction Kit	100 preps
GF-BT-050	GF-1 Tissue Blood Combi DNA Extraction Kit	50 preps
GF-BT-100	GF-1 Tissue Blood Combi DNA Extraction Kit	100 preps
GF-PT-050	GF-1 Plant DNA Extraction Kit	50 preps
GF-PT-100	GF-1 Plant DNA Extraction Kit	100 preps
GF-PL-050 GF-PL-100 GF-PL-200	GF-1 Plasmid DNA Extraction Kit GF-1 Plasmid DNA Extraction Kit GF-1 Plasmid DNA Extraction Kit	50 preps 100 preps 200 preps

GF-1 PCR Clean-up Kit

Description

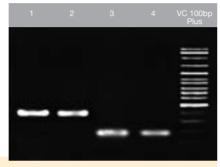
The GF-1 PCR Clean-up Kit is designed for rapid and efficient clean up of DNA ranging from 100bp to 20kb. The kit efficiently removes dNTPs, short oligo fragments, mineral oil, enzymes from a PCR reaction product, removes proteins after restriction enzyme treatment and dephosphorylation, residual dye and ethidium bromide. This kit is also suited for concentrating DNA, changing of buffers and desalting.

Features

- Indicator dye for easy pH determination
- Purification process less than 15 minutes
- Highly pure DNA ready to use for routine molecular biology applications such as restriction enzymes digestion, PCR, ligation, DNA sequencing and probe preparations.

Kit Components

- Buffer PCR
- Wash Buffer (concentrate)
- Elution Buffer



Clean up and recovery of DNA fragments from PCR reactions using the GF-1 PCR Clean-up Kit.

- 1. PCR product of 400bp before clean-up
- 2. PCR Cleaned-up product of 400bp
- 3. PCR product of 200bp before clean-up
- 4. PCR Cleaned-up product of 200bp

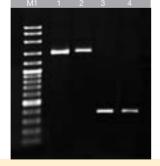
GF-1 Gel DNA Recovery Kit

Description

The GF-1 Gel DNA Recovery Kit is a system designed for rapid purification of DNA bands ranging from 100bp to 10kb from all grades of agarose gel in TAE (Tris-acetate / EDTA) or TBE (Tris-borate / EDTA). Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure DNA.

Features

- 90% of recovery achievable
- Purification process less than 15 minutes
- High pure DNA ready-to-use for routine molecular biology applications such as restriction enzyme digestion, PCR, ligation, DNA sequencing, probe preparations, southern blotting and DNA fingerprinting.



Recovery of small DNA fragments from 1.0% TBE agarose gel. M1. VC 100bp Plus DNA Ladder

- 1. DNA fragment, 1.5kb
- Recovered 1.5kb DNA fragment
- DNA fragment, 0.36kb
- 4. Recovered 0.36kb DNA fragment

Kit Components

- Buffer GB
- Wash Buffer (concentrate)
- Elution Buffer

M2 A B C D

Recovery of large DNA fragments from 0.7% TBE agarose gel. M2. VC 1kb DNA Ladder

- Skb DNA fragment, before recovery
- B. Recovered 5kb DNA fragment
- C. 10kb DNA fragment, before recovery
- D. Recovered 10kb DNA fragment

GF-1 AmbiClean Kit (Gel & PCR)

Description

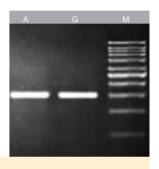
The GF-1 AmbiClean Kit (Gel & PCR) is designed for rapid DNA recovery from agarose gel and PCR clean-up of DNA bands ranging from 100bp to 20kb. Special buffer provide the correct salt concentration and pH for efficient recovery (80-90%) of DNA from both PCR product and agarose gel from TAE or TBE buffers. The kit is well suited for the removal of agarose, excess dNTPs, short oligo fragments, mineral oil, enzymes from a PCR reaction product, proteins after restriction enzyme treatment and dephosphorylation, residual dye and ethidium bromide. This kit also allows for concentration of DNA, changing of buffers and desalting.

Features

- 90% of recovery achievable
- Purification process less than 15 minutes
- High pure DNA ready-to-use for routine molecular biology applications such as restriction enzyme digestion, PCR, ligation, DNA sequencing, probe preparations, etc.

Kit Components

- Buffer DB
- Wash Buffer (concentrate)
- Elution Buffer



- M: VC 1kb DNA ladder
- A: PCR product (1.5kb) before purification
- P: Purified PCR product using GF-1 AmbiClean Kit (PCR clean-up)
- G: Purified PCR product using GF-1 AmbiClean Kit (Gel Recovery)

Catalog No	Description	Pack Size
GF-PC-050	GF-1 PCR Clean-up Kit	50 preps
GF-PC-100	GF-1 PCR Clean-up Kit	100 preps
GF-PC-200	GF-1 PCR Clean-up Kit	200 preps
GF-GP-050	GF-1 Gel DNA Recovery Kit	50 preps
GF-GP-100	GF-1 Gel DNA Recovery Kit	100 preps
GF-GP-200	GF-1 Gel DNA Recovery Kit	200 preps
GF-GC-050	GF-1 AmbiClean Kit (Gel & PCR)	50 preps
GF-GC-100	GF-1 AmbiClean Kit (Gel & PCR)	100 preps
GF-GC-200	GF-1 AmbiClean Kit (Gel & PCR)	200 preps

GF-1 Forensic DNA Extraction Kit

Description

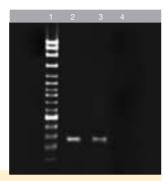
The GF-1 Forensic DNA Extraction Kit is designed for rapid and efficient purification of DNA from traces of biological materials such as blood stains, saliva, semen, hair and nail for clinical and forensic analysis. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of DNA. Special buffers provided in the kit are optimized to enhance the binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure DNA.

Features

- No organic-based extraction required.
- Highly pure DNA ready to use for routine molecular biology applications such as restriction enzymes digestion, PCR, protein-DNA interactions and blotting.

Kit Components

- Buffer STL
- Buffer BL
- HB Buffer
- Wash Buffer (concentrate)
- Elution Buffer
- OB Protease



Amplification of β-actin gene from human genomic DNA in a 50µl PCR reaction mixture. 2μl from 50μl of eluted sample is used as template in PCR. 3μl of PCR product is loaded per lane and electrophoresed in a 1.0% TBE agarose gel. Expected PCR product size amplified from human genomic DNA is 250bp.

- VC 100bp Plus DNA Ladder
- Human blood stains
- Human whole blood
- Negative control

GF-1 Soil Sample DNA Extraction Kit

Description

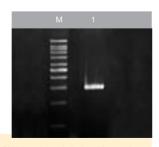
The GF-1 Soil Sample DNA Extraction Kit is designed for rapid and efficient purification of bacterial DNA from up to 1 gram soil sample. The purification is based on the usage of denaturing agents to provide efficient cell lysis, and to eliminate PCR inhibitory compounds like humid acid. Special buffers provided in the kit are optimized to enhance the binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure DNA.

Features

- No organic-based extraction required.
- Highly pure DNA ready to use for routine molecular biology applications such as PCR.

Kit Components

- Glass Beads
- HTR Reagents
- Buffer DS
- Buffer SLX MINS
- P2 Buffer
- XP1 Buffer
- SPW Wash Buffer
- Elution Buffer



Amplification of conserved region of 16S rDNA from bacteria isolated from soil sample in a 25µl reaction mixture. 2µl from 50µl of eluted sample is used as template in PCR. 3µl of PCR product is loaded per lane and electrophoresed in a 1.0% TBE agarose gel. Expected PCR product size amplified from bacteria isolated from soil is 1.5kb. M - VC 1kb DNA Ladder

1 - Soil sample

GF-1 Food DNA Extraction Kit

Description

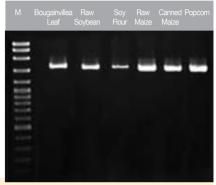
The GF-1 Food DNA Extraction Kit is designed for rapid and efficient purification of DNA from raw or processed food from plant, animal or mixed origins. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of DNA. Special buffers provided in the kit are optimized to enhance the binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure DNA.

Features

- Yields up to 20µg of DNA.
- · No organic-based extraction required.
- · Highly pure DNA ready to use for routine molecular biology applications such as PCR.

Kit Components

- Buffer FL
- Buffer FB
- Wash Buffer 1 (concentrate)
- Wash Buffer 2 (concentrate)
- Elution Buffer
- Proteinase K



Amplification of chloroplast DNA from DNA purified from soybean and maize varieties in a 25µl reaction mixture. 1µl to 4µl of 200µl eluted sample is used as template in PCR. 2µl of PCR product is loaded per lane and electrophoresed in a 1.0% TBE agarose gel. Expected PCR product size amplified from soybean and maize varieties is 1.5kb. M. VC 100bp Plus DNA Ladder

Catalog No	Description	Pack Size
GF-FD-025	GF-1 Forensic DNA Extraction Kit	25 preps
GF-SD-025	GF-1 Soil Sample DNA Extraction Kit	25 preps
GF-FE-025	GF-1 Food DNA Extraction Kit	25 preps
GF-FE-100	GF-1 Food DNA Extraction Kit (Proteinase K included)	100 preps

GF-1 microTotal RNA Extraction Kit

Description

The GF-1 microTotal RNA Extraction Kit is designed for rapid and efficient purification of RNA from various types of samples including animal cells, cultured cells (monolayer cells, suspension cells), blood. plasma, serum, buffy coat, biological fluids like saliva and semen. The purification is based on the usage of denaturing agents to provide lysis of cells, denaturation of proteins and subsequently release of RNA. Special buffers provided in the kit are optimized to enhance binding of RNA onto a specially-treated glass filter membrane for efficient recovery of highly total RNA. Simple centrifugation steps allow multiples sample processing and remove contaminants to yield highly pure total RNA which is suitable to be used in various down-stream applications.

Features

- Purify large RNA, siRNA and microRNA
- Yields up to 3µg of total RNA.
- No organic-based extraction required.
- Highly pure RNA ready to use for RT-PCR, Northern Blotting, polyA RNA (mRNA) purification, nuclease protection and in vitro translation.

Kit Components

- Buffer TS
- Wash Buffer (concentrate)
- RNase-free Water

GF-1 Total RNA Extraction Kit

Description

The GF-1 Total RNA Extraction Kit is designed for rapid and efficient purification of RNA from various types of samples including cultured animal cells, bacterial cells, animal and plant tissues, and yeast cells. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of RNA. Special buffers provided in the kit are optimized to enhance the binding of RNA onto a specially-treated glass filter membrane for efficient recovery of highly pure RNA.

Features

- Yields up to 3µg of total RNA.
- No organic-based extraction required.
- Highly pure RNA ready to use for RT-PCR, Northern Blotting, polyA RNA (mRNA) purification, nuclease protection and in vitro translation.

(concentrate) • Wash Buffer (concentrate) Digestion Buffer

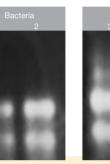
• Digestion Enhancer

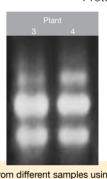
Inhibitor Remover Buffer

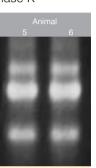
Kit Components

• Buffer TR

- RNase-free Water
- DNase I
- Proteinase K







Total RNA extracted from different samples using GF-1 Total RNA Extraction Kit. 5µl of eluted RNA is loaded per lane and electrophoresed in a 1% TBE agarose gel.

- 1. E.coli Top 10F'
- 3. Orchid leaf
- Mouse kidney
- Bacillus subtilis
- Orchid root
- Mouse liver

Catalog No	Description	Pack Size
GF-MT-025 GF-MT-050 GF-MT-100	GF-1 microTotal RNA Extraction Kit GF-1 microTotal RNA Extraction Kit GF-1 microTotal RNA Extraction Kit	25 preps 50 preps 100 preps
GF-TR-025 GF-TR-050 GF-TR-100	GF-1 Total RNA Extraction Kit GF-1 Total RNA Extraction Kit GF-1 Total RNA Extraction Kit	25 preps 50 preps 100 preps

GF-1 Blood Total RNA Extraction Kit

Description

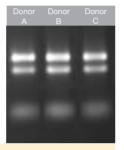
The GF-1 Blood Total RNA Extraction Kit is designed for rapid and efficient purification of total RNA from up to 1ml fresh and frozen anti-coagulated whole blood. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of RNA. Special buffers provided in the kit are optimized to enhance the binding of RNA onto a specially-treated glass filter membrane for efficient recovery of highly pure RNA.

Features

- Yields up to 3µg of total RNA.
- No organic-based extraction required.
- Highly pure total RNA, ready to use for RT-PCR, Northern Blotting, polyA RNA (mRNA) purification, nuclease protection and in vitro translation.

Kit Components

- Buffer BR
- Inhibitor Remover Buffer (concentrate)
- Wash Buffer (concentrate)
- Digestion Buffer
- Digestion Enhancer
- RNase-free Water
- DNase I
- Proteinase K



Total RNA extracted from whole blood of different donors using GF-1 Blood Total RNA Extraction Kit. 5µl of eluted RNA is loaded per lane and electrophoresed in a 1% TBE agarose gel.

GF-1 Viral Nucleic Acid Extraction Kit

Description

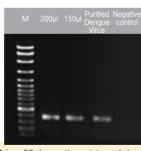
The GF-1 Viral Nucleic Acid Extraction Kit is designed for rapid and efficient purification of viral DNA/RNA from samples such as serum, plasma, body fluid or virus-infected cell culture supernatant. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of DNA or RNA. Special buffers provided in the kit are optimized to enhance the binding of DNA or RNA onto a specially-treated glass filter membrane for efficient recovery of highly pure DNA or RNA.

Features

- No organic-based extraction required.
- Highly pure DNA or RNA, ready to use for routine molecular biology applications such as PCR and RT-PCR.

Kit Components

- Buffer VL
- Wash Buffer 1 (concentrate)
- Wash Buffer 2 (concentrate)
- Elution Buffer
- Proteinase K
- Carrier RNA



RT-PCR products amplified from dengue virus-infected cell culture supernatant total RNA in a 50µl reaction mixture. 1µl of 30µl eluted sample is used as template in RT-PCR. 3µl of RT-PCR product is loaded per lane and electrophoresed in a 1.0% TBE agarose gel. Expected RT-PCR product size amplified from dengue virus total RNA is 362bp.

M. VC 100bp Plus DNA Ladder

Ordering Information

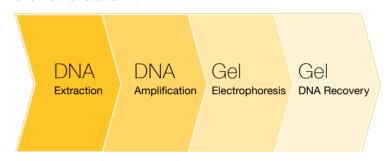
C	Catalog No	Description	Pack Size
_	GF-TB-025	GF-1 Blood Total RNA Extraction Kit	25 preps
	GF-TB-100	GF-1 Blood Total RNA Extraction Kit	100 preps
	GF-RD-025	GF-1 Viral Nucleic Acid Extraction Kit	25 preps
	GF-RD-050	GF-1 Viral Nucleic Acid Extraction Kit	50 preps
	GF-RD-100	GF-1 Viral Nucleic Acid Extraction Kit	100 preps

GF-1 Starter Kit: From Sample to Gene Isolation

Description

GF-1 Starter kit offers a comprehensive approach for the isolation of high quality DNA from a variety of sources for subsequent amplification, visualization and purification of desired genes. This kit is specially prepared and combines our state-of-the-art DNA extraction kits with *Taq* DNA Polymerase/Chromo *Taq* DNA Polymerase, DNA ladders, and gel extraction kit eliminating the need for separate orders for the completion of the experiment. Appropriate positive controls are made available to verify that correct experimental approaches have been taken throughout the whole procedure.

Overview of Starter Kit



Kit Components

- 25 preps of GF-1 DNA Extraction Kit
- 25 preps of GF-1 Gel DNA Recovery Kit
- 200u of Taq DNA Polymerase / Chromo Taq DNA Polymerase
- 1ml 10X ViBuffer A
- 1ml 10X ViBuffer S
- 1ml 50mM MgCl₂
- 0.25ml of 2mM dNTP mix
- 100ng of control DNA
- 25µl of each 10µM forward and reverse primers for control DNA
- 50 applications of readyto-use VC 100bp Plus DNA Ladder
- 50 applications of readyto-use VC 1kb DNA Ladder
- 100µl of 6X Loading dye

Catalog No	Description	Pack Size
GF-BA-K GF-BA-KW	GF-1 Bacterial Starter Kit / Taq DNA Polymerase GF-1 Bacterial Starter Kit / Chromo Taq DNA Polymerase	25 preps 25 preps
GF-BD-K GF-BD-KW GF-PT-K	GF-1 Blood Starter Kit / <i>Taq</i> DNA Polymerase GF-1 Blood Starter Kit / Chromo <i>Taq</i> DNA Polymerase GF-1 Plant Starter Kit / <i>Taq</i> DNA Polymerase	25 preps 25 preps 25 preps
GF-PT-KW GF-TD-K	GF-1 Plant Starter Kit / Chromo Taq DNA Polymerase GF-1 Tissue Starter Kit / Taq DNA Polymerase	25 preps 25 preps
GF-TD-KW GF-PL-K GF-PL-KW	GF-1 Tissue Starter Kit / Chromo <i>Taq</i> DNA Polymerase GF-1 Plasmid Starter Kit / <i>Taq</i> DNA Polymerase GF-1 Plasmid Starter Kit / Chromo <i>Taq</i> DNA Polymerase	25 preps 25 preps 25 preps
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GF-1 96-well Nucleic Acid Extraction Kit

GF-1 96-well Bacterial Genomic DNA Extraction Kit

Description

The GF-1 96-well Bacterial Genomic DNA Extraction Kit is designed for rapid and high-throughput purification of bacterial genomic DNA, up to 96 samples simultaneously. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of genomic DNA. Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure genomic DNA.

Features

- Yields up to 5-10µg of genomic DNA.
- Purification process takes less than 30 minutes.
- No organic-based extraction required.
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting, DNA fingerprinting, etc.

GF-1 96-well Tissue DNA Extraction Kit

Description

The GF-1 96-well Tissue DNA Extraction Kit is designed for rapid and high-throughput purification of animal tissues, cultured animal cells and paraffin-embedded tisues, up to 96 samples simultaneously. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of genomic DNA. Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure genomic DNA.

Features

- Yields up to 5-10µg of genomic DNA.
- No organic-based extraction required.
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern Blotting, DNA fingerprinting, etc.

GF-1 96-well PCR Clean-up Kit

Description

The GF-1 96-well PCR Clean-up Kit is designed for rapid and high-throughput purification of DNA ranging from 100bp to 20kb, up to 96 samples simultaneously. The kit efficiently removes dNTPs, short oligo fragments, mineral oil, enzymes from a PCR reaction product, proteins after restriction enzyme treatment and dephosphorylation, residue of dye and ethidium bromide. The kit is also suitable for concentrating DNA, changing of buffers and desalting.

Features

- Up to 90% recovery of DNA.
- Puri cation process of about 30 minutes.
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern Blotting, DNA ngerprinting, etc.

GF-1 96-well Nucleic Acid Extraction Kit (cont'd)

GF-1 96-well Plasmid DNA Extraction Kit

Description

The GF-1 96-well Plasmid DNA Extraction Kit is designed for rapid and high-throughput purification of high copy number and low copy number plasmid DNA from 1-2ml of bacteria culture, up to 96 samples simultaneously. The kit uses alkaline lysis-SDS method to lyse cells and release plasmid DNA. Special buffers provided in the kit are optimized to enhance binding of plasmid DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure plasmid DNA.

Features

- Yields up to 5-10µg and 0.5-5µg of DNA for high copy number and low copy number plasmid, respectively.
- Purification process takes less than 60 minutes.
- No organic-based extraction required.
- Highly pure genomic DNA ready to use for routine molecular biology applications such as restriction enzyme digestion, PCR, Southern blotting, DNA fingerprinting, etc.

GF-1 96-well Total RNA Extraction Kit

Description

The GF-1 96-well Total RNA Extraction Kit is designed for rapid and high-throughput purification of total RNA from bacterial cultures, viruses, plant and animal tissues, up to 96 samples simultaneously. The purification is based on the usage of denaturing agents to provide efficient cell lysis, denaturation of proteins and subsequent release of RNA. Special buffers provided in the kit are optimized to enhance binding of DNA onto a specially-treated glass filter membrane for efficient recovery of highly pure total RNA.

Features

- Purification process takes less than 60 minutes.
- No organic-based extraction required.
- Highly pure total RNA ready to use for routine molecular biology applications such as cDNA synthesis, RT-PCR, etc.

Catalog No	Description	Pack Size
GF-96-G05 GF-96-G10	GF-1 96-well Bacterial Genomic DNA Extraction Kit GF-1 96-well Bacterial Genomic DNA Extraction Kit	96 x 5 plates 96 x 10 plates
GF-96-T05 GF-96-T10	GF-1 96-well Tissue DNA Extraction Kit GF-1 96-well Tissue DNA Extraction Kit	96 x 5 plates 96 x 10 plates
GF-96-C05 GF-96-C10	GF-1 96-well PCR Clean-up Kit GF-1 96-well PCR Clean-up Kit	96 x 5 plates 96 x 10 plates
GF-96-P05 GF-96-P10	GF-1 96-well Plasmid DNA Extraction Kit GF-1 96-well Plasmid DNA Extraction Kit	96 x 5 plates 96 x 10 plates
GF-96-R05 GF-96-R10	GF-1 96-well Total RNA Extraction Kit GF-1 96-well Total RNA Extraction Kit	96 x 5 plates 96 x 10 plates



& Markers / Nucleic Acids / Nucleotides

Ladders & Markers / Nucleic Acids / Nucleotides

DNA Ladders & Markers

Protein Ladders & Markers

Nucleic Acids

Nucleotides

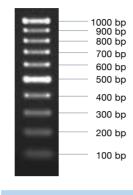
VC 100bp DNA Ladder

Vivantis Technologies . Product Catalog Volume IV

Description

Serves as molecular weight standards for electrophoresis for both agarose and polyacrylamide gels. Suitable for sizing of PCR products or other double-stranded DNA fragments. Fragments with size 500bp, and 1000bp are higher in intensity in comparison to other bands to serve as orientation points.

Storage Buffer 10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol. Usage Recommendation
Use 0.05-0.10µg of the
DNA ladder per 1mm
width of gel lane.



1.0% Agarose in 1X TBE

VC 100bp Plus DNA Ladder

Description

Serves as molecular weight standards for electrophoresis for both agarose and polyacrylamide gel electrophoresis. Suitable for sizing of PCR products or other double-stranded DNA fragments. Fragments with size 500bp, and 1000bp are higher in intensity in comparison to other bands to serve as orientation points.

Storage Buffer

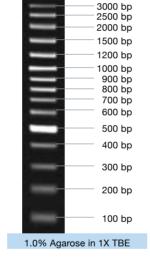
10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

6X Loading Dye Solution

10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol.

Usage Recommendation

Use 0.05-0.10µg of the DNA ladder per 1mm width of gel lane.



Ordering Information

Catalog No	Description	Pack Size
NL1401	VC100bp DNA Ladder	50 μg
NL1402	VC100bp DNA Ladder	5 x 50 μg
NL1403	VC100bp DNA Ladder (ready-to-use)	50 μg
NL1404	VC100bp DNA Ladder (ready-to-use)	5 x 50 μg
NL1405	VC 100bp Plus DNA Ladder	50 μg
NL1406	VC 100bp Plus DNA Ladder	5 x 50 μg
NL1407	VC 100bp Plus DNA Ladder (ready-to-use)	50 μg
NL1408	VC 100bp Plus DNA Ladder (ready-to-use)	5 x 50 μg

VC 1kb DNA Ladder

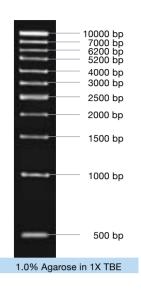
Description

Serves as molecular weight standards for agarose gel electrophoresis. Suitable for sizing of PCR products or other double-stranded DNA fragments. Fragments with size 2500bp are higher in intensity in comparison to other bands to serve as orientation points.

Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol. Usage Recommendation Use 0.05-0.10µg of the DNA ladder per 1mm

DNA ladder per 1mm width of gel lane.



VC 1kb-Ex DNA Ladder

Description

Serves as molecular weight standards for agarose gel electrophoresis. Suitable for sizing of PCR products or other double-stranded DNA fragments. Fragments with size 2500bp are higher in intensity in comparison to other bands to serve as orientation point.

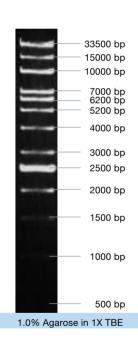
Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03%

bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol

Usage Recommendation Use 0.05-0.10µg of the

DNA ladder per 1mm width of gel lane



Catalog No	Description	Pack Size
NL1409 NL1410 NL1411 NL1412	VC 1kb DNA Ladder VC 1kb DNA Ladder VC 1kb DNA Ladder (ready-to-use) VC 1kb DNA Ladder (ready-to-use)	50 μg 5 x 50 μg 50 μg 5 x 50 μg
NL1413 NL1414 NL1415 NL1416	VC 1kb-Ex DNA Ladder VC 1kb-Ex DNA Ladder VC 1kb-Ex DNA Ladder (ready to use) VC 1kb-Ex DNA Ladder (ready to use)	50 μg 5 x 50 μg 50 μg 5 x 50 μg

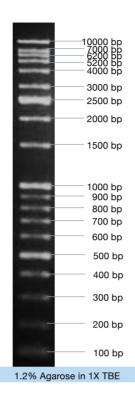
VC DNA Ladder Mix

Vivantis Technologies . Product Catalog Volume IV

Description

Serves as molecular weight standards for electrophoresis for both agarose and polyacrylamide gels. Suitable for sizing of PCR products or other double-stranded DNA fragments. Fragments with size 500bp, 1000bp, 2500bp are higher in intensity in comparison to other bands to serve as orientation points.

Storage Buffer 10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol Usage Recommendation
Use 0.05-0.10µg of the
DNA ladder per 1mm
width of gel lane



Ordering Information

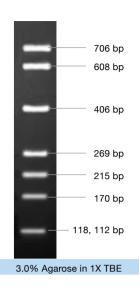
Catalog No	Description	Pack Size
NL1417	VC DNA Ladder Mix (ready-to-use)	50 μg
NL1418	VC DNA Ladder Mix (ready-to-use)	5 x 50 μg
NL1419	VC DNA Ladder Mix (ready-to-use)	50 μg
NL1420	VC DNA Ladder Mix (ready-to-use)	5 x 50 μg

CentiMark PCR Marker

Description

PCR marker is suitable for sizing linear double-stranded DNA fragment or PCR products in agarose. Plasmid is completely digested and yields 11 fragments in size (in bp): 706, 608, 406, 269, 215, 170, 118, 112, 43, 26, 13 (obscure bands).

Storage Buffer 10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C. 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol. Usage Recommendation
Use 0.05-0.10µg of the
DNA ladder per 1mm
width of gel lane.



MilliMark PCR Marker

Description

PCR marker is suitable for sizing linear double-stranded DNA fragment or PCR products in agarose. Plasmid is completely digested and yields 8 fragments in size (in bp): 1161, 943, 718, 585, 497, 341, 267, 225, 153, 105, 85, 78, 75, 46, 36, 18, 17, 12, 11, 8 (obscure bands).

Storage Buffer

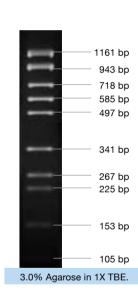
10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

6X Loading Dye Solution 10mM Tris-HCI (pH 8.0),

60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol.

Usage Recommendation

Use 0.05-0.10µg of the DNA ladder per 1mm width of gel lane.



	Catalog No	Description	Pack Size
-	NM2417 NM2418 NM2419 NM2420	CentiMark PCR Marker CentiMark PCR Marker CentiMark PCR Marker (ready-to-use) CentiMark PCR Marker (ready-to-use)	50 µg 5 x 50 µg 50 µg 5 x 50 µg
	NM2421 NM2422 NM2423 NM2424	MilliMark PCR Marker MilliMark PCR Marker MilliMark PCR Marker (ready-to-use) MilliMark PCR Marker (ready-to-use)	50 μg 5 x 50 μg 50 μg 5 x 50 μg

VC Lambda/BssT1l Marker

Description

This lambda marker is suitable for sizing linear double-stranded DNA fragments in agarose. The lambda DNA is completely digested with BssT1l and yields 11 fragments in size (in bp): 19329, 7743, 6223, 4254, 3472, 2690, 1882, 1489, 925, 421, 74 (obscure band).

Storage Buffer

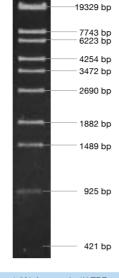
10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

6X Loading Dye Solution

10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol

Usage Recommendation

Use 0.05-0.10µg of the DNA ladder per 1mm width of gel lane



1.0% Agarose in 1X TBE

Ordering Information

Description	Pack Size
VC Lambda/ <i>Bss</i> T1I Marker	50 μg
VC Lambda/BssT1I Marker	5 x 50 μg
VC Lambda/BssT1I Marker (ready to use)	50 μg
VC Lambda/BssT1I Marker (ready to use)	5 x 50 μg
	VC Lambda/ <i>Bss</i> T1I Marker (ready to use)

VC Lambda / EcoRI Marker

Description

This lambda marker is suitable for sizing linear double-stranded DNA fragments in agarose gel. The lambda DNA is completely digested with *EcoR* I and yields 6 fragments (in bp): 21226, 7421, 5804, 5643, 4878, 3530.

Storage Buffer 10mM Tris-HCl (pH 8.0) and 1mM EDTA. Store 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol. Usage Recommendation Use 0.05-0.10µg of the DNA ladder per 1mm width of gel lane.

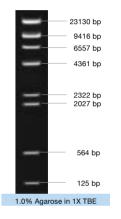
VC Lambda / Hind III Marker

Description

at -20°C.

This lambda marker is suitable for sizing linear double-stranded DNA fragments in agarose gel. The lambda DNA is completely digested with *Hind* III and yields 8 fragments (in bp): 23130, 9416, 6557, 4361, 2322, 2027, 564, 125.

Storage Buffer 10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C. 6X Loading Dye Solution 10mM Tris-HCl (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol. Usage Recommendation Use 0.05-0.10µg of the DNA ladder per 1mm width of gel lane.



VC Lambda / EcoRI + HindIII Marker

Description

This lambda marker is suitable for sizing linear double-stranded DNA fragments in agarose gel. The lambda DNA is completely digested with *Eco*R I and *Hind* III and yields 12 fragments in size (in bp): 21226, 5148, 4973, 4268, 3530, 2027, 1904, 15844, 1375, 947, 831, 564.

Storage Buffer

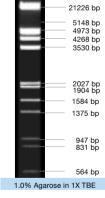
10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C

6X Loading Dye Solution 10mM Tris-HCl (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60%

glycerol.

Usage Recommendation

Use 0.05-0.10µg of the DNA marker per 1mm width of gel lane.



Catalog No	Description	Pack Size
NM2401	VC Lambda / <i>Eco</i> R I Marker	50 µg
NM2402	VC Lambda / <i>Eco</i> R I Marker	5 x 50 μg
NM2403	VC Lambda / EcoR I Marker (ready-to-use)	50 μg
NM2404	VC Lambda / EcoR I Marker, (ready-to-use)	5 x 50 μg
NM2405	VC Lambda / Hind III Marker	50 μg
NM2406	VC Lambda / Hind III Marker	5 x 50 μg
NM2407	VC Lambda / Hind III Marker (ready-to-use)	50 μg
NM2408	VC Lambda / Hind III Marker (ready-to-use)	5 x 50 μg
NM2409	VC Lambda/ EcoR I + Hind III Marker	50 μg
NM2410	VC Lambda/ <i>Eco</i> R I + <i>Hin</i> d III Marker	5 x 50 μg
NM2411	VC Lambda/ <i>Eco</i> R I + <i>Hin</i> d III Marker (ready-to-use)	50 μg
NM2412	VC Lambda/ EcoR I + Hind III Marker (ready-to-use)	5 x 50 μg

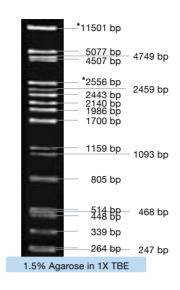
VC Lambda / Pst | Marker

Description

This lambda marker is suitable for sizing linear double-stranded DNA fragments in agarose and polyacrylamide gels. The lambda DNA is completely digested with *Pst*l and yields 29 fragments in size(in bp): *11501, 5077, 4749, 4507, *2556, 2459, 2443, 2140, 1986, 1700, 1159, 1093, 805, 514, 468, 448, 339, 264 and 247. 216, 211, 200, 164, 150, 94, 87, 72, 15 (obscure bands)

Storage Buffer 10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C 6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol.

Usage Recommendation
Use 0.05-0.10µg of the
DNA ladder per 1mm
width of gel lane.



VC pUC19 / MspI Marker

Description

Serves as molecular weight (in bp) standards for electrophoresis for both agarose and polyacrylamide gels. Suitable for sizing of double-stranded DNA fragments. Please note that bands indicated by *migrate anomalously. The pUC19 DNA is completely digested with *Msp*l and yields 13 fragments in size (bp): 501, 489, 404, 331, 242, 190, 147, 111, 110. 67, 34, 26 (obscure bands)

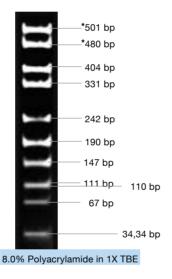
Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol.

Usage Recommendation Use 0.05-0.10µg of the

DNA ladder per 1mm width of gel lane.



VC pBR322 / Hae III Marker

Description

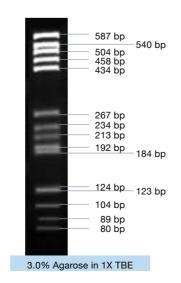
The DNA marker is suitable for sizing linear double-stranded DNA fragments in agarose gel electrophoresis. The pBR322 DNA is completely digested with III and yields 22 fragments in size (in bp): 587, 540, 504, 458, 434, 267, 234, 213, 192, 184, 124, 123, 104, 89, 80. 64, 57, 51, 21, 18, 11, 8 (obscure bands).

Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C

6X Loading Dye Solution 10mM Tris-HCI (pH 8.0), 60mM EDTA, 0.03% bromophenol blue, 0.03% xylene cyanol FF and 60% glycerol. Usage Recommendation
Use 0.05-0.10µg of the
DNA ladder per 1mm

width of gel lane.



Custom Made DNA Markers

We also provide customized sizing of DNA markers using desired restriction endonucleases on DNA templates as listed below.

Lambda DNA
 pBR322
 pUC18
 pUC19

Kindly contact your local distributor for more details.

Catalog No	Description	Pack Size
NM2425 NM2426 NM2427 NM2428	VC Lambda / Pst Marker VC Lambda / Pst Marker VC Lambda / Pst Marker (ready-to-use) VC Lambda / Pst Marker (ready-to-use)	50 µg 5 x 50 µg 50 µg 5 x 50 µg
NM2413 NM2414 NM2415 NM2416	VC pUC19 / Msp I Marker VC pUC19 / Msp I Marker VC pUC19 / Msp I Marker (ready-to-use) VC pUC19 / Msp I Marker (ready-to-use)	50 μg 5 x 50 μg 50 μg 5 x 50 μg
NM2429 NM2430 NM2431 NM2432	VC pBR 322 / HaeIII Marker VC pBR 322 / HaeIII Marker VC pBR 322 / HaeIII Marker (ready-to-use) VC pBR 322 / HaeIII Marker (ready-to-use)	50 µg 5 x 50 µg 50 µg 5 x 50 µg

Lambda DNA

(dam- and dcm-)

Description

The lambda DNA was isolated from bacteriophage lambda (cl857 ind 1 Sam 7) obtained from the heat inducible lysogenic *E. coli* strain (dam⁻ and dcm⁻).

Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

Quality Control

Purified DNA is assayed for contaminating exonuclease,

non-specific nucleases and phosphatase.

Lambda DNA

Description

The lambda DNA was isolated from bacteriophage lambda (cl857 1 7) obtained from the heat inducible lysogenic strain (dam+ and dcm+).

Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

Quality Control

Purified DNA is assayed for contaminating exonuclease,

non-specific nucleases and phosphatase.

pBR322 DNA

Description

pBR322 DNA is a commonly used plasmid cloning vector. Isolated from *E. coli* strain Top10F'.

Storage Buffer

Quality Control

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

Gel analysis for purity. Digested with Alul, BsnI and

HindIII for cut pattern confirmation.

pUC18 DNA

Description

pUC18 is commonly used plasmid cloning vectors. Isolated from strain Top10F'.

Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

Quality Control

Gel analysis for purity. Digested with *EcoRI*, *HindIII* and *MspI* for cut pattern confirmation.

pUC19 DNA

Description

pUC19 is commonly used plasmid cloning vectors. Isolated from *E. coli* strain Top10F'. Multiple Cloning site in opposite orientation from pUC18.

Storage Buffer

10mM Tris-HCI (pH 8.0) and 1mM EDTA. Store at -20°C.

Quality Control

Gel analysis for purity. Digested with EcoR I, HindIII and

DTA. Store Mspl for cut pattern confirmation.

Catalog No	Description	Pack Size
NN1401	Lambda DNA (dam ⁻ and dcm ⁻), 0.3μg/μl	500 µg
NN1402	Lambda DNA, 0.3μg/μl	500 μg
NN1404	pBR322 DNA, 0.2-0.5µg/µl	100 µg
NN1405	pUC18 DNA, 0.2-0.5µg/µl	50 μg
NN1406	pUC19 DNA, 0.2-0.5µg /µl	50 μg

Ladders & Markers / Nucleic Acids / Nucleotides

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Ladders & Markers / Nucleic Acids / Nucleotides

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dATP*

Description

dATP (2'-deoxyadenosine 5'-triphosphate) is supplied as a 100mM aqueous solution at pH 7.0.

Storage Buffer **Quality Control**

Funtionally tested in PCR with Tag and Pfu DNA Store at -20°C.

Polymerase. Purity at the nucleotide is >98% by HPLC.

dCTP*

Description

dCTP (2'-deoxycystidine 5'-triphosphate) is supplied as a 100mM aqueous solution at pH 7.0.

Storage Buffer **Quality Control**

Store at -20°C. Funtionally tested in PCR with Tag and Pfu DNA

Polymerase. Purity at the nucleotide is >98% by HPLC.

dGTP*

Description

dGTP (2'-deoxguanosine 5'-triphosphate) is supplied as a 100mM aqueous solution at pH 7.0.

Storage Buffer **Quality Control**

Store at -20°C. Funtionally tested in PCR with Taq and Pfu DNA

Polymerase. Purity at the nucleotide is >98% by HPLC.

dTTP*

Description

dTTP (2'-deoxythymidine 5'-triphosphate) is supplied as a 100mM aqueous solution at pH 7.0.

Storage Buffer **Quality Control**

Store at -20°C. Funtionally tested in PCR with Tag and Pfu DNA

Polymerase. Purity at the nucleotide is >98% by HPLC.

dUTP*

dUTP (2'-deoxyuridine 5'-triphosphate) is supplied as a 100mM aqueous solution at pH 7.0.

Storage

Quality Control

Application

Store at -20°C. Functionally tested in PCR with Tag and Pfu DNA Polymerases.

Generally used in various PCR, RT-PCR, cDNA

Purify of each > 98% by

synthesis and primer extension.

HPLC.

dNTP Set*

Description

dNTP Set consists of 100mM aqueous solution of dATP, dCTP, dGTP and dTTP, each in a separate vial.

Storage Buffer

Store at -20°C. Funtionally tested in PCR with Tag and Pfu DNA

Polymerase. Purity at the nucleotide is >98% by HPLC.

dNTP Mix*

Description

dNTP Mix is an aqueous solution containing dATP, dCTP, dGTP and dTTP, each in a final concentration as indicated.

Available Concentration

 10 mM 2 mM

25 mM

* Storage Store at -20°C. * Quality Control

Functionally tested in

PCR with Tag and Pfu DNA Polymerase. Purify

of each dNTP nucleotide >98% by HPLC.

* Application

Generally used in various PCR applications, cDNA synthesis, primer extension, DNA sequencing and DNA

labeling reactions.

Catalog No	Description	Pack Size
NP2401	dATP, 100mM	0.25 ml
NP2402	dCTP, 100mM	0.25 ml
NP2403	dGTP, 100mM	0.25 ml
NP2404	dTTP, 100mM	0.25 ml
NP2405	dUTP, 100mM	0.25 ml
NP2406	dNTP Set, 100mM	4 x 0.25 ml
NP2407	dNTP Set, 100mM	4 x 1 ml
NP2408	dNTP Set, 100mM	4 x 5 ml
NP2409	dNTP Mix, 10mM	0.25 ml
NP2410	dNTP Mix, 10mM	1 ml
NP2411	dNTP Mix, 2mM	1 ml
NP2412	dNTP Mix, 2mM	5 x 1 ml
NP2413	dNTP Mix, 25mM	0.25 ml
NP2414	dNTP Mix, 25mM	5 x 0.25 ml

Chromatein Prestained Protein Ladder

Description

Chromatein Prestained Protein Ladder contains 11 proteins that resolve into sharp, tight bands in the range of 10-175 kDa. It is supplied in a loading buffer for direct loading on gels. Allows monitoring molecular weight separation during electrophoresis, estimation of molecular weights of interest and evaluate western transfer ef ciency.

Feature

Broad Range: 10-175kDa Convenient: Supplied in a loading buffer for direct loading.

Easy Identification: ~10, ~40, and ~90kDa

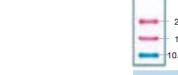
reference bands coupled with a blue dye.

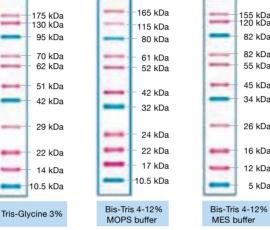
Quality Control

Tested in SDSpolyacrylamide gel electrophoresis and western blotting

Storage

Stable at 4°C for 3 months Store at -20°C for 24 months





Tricolor Broad Range Prestained Protein Ladder

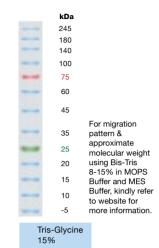
Description

Tricolor Broad Range Prestained Protein Ladder contains 13 proteins that resolve into sharp, tight bands in the range of 5-245kDa.lt can be used to monitor molecular weight separation during electrophoresis, estimate molecular weights of proteins of interest, and evaluate western transfer efficiency. It contains 2 reference bands ~25 and ~75 kDa coupled with blue chromophore as well as red dye and green dye for easy identification. It can be used on PVDF and nylon membrane.

Storage Buffer **Quality Control**

Store at -20°C. Tested in SDS-polyacrylamide gel electrophoresis and

western blotting



Whole Blue Range Prestained Protein Ladder

Description

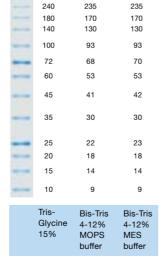
Whole Blue Range Prestained Protein Ladder contains 12 proteins that resolve into sharp, tight bands in the range of 10-240kDa.lt can be used to monitor molecular weight separation during electrophoresis, estimate molecular weights of proteins of interest, and evaluate western transfer efficiency. It contains 2 reference bands ~25 and ~72 kDa coupled with blue chromophore for easy identification. It can be used on PVDF and nylon membrane.

Storage Buffer

Quality Control

Store at -20°C

Tested in SDS-polyacrylamide gel electrophoresis and western blotting



82 kDa

82 kDa

55 kDa

45 kDa

34 kDa

26 kDa

16 kDa

12 kDa

5 kDa

kDa

Catalog No	Description	Pack Size
PR0602	Chromatein Prestained Protein Ladder (ready-to-use)	2 x 250 µl
PR0623	Whole Blue Range Prestained Protein Ladder	2 x 250 µl
PR0624	Tricolor Broad Range Prestained Protein Ladder	2 x 250 µl



Polymerases & Modifying Enzymes

Polymerases & Modifying Enzymes

Reverse Transcriptases

Modifying Enzyme

Ribonuclease Inhibitor RNase-Free

T4 DNA Ligase

AMV Reverse Transcriptase (Recombinant)



Description

Avian Myeloblastosis Virus (AMV) Reverse Transcriptase is an RNA-dependent DNA polymerase (aß holoenzyme) with molecular weight of 157 kDa. It synthesizes a complementary DNA strand initiating from a primer using either RNA (cDNA synthesis) or single-stranded DNA as a template.

 $20 u/\mu l$

Assay Condition

50mM Tris- HCI (pH8.3), 6mM MgCl₂, 40mM KCl, 4mM DTT, 0.5mM [3H]-TTP (10-20c/m/pmol), 0.4mM $Poly(rA) \bullet (dT)_{12-18}$, in a reaction volume of 25µl.

10X Buffer AMV-RT 250mM Tris-HCI (pH 8.3), 500mM KCI, 50mM MgCl₂ and 20mM DTT.

Storage Buffer

200mM Potassium Phosphate (pH 7.2), 0.2% Triton™ X-100, 2mM DTT and 50% glycerol. Store at -70°C for long term periods. Store at -20°C for short term (1 month)

Thermal Inactivation

80°C for 10 minutes

1u is defined as the amount of enzyme that is required to incorporate 1nmol of dMTP into an acid-insoluble material in 10 minutes at 37°C using Poly(rA) • (dT)₁₂₋₁₈ as a template primer.

Application

- First strand synthesis of cDNA.
- Synthesis & cDNA for cloning.
- DNA labelling.
- Primer extensions and RNA sequencing.
- RT-PCR
- Dideoxy sequencing of DNA & RNA

All preparations are assayed for contaminating endonuclease, exonuclease and non-specific RNase activities.





M-MuLV Reverse Transcriptase {RNase H - }



Description

Molonev Murine Leukemia Virus (M-MuLV) Reverse Transcriptase is an RNA dependent DNA polymerase. It can synthesize a complementary DNA strand initiating from a primer using either RNA or single-stranded DNA as a template. The absence of RNase H activities enhances the synthesis of long cDNAs and therefore the enzyme is recommended for preparing long cDNAs.

Concentration

100 - 500u/µl

Assay Condition

50mM Tris- HCI (pH8.3), 6mM MgCl₂, 10mM DTT, 0.4mM Poly(rA) • (dT)₁₂₋₁₈, in a reaction volume of 50µl.

10X Buffer M-MuLV RT 500mM Tris-HCI (pH 8.8 at 25°C), 67mM MgCl₂, 3mM MgCl₂ and 10mM DTT. Store at -20°C.

Storage Buffer

10mM K-phosphate (pH 7.5), 0.1mM EDTA, 200mM NaCl, 7mM 2-mercaptoethanol and 50% glycerol.Store at -20°C.

Thermal Inactivation

70°C for 10 minutes

1u is defined as the amount of enzyme that is required to incorporate 1nmol of dTTP into an acid-insoluble material in 10 minutes at 37°C using Poly(rA) • oligo(dT).

Application

- First strand cDNA synthesis.
- DNA labeling.
- RNA analysis by primer extension.

Purified free of detectable levels of RNase, endonuclease and exonuclease activities.

Catalog No	Description	Pack Size
ME2301 ME2302	AMV Reverse Transcriptase (Recombinant) AMV Reverse Transcriptase (Recombinant)	500u 2500u
ME2305 ME2306	M-MuLV Reverse Transcriptase (RNase H ⁻) M-MuLV Reverse Transcriptase (RNase H ⁻)	10000u 50000u





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T4 DNA Ligase





T4 DNA Ligase catalyzes the formation of a phosphodiester bond between juxtaposed 5'-phosphate and 3'-hydroxyl termini in duplex DNA or RNA. This enzyme will join blunt-end and cohesive end termini as well as repair single stranded nicks in duplex DNA, RNA, or DNA-RNA hybrids.

50 - 200u/µl

Features

- Seals single-stranded nicks in duplex DNA, RNA or DNA-RNA hybrids.
- ATP is an essential cofactor for the reaction.
- Ultrapure recombinant protein.

10X Buffer T4 Ligase 50mM Tris-HCI (pH7.8 at 25°C), 10mM MgCl₂, 10mM DTT, 1mM ATP and 25µg/ml BSA. Store at -20°C

Thermal Inactivation

65°C for 15 minutes.

1u (*Cohesive End Ligation Unit) is defined as the amount of enzyme that is required to give 50% ligation of Hind III fragments of lambda DNA (5' DNA termini concentration of 0.12uM [300ua/ml]) in 20ul of 1X T4 DNA Ligase Buffer in 30 minutes at 16°C.

* One Cohesive End Ligation Unit is equal to 0.015 Weiss units. Equivalently, one Weiss unit is equal to 67 Cohesive End Ligation Units.

Storage Buffer

10mM Tris-HCI (pH7.5), 50mM NaCl, 0.1mM EDTA, 10mM 2-mercaptoethanol and 50% glycerol. Store at -20°C.

Application

- Catalyzes the linkage of 5' or 3' blunt/cohesive ends of doublestranded DNA by formation of phosphodiester bond.
- Joining of oligonucleotide linkers or adapters to blunt ends.
- Repair nicks formation in duplex nucleic acids.

All preparations are assayed for contaminating endonuclease, exonuclease and non-specific DNase activities.





Ribonuclease Inhibitor RNase-Free

Description

Ribonuclease Inhibitor RNase-free inhibits the activity of RNase A, B, C by binding them in a noncompetitive mode at a 1:1 ratio. It does not inhibit RNase 1, T1, T2, H, U1, U2, CL3 and other enzymes.

40 u/µl

Features

- · Performs under a wide range of reaction conditions.
- Protects RNA from degradation at temperature up to 55°C.
- Increase the time RNA can be safely stored.

Assay Conditions

100mM Tris-HCI (pH 7.5), 1.2mM EDTA, 0.1mg/ml BSA, 100ng/ml RNase, 0.1mg/ml E.coli [5H] RNA, 50mg/ml yeast RNA and 8mM DTT.

Storage Buffer

20mM HEPES-KOH (pH7.5), 50mM KCI, 5mM DTT and 50% glycerol.

1u is defined as the amount of ribonuclease inhibitor that inhibits the activity of 5ng of Ribonuclease A by 50%.

- Applied in procedures where RNase contamination constitutes a problem:
- in vitro transcription.
- in vitro translation. - cDNA synthesis.

- separation and identification of specific ribonuclease activities.

fractions that contain mRNA-

- isolation of mammalian cell

protein complex.

Quality Control

The absence of endodeoxyribonucleases, exodeoxyribonucleases, phosphatases and ribonucleases confirmed by appropriate quality tests. Functionally tested in RNA and cDNA synthesis.

Catalog No	Description	Pack Size
ME4303	T4 DNA Ligase	4000u
ME4304	T4 DNA Ligase	20000u
ME4309	Ribonuclease Inhibitor RNase-Free	2500u
ME4310	Ribonuclease Inhibitor RNase-Free	4 x 2500u



Cloning Kits pTG19-T 2680 bp

Flex-C Cloning Kits

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Description

Flexi-C Cloning Kit is a high efficient, rapid and easy-to-use PCR cloning kit. The Flexi-C Enzyme allows direct cloning of any PCR fragments to any linearized expression vector at any site in a single 20-minute reaction.

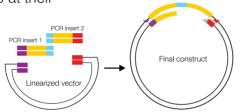
The application protocol is simple. The PCR fragments can be generated by PCR Polymerase (Tag DNA Polymerase) with primers that are designed to have at least 10 bases of homology at their linear ends. No additional treatment of the PCR fragment is required (such as restriction digestion, ligation, phosphorylation, or blunt-end polishing). The linearized vector can be generated by PCR or restriction enzymes (single or double cut). The Flexi-C Enzyme joins PCR fragments and linearized vectors accurately and efficiently by recognizing the 10bp overlap at their ends.

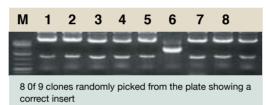
Features

- · Clone any insert, at any site within any vector
- Restriction enzyme, phosphatase and ligase free system
- Joining multiple fragments at once
- Broad PCR size up to 10kb
- Good for 5' overhangs, 3' overhangs, blunt end
- Precise insertion at a desired orientation
- High Efficiency with > 95% positive clones
- Multiple applications:
 - adding adaptor, linker and tag before or after the insert
 - mutation generation
 - gene synthesis
- High through put application

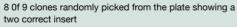
Kit Components

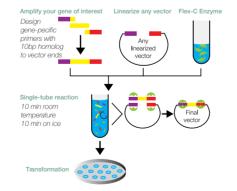
- Flex-C Cloning reagents
- 500u of Tag DNA Polymerase and buffers
- 0.25ml of 10mM dNTP Mix
- 1ml of nuclease-free water











Ordering Information

Catalog No	Description	Pack Size
MEPL01 MEPL02	Flex-C Cloning Kit (Without Competant Cell) Flex-C Cloning Kit (With Competant Cell)	20 applications 20 applications

pTG19-T PCR Cloning Vector

Description

The pTG19-T vector is designed for rapid and efficient cloning of PCR products with 3'dA overhangs. The linearized pTG19-T vector with 3'-dT overhangs prevent vector recircularization, therefore resulting in high percentage of recombinant clones and low background.

Features

- Convenient ready-to-use linearized 3'dT overhang pTG19-T vector.
- Efficient -more than 80% of the recombinant clones contain the target DNA
- Rapid clone selection:
 - -lacZ gene for blue/white selection.
 - -M13 primer sites for PCR screening and sequencing.
 - -BamHI restriction enzyme can be used to release the insert from the pTG19-T vector.

Kit Components

- pTG19-T vector
- Control insert

Quality Control

• More than 80% clones are white with control insert.

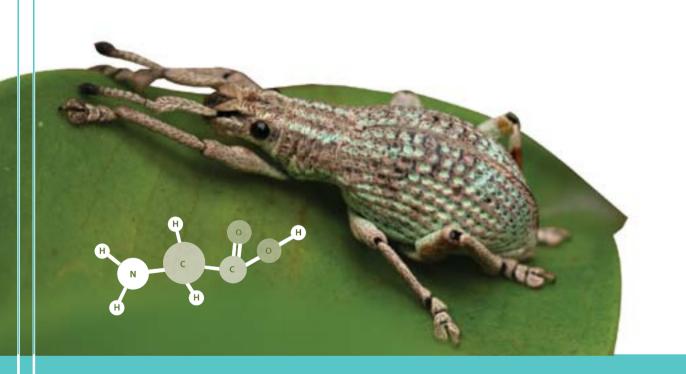
Cloning Kits

- More than 85% of white clones are positively by restriction endonuclease digestion.
- The 3'dT overhangs for every batch vectors is confirmed by sequencing of five recombinant clones.

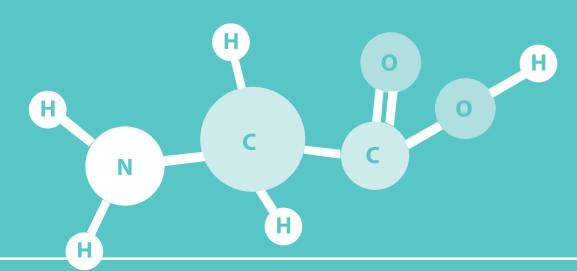
Storage & Stability

- All components are stable at -20°C for one year if properly stored.
- To avoid frequent feezethaw cycles, keeping small aliquots at -20°C is recommended.

Catalog No	Description	Pack Size
TA010	pTG19-T Cloning Vector	20 applications



Biochemicals



Biochemicals

Biochemicals

Protein Biochemicals

Storage

Store at -20°C

Store at -20°C

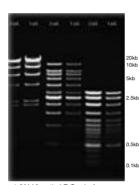
Vivantis Technologies . Product Catalog Volume IV

Agarose (Molecular Biology Grade)

Description

Vivantis Agarose is ideal for routine analysis of nucleic acids by gel eletrophoresis and blotting. Each gel sharply resolve DNA and provides consistent resolution from lot-to-lot. This molecular biology grade agarose has no detectable DNase or RNase activity and forms strong gels with low background upon ethidium bromide, SYBR® Green, or GelStar® staining.





1.0% Vivantis LE Grade Agarose 1X TAE Buffer

1.0% Vivantis LE Agarose, 1X TAE Buffer Lane 1,2 = Lambda / *Hind* III DNA Marker (#NM2405) Lane 3,4 = VC 1kb DNA Ladder (#NL1409) Lane 5,6 = VC 100bp Plus DNA Ladder (#NL1405)

Features:

- None RNase/ DNase Activities
 Consistent Resolution for
- Consistent Resolution for lot-to-lot assay
- High Gel Strength

Application:

- Used for analysis and recovery of DNA and RNA
- Recommended for protein gel electrophoresis applications such as Ouchterlony (antigen- antibody interaction assay) and radial immunodiffusion (RID) (antigen quantitation assay)

Typical Properties of Agarose:

- Gelling temp : 36° ± 1.0°C
- Melting temp : ≥ 90°C
- Moisture content :10%
- Sulfate: 0.15%
- EEO, (-mr): 0.09 0.12
- Gel strength : 1200g/cm²
- RNase/DNase Activity:

None detected

Storage:

Store at RT

Ordering Information

Catalog No	Description	Pack Size
PC0701-100g	Agarose (Molecular Biology Grade)	100g
PC0701-500g	Agarose (Molecular Biology Grade)	500g
PC0701-1kg	Agarose (Molecular Biology Grade)	1kg

Biochemicals

(All Vivantis biochemicals are of Molecular Grade)

Adenosine 5'-Monophosphate, Disodium Salt (AMP.2Na)

Ordering Information

Catalog No	Pack Size
PC1001-25g	25g
PC1001-100g	100g

Adenosine 5'-Diphosphate, Disodium Salt (ADP.2Na)

Ordering Information

Catalog No	Pack Size
PC1002-5g	5g
PC1002-25g	25g

Adenosine 5'-Triphosphate, Disodium Salt (ATP.2Na)

Ordering Information

Catalog No	Pack Size
PC1003-5g	5g
PC1003-25g	25g

Ammonium Sulfate

Ordering Information

Catalog No	Pack Size
PC0902 - 500g	500g

Storage

Storage

Store at -20°C

Store at RT

Bovine Serum Albumin (BSA)

Ordering Information

Catalog No	Pack Size
PC0903 - 25g	25g

Storage

Store at 2-8°C



Calcium Chloride

Vivantis Technologies . Product Catalog Volume IV

Ordering Information

Catalog No	Pack Size
PC0904 - 500g	500g

Storage Store at RT

Ordering Information

Storage Store at RT

Storage

Storage

Storage

Store at RT

Store at -20°C

Store at 2-8°C

Store at RT

Catalog No Pack Size PC0706 - 500g 500g PC0706 - 1kg 1kg

EDTA, Disodium Salt, Dihydrate

3-[(3-cholamidopropyl)dimethyl-ammonio]-1-propanesulfonate (CHAPS)

Ordering Information

Catalog No	Pack Size
PC0925-5g	5g

Ficoll 400

Ordering Information

Formamide

Catalog No

Ordering Information

PC0907 - 500ml

Catalog No	Pack Size
PC0920-100g	100g

Pack Size

Pack Size

Pack Size

1g

5g

10g

5g

10g

500ml

Deoxyribonuclease I (DNase I)

Ordering Information

Ordering information	
Catalog No	Pack Size
PC0704 - 25mg	25mg
PC0704 - 50mg	50mg
PC0704 - 100mg	100mg
PC0704 - 1g	1g

Storage

Store at -20°C

Deoxyribonuclease I (DNase I) - For RNA work

Ordering Information

Catalog No	Pack Size
PC0719 - 50kU	50kU
PC0719 - 100kU	100kU

Storage

Store at RT

Storage Store at -20°C

Pack Size	
50kU	

Glycerol

Ordering Information

Ordering Information

Catalog No

PC0708 - 1g

PC0708 - 5g

PC0708 - 10g

Ordering Information

Catalog No

PC0710 - 1g

PC0710 - 5g

PC0710 - 10g

Catalog No	Pack Size
PC0908 - 500ml	500ml
PC0908 - 1L	1L

Isopropyl-beta-D-thiogalactopyranonside (IPTG)

Diethylpyrocarbonate (DEPC)

Ordering Information

Catalog No	Pack Size
PC0905 – 25ml	25ml
PC0905 - 100ml	100ml

Storage

Store at 2-8°C

Dimethysulfoxide (DMSO)

Ordering Information

Catalog No	Pack Size
PC0906 - 500ml	500ml

Storage

DL-Dithiothreitol (DTT)

Ordering Information

Catalog No	Pack Size
PC0705 - 5g	5g
PC0705 - 25g	25g
PC0705 - 100g	100g

Store at RT

Storage

Store at -20°C

2-(N-Morpholino) Ethanesulfonic Acid (MES)

Lysozyme, Egg White

Ordering Information

Ordering information	
Catalog No	Pack Size
PC0927-100g	100g

Storage Store at -20°C

Storage

Protein Biochemicals

Vivantis Technologies . Product Catalog Volume IV

(All Vivantis protein biochemicals are of Molecular Biology Grade)

Acrylamide Ordering Information Storage

Store at RT

Catalog No	Pack Size
PR0603 – 100g	100g
PR0603 – 500g	500g
PR0603 – 1kg	1kg

Acryl / Bis 37.5:1(30:0.5) (Premixed Powder)

Storage

Ordering Information

Catalog No Pack Size 40g PR0604 - 40g PR0604 - 200g 200g

APS (Ammonium Persulfate)

Ordering Information

Catalog No	Pack Size
PR0605 – 25g	25g
PR0605 – 100g	100g

Bis-Acrylamide

Ordering Information

Catalog No	Pack Size
PR0606 – 50g	50g
PR0606 – 250g	250g

Blue Lightning Stain II

Ordering Information

Catalog No	Pack Size
PR06017 - 1L	1L

Boric Acid

Ordering Information

Catalog No	Pack Size
PR0607 – 500g	500g
PR0607 – 1kg	1kg

Glycine

Ordering Information

Catalog No	Pack Size
PR0608 - 1kg	1kg

Store at RT

Storage Store at RT

Storage

Store at RT

Storage

Store at RT

Storage

Store at RT

Storage

Store at RT

3-(N-Morpholino) propane-sulfonic acid (MOPS) Store at RT

Ordering Information

Catalog No	Pack Size
PC0928-100g	100g

PEG 8000

Ordering Information

Catalog No	Pack Size
PC0921-1kg	1kg

Phosphate Buffered Saline (PBS) (Powder and 10X Ready Pack)

*1X PBS (pH7.3-7.5 at 2°C) solution contains 137mM NaCl, 2.7mM KCL and 10mM Phosphate buffer. There is sufficient powder to prepare 10L by using 9.88g/L. Each pack prepares 1L of 10X Concentrate.

Ordering Information

Catalog No	Pack Size
PC0711 - 2pk	2pks

Potassium Acetate

Ordering Information

Catalog No	Pack Size
PC0909 – 500g	500g

Potassium Chloride

Ordering Information

Ordering information	
Catalog No	Pack Size
PC0910 - 500g	500g

Potassium Phosphate, Dibasic, Anhydrous

Ordering Information

Catalog No	Pack Size
PC0911 - 500g	500g

Potassium Phosphate, Monobasic, Anhydrous

Ordering Information

Ordoning information	
Catalog No	Pack Size
PC0912-500g	500g

Storage

Storage

Store at RT

Storage

Storage

Store at RT

Store at RT

Store at RT

Storage

Store at RT

Storage

Vivantis Technologies . Product Catalog Volume IV

Proteinase K

Ordering Information

Catalog No	Pack Size
PC0712 - 100mg	100mg
PC0712 - 1g	1g
PC0712 - 1ml	1ml

Storage

Store at -20°C or RT

Ribonuclease A (RNase A)

Ordering Information

ordoring information	
Catalog No	Pack Size
PC0713 - 250mg	250mg
PC0713 - 500mg	500mg
PC0713 - 1g	1g
PC0713 - 1ml	1ml
PC0715 - 1ml	1ml

Storage

Store at -20°C

Sodium Acetate, Anhydrous

Ordering Information

Catalog No	Pack Size
PC0913 - 500g	500g

Storage

Store at RT

Sodium Chloride

Ordering Information

Catalog No	Pack Size
PC0914 - 500g	500g
PC0914 - 1kg	1kg

Storage

Store at RT

Sodium Phosphate, Dibasic, Anhydrous

Ordering Information

Catalog No	Pack Size
PC0916 - 500g	500g

Storage Store at RT

Sodium Phosphate, Monobasic, Anhydrous

Ordering Information

Catalog No	Pack Size
PC0917 - 500g	500g

Storage

Store at RT

Sucrose

Ordering Information

Ordering information	
Catalog No	Pack Size
PC0918 - 500g	500g
PC0918 - 1kg	1kg

Storage

Store at RT

Triton X-100

Ordering Information

Catalog No	Pack Size
PC0923-1L	1L

Tween 20

Ordering Information

Catalog No	Pack Size
PC0919 - 500ml	500ml
PC0919 - 1L	1L

Tween 80

Ordering Information

Catalog No	Pack Size
PC0924-1L	1L

X-Gal (5-Bromo-4-Chloro-3-Indolyl-beta-D-galactopyranoside)

Ordering Information

Catalog No	Pack Size
PC0716 - 100mg	100mg
PC0716 - 1g	1g
PC0716 - 500mg	500mg

Xylene Cyanol FF

Ordering Information

Ordering information	
Catalog No	Pack Size
PC0718 - 20g	20g

Storage

Store at RT

Storage

Store at RT

Storage

Store at RT

Storage

Store at -20°C

Storage

Biochemicals

Vivantis Technologies . Product Catalog Volume IV

Protein Biochemicals

(All Vivantis protein biochemicals are of Molecular Biology Grade)

Silver Nitrate

Ordering Information

Catalog No	Pack Size
PR0610 – 25g	25g
PR0610 – 100g	100g

Storage

Store at RT

SDS (Sodium Dodecyl Sulfate) Ordering Information

Catalog No	Pack Size
PR0611 – 100g	100g
PR0611 – 250g	250g
PR0611 – 500g	500g
PR0611 – 1kg	1kg

Storage

Store at RT

TEMED

(N,N,N',N'-Tetramethylethylene-Diamine)

Ordering Information

Catalog No	Pack Size
PR0616 – 25ml	25ml
PR0616 – 50ml	50ml
PR0616 – 100ml	100ml

Storage Store at RT

Tris Base

Ordering Information

Catalog No	Pack Size
PR0612 – 500g	500g
PR0612 – 1kg	1kg

Storage

Store at RT

Tris-HCI

Ordering Information

Catalog No	Pack Size
PR0614 – 500g	500g
PR0614 – 1kg	1kg

Storage

Store at RT

Urea

Ordering Information

Catalog No	Pack Size
PR0615 - 500g	500g
PR0615 – 1kg	1kg

Storage





Ready Made Buffers



Ready Made Buffers

Vivantis Technologies	. Product	Catalog	Volume I	IV
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Catalog No	Description	Pack Size	Storage
PB0100 – 500ml	Nuclease-free, Water	500ml	Store at RT
PB0245 – 500ml	0.5M EDTA, PH 8.0, Ultra Pure Grade	500ml	Store at RT
PB0312-1L	1X Phosphate Buffered Saline, pH7.2; Ultra Pure Grade	1L	Store at RT
PB0314-1L	1X Phosphate Buffered Saline, pH7.4; Ultra Pure Grade	1L	Store at RT
PB0315-1L	1X Phosphate Buffered Saline, pH8.0; Ultra Pure Grade	1L	Store at RT
PB0342-1L	10X Phosphate Buffered Saline; Ultra Pure Grade *1X Phosphate Buffered Saline, pH7.2	1L	Store at RT
PB0344 – 1L	10X Phosphate Buffered Saline; Ultra Pure Grade *1X Phosphate Buffered Saline, pH7.4	1L	Store at RT
PB0345-1L	10X Phosphate Buffered Saline; Ultra Pure Grade *1X Phosphate Buffered Saline, pH8.0	1L	Store at RT
PB0354-1L	20X Phosphate buffered Saline; Ultra Pure Grade *1X Phosphate Buffered Saline, pH7.4	1L	Store at RT
PB0355-1L	20X Phosphate buffered Saline; Ultra Pure Grade *1X Phosphate Buffered Saline, pH8.0	1L	Store at RT
PB0461-500ml	3M Sodium Acetate, pH5.2, Biotechnology Grade	500ml	Store at RT
PB0570-500ml	5M Sodium Chloride; Biotechnology Grade	500ml	Store at RT
PB0640-500ml	10% Sodium Dodecyl Sulfate (SDS) Solution; Biotechnology Grade	500ml	Store at RT
PC0729-1L	0.05M Sodium Pyrophosphate Buffer, pH7.5; Ultra Pure Grade	1L	Store at RT
PC0730-1L	0.05M Sodium Pyrophosphate Buffer, pH9.0; Ultra Pure Grade	1L	Store at RT
PB0735-1L	0.05 Sodium Hydroxide; Biotechnology Grade	1L	Store at RT
PB0750-500ml	1M Sodium Hydroxide, Biotechnology Grade	500ml	Store at RT
PB0852-1L	1M Tris Buffer, pH7.4; Ultra Pure Grade	1L	Store at RT
PB0853-1L	1M Tris Buffer, pH7.0; Ultra Pure Grade	1L	Store at RT
PB0854-1L	1M Tris Buffer, pH7.5; Ultra Pure Grade	1L	Store at RT
PB0855-1L	1M Tris Buffer, pH8.0; Ultra Pure Grade	1L	Store at RT

Catalog No	Description	Pack Size	Storage
PB0858-1L	1M Tris Buffer, pH6.8, Ultra Pure Grade	1L	Store at RT
PB0898-1L	1.5M Tris Buffer, pH8.8; Ultra Pure Grade	1L	Store at RT
PB0940-1L	10X Tris-Acetate-EDTA (TAE) Buffer; Ultra Pure Grade *1X Tris-Acetate-EDTA Buffer, pH8.0	1L	Store at RT
PB0950-1L	20X Tris-Acetate-EDTA (TAE) Buffer; Ultra Pure Grade *1X Tris-Acetate-EDTA Buffer, pH8.0	1L	Store at RT
PB0990-1L	50X Tris-Acetate-EDTA (TAE) Buffer; Ultra Pure Grade *1X Tris-Acetate-EDTA Buffer, pH8.0	1L	Store at RT
PB1030-1L	5X Tris-Borate-EDTA (TBE) Buffer; Ultra Pure Grade *1X Tris-Borate-EDTA Buffer, pH8.3	1L	Store at RT
PB1040-1L	10X Tris-Borate-EDTA (TBE) Buffer; Ultra Pure Grade *1X Tris-Borate-EDTA Buffer, pH8.3	1L	Store at RT
PB1110-1L	1X Tris Buffered Saline (TBS) pH7.4; Ultra Pure Grade	1L	Store at RT
PB1140-1L	10X Tris Buffered Saline (TBS), pH7.4; Ultra Pure Grade *1X Tris Buffered Saline, pH7.4	1L	Store at RT
PB1214-500ml	1X Tris-EDTA (TE) Buffer, pH7.5; Ultra Pure Grade	500ml	Store at RT
PB1215-500ml	1X Tris-EDTA (TE) Buffer, pH8.0; Ultra Pure Grade	500ml	Store at RT
PB1244-100ml	10X Tris-EDTA (TE) Buffer, Ultra Pure Grade *1X Tris-EDTA Buffer, pH7.5	100ml	Store at RT
PB1245-100ml	10X Tris-EDTA (TE) Buffer, Ultra Pure Grade *1X Tris-EDTA Buffer, pH8.0	100ml	Store at RT
PB1330-1L	5X Tris Phosphate-EDTA (TPE); Ultra Pure Grade	1L	Store at RT
PB1340-1L	10X Tris-Glycine (TG) Buffer, Ultra Pure Grade	1L	Store at RT
PB1410-1L	1X Tris-Glycine (TG) Buffer; Ultra Pure Grade	1L	Store at RT
PB1440-1L	10X Tris-Glycine (TG) Buffer; Ultra Pure Grade	1L	Store at RT
PB1510-1L	1X Tris-Glycine-Sodium Dodecyl Sulfate (TG-SDS) Buffer; Ultra Pure Grade	1L	Store at RT
PB1540-1L	10X Tris-Glycine-Sodium Dodecyl Sulfate (TG-SDS) Buffer; Ultra Pure Grade	1L	Store at RT
PB1610-1L	1X Tris Bufferred Saline-Tween 20 (TBST), Ultra Pure Grade	1L	Store at RT
PB1640-1L	10X Tris Bufferred Saline-Tween 20 (TBST), Ultra Pure Grade	1L	Store at RT



Tracking Dye 6x loading dye 6x loading dye with SDS **Nucleic Acid Dye** ViSafe Green Gel Stain ViSafe Red Gel Stain Viva SybrGreen Nucleic Acid Stain qPCR Dye Viva qGreen I Fluorescent Dye (Equivalent to SYBR® Green Dye) Viva qGreen II Fluorescent Dye (Equivalent to EvaGreen® Dye)

Dves

Tracking Dye

6X Loading Dye

(with xylene cyanol gel loading dye)

Vivantis Technologies . Product Catalog Volume IV

Used for loading DNA markers and samples in agarose gel. Contains 2 dyes; bromophenol blue and xylene cyanol FF to track DNA migration during electrophoresis. Bromophenol blue migrates with the 300bp fragment while xylene cyanol FF migrates with the 4000bp fragment.

Storage

- Store at 4°C for 1 week
- Store at -20°C for long storage

6X Loading Dye with SDS

This product is specially designed for loading DNA samples that contains high amount of proteins that may form complexes with DNA during gel electrophoresis. This product is suitable for use in prevention of band-shift (due to protein binding) or annealing of DNA during both agarose and polyacrylamide gel electrophoresis. The 6X Loading Dye with SDS contains 2 dyes; bromophenol blue and xylene cyanol FF to track DNA migration during electrophoresis.

Storage

Store at 4°C for 1 week

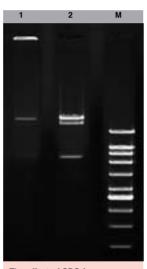
Store at -20°C for long storage

Application

Recommended for electrophoretic DNA sample after digestion with restriction endonuclease, ligation or diphosphorylation reactions.

Usage Recommendation

Add 5 volume of DNA sample to 1 volume of 6X Loading dye with SDS and mix well heat at 65°C for 10 minutes and chill on ice for 3 minutes prior to loading.



The effect of SDS for electrophoresis of DNA samples that contain high amount of DNA-protein complex Lane 1: Lambda DNA digested with Psp C I (with Buffer V1) loaded with 6X loading dye. Lane 2: Lambda DNA digested with Psp C I (with Buffer V1) loaded with 6X loading dye with Lane M: VC 1kb DNA Ladder

Ordering Information

Catalog No	Description	Pack Size
NM0410	6X Loading Dye (with xylene cyanol gel loading dye)	5 x 1 ml
NM0416	6X Loading Dye with SDS	5 x 1 ml

Nucleic Acid Dye

ViSafe Green Gel Stain (10000x in water)

Description

ViSafe Green Gel Stain is a stable, sensitive and environmentally safe fluorescent nucleic acid dye for staining double stranded DNA (dsDNA), single stranded DNA (ssDNA) or RNA in agarose gels or polyacrylamide gels.

The stain is noncytotoxic & nonmutagenic shown by Ames tests.

More sensitive compared to EtBr or Viva SybrGreen Nucleic Acid Stain.

Stable at room temperature for long-term storage. Stable to be microwaved or being heated. The working solution is stable at room temperature when kept in dark.

Suitable to stain dsDNA, ssDNA and RNA. Suitable to use in agarose gel or polyacrylamide gel. Compatible with downstream applications, such as gel recovery & cloning.

Easy precast gel staining and post-staining procedures.

Gel can be viewed with standard UV transilluminator, visible light gel reader or other gel imaging system.

Storage

Shipped at ambient temperature

Stored at 2-8°C or -20°C

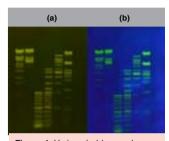
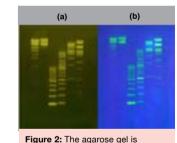


Figure 1: Various ladders and makers run at 1.5% TBE agarose gel. The agarose gel is poststained with ViSafe Green Gel Stain. The gel is visualized using transilluminator with (a) blue light: (b) UV light.



pre-stained with ViSafe Green Gel Stain Various ladders and makers run at 1.5% TBE pre-stained agarose gel. The gel is visualized using transilluminator with (a) blue light; (b) UV light.

ViSafe Red Gel Stain (10000x in water)

Vivantis Technologies . Product Catalog Volume IV

Description

ViSafe Red Gel Stain is a stable, sensitive and environmentally safe fluorescent nucleic acid dye for staining double stranded DNA (dsDNA), single stranded DNA (ssDNA) or RNA in agarose gels or polyacrylamide gels.

Features

The red gel stain is non cytotoxic & non mutagenic shown by

More sensitive compared to EtBr or Viva SybrGreen Nucleic Acid Stain.

Stable at room temperature for long-term storage. Stable to be microwaved or being heated. The working solution is stable at room temperature when kept in dark.

Suitable to stain dsDNA, ssDNA and RNA. Suitable to use in agarose gel or polyacrylamide gel. Compatible with downstream applications, such as gel recovery & cloning.

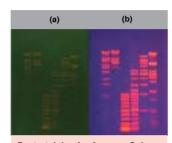
Easy precast gel staining & post-staining procedures.

Gel can be viewed with standard UV transilluminator, visible light gel reader, or other gel imaging system.

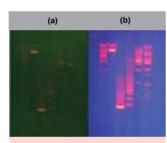
Storage

Shipped at ambient temperature

Stored at 2-8°C or -20°C



Post-staining for Agarose Gel Figure 1: Various ladders and markers run at 1.5% TBF agarose. gel. The agarose gel is post-stained with ViSafe Red Gel Stain. The gel is visualized using transilluminator with (a) blue light; (b) UV light.



Precast for Agarose Gel Figure 2: The agarose gel is prestained with ViSafe Red Gel Stain. Various ladders and markers run at 1.5% TBE pre-stained agarose gel. The gel is visualized using transilluminator with (a) blue light; (b) UV light.

Viva SybrGreen Nucleic Acid Stain (10000X in DMSO)

Description

Viva Sybr Green Nucliec Acid Stain in one of type of novel generation of fluorescent nucleic acid gel stains used to detect double-stranded DNA in agarose and polyacrylamide gels. The detection limit using sthe Sybr Green stain is as low as 60pg of duoble stranded DNA using 300nm transillu- mination. Double stranded DNA as little as 20pg can be detected with 254nm ilumination. Single-stranded DNA and RNA can be detected as well although the sensitivity is lower. need approximately 100 to 300pg per well.

Features

The dye is noncytotoxic & nonmutagenic shown by Ames tests.

5-10X more sensitive than EtBr under UV and 8-20X more sensitive than EtBr in visible light.

Stable at room temperature for long-term storage. Stable to be microwaved or being heated. The working solution is stable at room temperature when kept in dark.

Suitable to stain dsDNA, ssDNA and RNA. Suitable to use in agarose gel or polyacrylamide gel. Compatible with downstream applications, such as gel recovery & cloning.

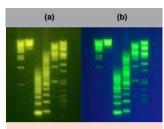
Easy precast gel staining, pre-staining and post-staining procedures.

Gel can be viewed with standard UV transilluminator, visible light gel reader or other gel imaging system.

Storage

Shipped at ambient temperature

Stored at 2-8°C or -20°C



Dves

Figure 1: The agarose gel is prestained with Viva Sybr Green® Nucleic Acid Stain. Various ladders and makers run at 1.5% TBE agarose gel. The gel is visualized using transilluminator with (a) blue light; (b) UV light.

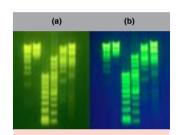


Figure 2: Samples and loading dye are mixed with Viva Sybr Green® Nucleic Acid Stain. Various ladders and makers run at 1.5% TBE agarose gel. The gel is visualized using transilluminator with (a) blue light: (b) UV light.

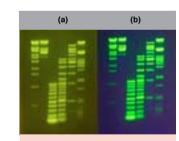


Figure 3: Various ladders and makers run at 1.5% TBE agarose gel. The agarose gel is post-stained with Viva Svbr Green® Nucleic Acid Stain. The gel is visualized using transilluminator with (a) blue light: (b) UV light.

Ordering Information

Catalog No	Description	Pack Size
SD0101	ViSafe Green Gel Stain (10000X in water)	500µl/pack
SD0103	ViSafe Red Gel Stain (10000X in water)	500µl/pack
SD0107	Viva SybrGreen Nucleic Acid Stain (10000X in DMSO)	1ml

qPCR Dye

Viva qGreen I Fluorescent Dye (Equivalent to SYBR® Green Dye)



Description

gPCR Viva gGreen I Fluorescent Dye (equivalent to SYBR® Green Dye) is a sensitive green fluorescent nucleic acid dye used for detection of double stranded DNA. The dye is widely used in non-specific detection of amplification in quantitative real-time PCR (qPCR) experiments. The detection is monitored by measuring the increase in fluorescence throughout the cycle.

Features

Easy and affordable:

Probes are not required, reduce assay setup and running cost; given that PCR primers are well designed and reaction is well characterized.

High sensitivity:

Increased fluorescence when bound to any double-stranded

Highly stable :

Stable during storage and under PCR condition, able to withstand repeated freeze-thaw cycles.

Versatile applications:

Can be used as a general double stranded DNA binding dye for common DNA quantification, melt curve analysis, etc.

Compatible with most system :

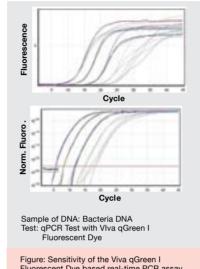
Compatible with major brands of gPCR instruments & enzyme systems.

Storage

Shipped at ambient temperature

Stored at 2-8°C or -20°C





rigure: Sensitivity of the viva defeet i
Fluorescent Dye based real-time PCR assay.
Amplification plot (cycle number versus
fluorescence) of known copies of DNA
standard (100ng - 0.01ng) was plotted with
three replicates.

1	100ng	Standard	5.47	Mean Ct: 5.537
2	100ng	Standard	5.59	
3	100ng	Standard	5.55	
4	10ng	Standard	8.07	Mean Ct: 8.077
5	10ng	Standard	8.03	
6	10ng	Standard	8.13	
7	1ng	Standard	11.63	Mean Ct: 11.653
8	1ng	Standard	11.87	
9	1ng	Standard	11.46	
10	0.1ng	Standard	15.09	Mean Ct: 14.880
11	0.1ng	Standard	14.58	
12	0.1ng	Standard	14.97	
13	0.01ng	Standard	18.34	Mean Ct: 17.567
14	0.01ng	Standard	17.76	
15	0.01ng	Standard	16.66	

Viva qGreen II Fluorescent Dye (Equivalent to EvaGreen® Dve)

Description

gPCR Viva gGreen II Fluorescent Dye (equivalent to EvaGreen® Dye) is one of the most sensitive dyes to detect double stranded DNA in quantitative real-time PCR (qPCR) experiments as well as high-resolution DNA melt curve analysis, vielding robust and reproducible results.

Features

Safer :

The dye is noncytotoxic & nonmutagenic for safe handling and easy disposal down to drain, completely impermeable to cell membrane.

Higher sensitivity:

Low PCR inhibitory and high concentration of dye used for maximal signal and high resolution DNA melt analysis.

Extremely stable:

Stable during storage and under PCR condition. No dye decomposition in PCR buffer at 95-100°C for 48 hours. Highly stable under alkaline or acidic condition and able to withstand repeated freeze-thaw cycles.

Versatile applications:

Used as a general double stranded DNA binding dye for DNA quantification, melt curve analysis and more.

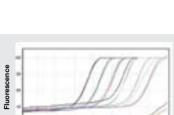
Excellent for qPCR and isothermal application :

Brighter and more sensitive than Viva aGreen I Fluorescent Dye (equivalent to SYBR® Green) for detecting amplification due to novel 'release on demand' DNA binding mechanism.

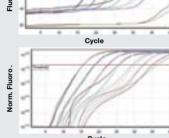
Storage

Shipped at ambient temperature

Stored at 2-8°C or -20°C



Dves



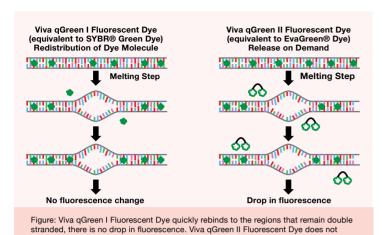
Sample of RNA: Dengue Virus RNA Test: RT-qPCR test with Viva qGreen II Fluorescent Dve

Figure: Sensitivity of the Viva qGreen II Fluorescent Dye based real-time PCR assay. Amplification plot (cycle number versus fluorescence) of known copies of DNA standard (100ng - 0.001ng) was plotted with three replicates.

No. Color	Name	туре	Ct	Ct Comment
1	100ng	Standard	15.62	Mean Ct: 15.623
2	100ng	Standard	15.61	
3	100ng	Standard	15.64	
4	10ng	Standard	19.06	Mean Ct: 18.953
5	10ng	Standard	18.94	
6	10ng	Standard	18.86	
7	1ng	Standard	22.48	Mean Ct: 22.557
8	1ng	Standard	22.63	
9	1ng	Standard	22.56	
10	0.1ng	Standard	25.50	Mean Ct: 25.913
11	0.1ng	Standard	26.12	
12	0.1ng	Standard	26.12	
13	0.01ng	Standard	29.59	Mean Ct: 29.737
14	0.01ng	Standard	30.27	
15	0.01ng	Standard	29.35	
16	0.001ng	Standard	32.66	Mean Ct: 32.533
17	0.001ng	NTC	32.58	
18	0.001ng	NTC	32.36	
19	NTC	NTC	35.17	
20	NTC	NTC	36.37	

Dyes

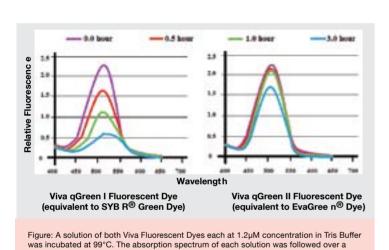
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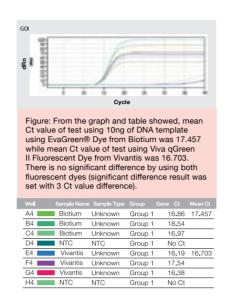


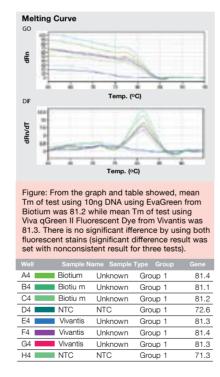
redistribute from the melted regions of single-stranded DNA back to double-stranded

DNA, resulting in a reduction of fluorescence. This difference gives the Viva qGreen II Fluorescent Dye the higher sensitivity in detecting amplification due to "release on

demand" DNA binding mechanism.







Ordering Information

period of 3 hours.

	Catalog No	Description	Pack Size
•	SD1101	Viva qGreen I Fluorescent Dye 20X in DMSO (equivalent to SYBR® Green Dye)	1ml/pack
	SD1103	Viva qGreen II Fluorescent Dye 20X in Water (equivalent to EvaGreen® Dye)	1ml/pack





Custom Polyclonal Antibody Development Peptide Route Whole Protein Route **Peptide Synthesis Service** Oligo Synthesis

Custom Polyclonal Antibody Development

Peptide Route

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Description

This package is designed for customers who have the peptide sequence and require us to begin from synthesizing it before immunization process. If the peptide can be supplied, Step 1 and 2 shall be skipped. For more details, please arrange a detailed consultation with us to determine the best package that suits your needs.

No.	Protocol details	Quality Control	Timeline	Notes
1	Peptide synthesis according to the sequence provided by customer	 Mass spectrometry to ensure molecular weight is correct HPLC to ensure purity greater` than 85%` 8mg for each peptide 	2 weeks	 Typical length for a peptide is approximately 15 amino acids Please notify if there is any special modification required
2	Protein carrier coupling (KLH/BSA)	No exposure to mercaptoethanol	2 days	
3	Antiserum preparation	ELISA titer for serum greater than 1:20,000 Serum volume is greater than 50ml per rabbit	10 weeks	Preimmune serum will be provided as control.
4	Protein A purified antiserum	>50mg total IgG	3 days	
5	Antigen affinity purification (serum)	Purified serum 40ml Antibody amount >3mg	3 days	ELISA titer for antiserum greater than 1:20,000

Note

- 1. Please notify if there is any modification required for peptide synthesis.
- Detailed consultation is required for phosphorylated antibodies, methylated or acetylated antibodies or special structures.

Requirements for customer-supplied peptide

- Peptide concentration more than 1mg/ml
- Peptide purity greater than 80%
- Peptide can be stored in leak-proof tube and ziplock bag. Deliver by using a Styrofoam box with ice packs is recommended

Deliverables upon completion

- 1. Preimmune serum (>50ml each rabbit)
- 2. Antigen affinity purified antibody (>3mg)
- 3. Test records: ELISA result

Whole Protein Route

Description

This package is designed for customers who can supply us with the whole protein. Polyclonal antibodies can be produced against soluble and non-soluble recombinant proteins, fusion proteins and gel strips. Please arrange a detailed consultation with us to decide the best package that suits your needs.

No.	Protocol details	Quality Control	Timeline	Notes
1	Antiserum preparation	ELISA titer for serum greater than 1:20,000 Serum volume is greater than 50ml per rabbit	10 weeks	Preimmune serum will be provided as control.
2	Protein A purified antiserum	>50mg total IgG	3 days	

Deliverables upon completio

- 1. Preimmune serum (>50ml each rabbit)
- 2. Protein A purified antiserum (>50mg)

Requirements for customer-supplied

full-length protein:

 Whole protein concentration greater than 1mg/ml is preferable

Peptide Synthesis Service

Description

Vivantis Technologies offers state of the art peptide synthesis technology with numerous platforms for various kinds of peptide, purity level, low to bulk quantities, excellent quality control and affordable.

What you get upon completion:

- Lyophilized form of peptide
- Certificate of Analysis with HPLC and MS analyses

Advantages

- High success rate and quality control
- Extensive range of peptide grades, quantity, formats and modifications according to customers' needs
- Competitive price
- Turnaround 3-4 weeks (subject to peptide complexity and length)

Services

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Oligo Synthesis

Type of Oligo Purification		tration	Sequence base
	nmole	OD	
HAP Purification	25	2	11-59 bases
	100	5	
	200	10	
PAGE Purification	100	2	11-59 bases
	200	5	
	400	10	
HPLC Purification	100	2	5-59 bases
	200	5	
	400	10	
Deoxylnosine with PAGE Purification	100	2	11-59 bases
	200	5	
	400	10	
Deoxylnosine with HPLC Purification	100	2	5-59 bases
	200	5	
	400	10	
DeoxyUridine with PAGE Purification	100	2	11-59 bases
	200	5	
	400	10	
DeoxyUridine with HPLC Purification	100	2	5-59 bases
	200	5	
	400	10	
Probe synthesis with different fluorophores with HPLC Purification	100	2	-
*FAM, HEX, Cy3, JOE, TAMRA, ROX etc.	200	5	
	400	10	









Compatible Ends Generated by Vivantis Restriction Endonucleases

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Enzyl	Acs	EcoF	Sse9	Bst2	Hind	Bsp1	Ama	AsiG	Bse1	Bsp1	Mro	Xma	BseF	Mlu	BstD	Ahl	Asp/	Asul	Xba	BssT	Erh	Bso3	BstN
									Se														
									nzym														
									a po	MK										*			
									nerated	∇⊥	ζ -										*	*	*
									gs ger	ن)	*	*	*	*	*	*	*	*	*			
									Z nucleotide 5' overhangs generated by enzymes	Recognition Sequence	5.	AAACGTT	TTAGGAA	ATACGAT	GRACGYC	0,000	090,9	0,000	TACGA	GT^MKAC	CA^TATG	T^TAA	AT^TAAT
									z nucleot	Fnzymo	LIIZYIIIC	4011	Bou14	BshVI	BssN	Hpall	HspAI	Mspl	Tagl	Fb/I	FauND I	Tru9 I	Vsp I

nzyn	F
by enz	7///
generated	(N
overhangs	ion Sequence
tide 5'	Recognit
3 nucleot	T 27/130

Enzyme	Recognition Sequence 5' 3'	ANT	GNC	GNC GWC	ΥΝ⊢	Z Z Z
Hinfl	G^ANTC	*				
AspS91	GAGNCC		*	*		
Bme18	GAGWCC			*		
Bpu101	CCTNAGC (-5/-2)				*	
Bse21	CCATNAGG				*	
Rsr2 I	CGAGWCCG		*	*		
Bsp1720 I	GC^TNAGC				*	
BstDEI	C^TNAG				*	
Bsf6 I	CTCTTC(1/4)	*	*	*	*	*

S	Z Z Z								*	
ızyme	ΑΝ⊢			*	*		*	*	*	
d by er	GWC	*	*			*			*	
nerated	GNC	*				*			*	
gs ger	ANT	*							*	
3 nucleotide 5' overhangs generated by enzymes	Recognition Sequence 5' — 3'	G^ANTC	G V G W C C	CCTNAGC (-5/-2)	CC^TNAGG	CGAGWCCG	GC^TNAGC	C^TNAG	CTCTTC(1/4)	
3 nucleot	Enzyme	Hinf I AspS9 I	Bme18	Bpu101	Bse21	Rsr2 I	Bsp1720 I	BstDEI	Bst6	

Blunt ends

* * * * *

* * * * *

Recognition Sequend 5' — 3'	GAGACTC	GATAATC	GTY^RAC	GTT^AAC	GAANN^NNT	TGGACCA	CMG^CKG	AGGACCT	CACAGTG	CAG^CTG	GT^AC	555,000	ATTT^AAAT	GACNN^NNR	AAT^ATT	GACAGTC
Enzyme	EcolCR I	EcoR V	Hind II	Hpa I	MroX I	Msp201	MspA11	Pce I	PspC I	Pvu II	Rsal	Smal	Smil	SmiM I	Ssp I	Zra I
Recognition Sequence 5' — 3'	TGC^GCA	CCGCTC(-3/-3)	AG^CT	AGT^ACT	GGN^NCC	GATNN^NNATC	00,00	GTA^TAC	YAC^GTR	90,90	GACNN^NNGTC	TACAGTA	CACGTC(-3/-3)	TCG^CGA	000,000	TTT^AAA
Enzyme	Acc161	AccBS I	Alu I	BmcA I	Bmil	Bse8 I	Bsnl	BssNAI	BstBAI	BstFNI	BstPAI	BstSNI	Btr I	BtuM I	Din I	Dra I

Z Z					*				*	*
R					*			*	*	*
N O					*		*	*	*	*
၁၅					*	*	*	*	*	*
N N				*	*				*	*
90		*	*	*	*				*	*
AT	*				*			*	*	*
Recognition Sequence 5'	CGATACG	0,000	0,000	GAATGC(1/-1)	GACNNNNVNNGTC	55,050	ACTGG(1/-1)	CGRYACG	GCAATG(2/0)	GGATG(2/0)
Enzyme	BpvU I	AspLEI	BstHHI	Pct I	DseD I	Sfr303 I	Bse11	BstMC I	Bse3D I	BstF51

Z Z Z	*	*	*	*	*
Recognition Sequence 5' — 3'	CCNNNNNANGG	CCANNNN'NTGG	CACNNN'GTG	GCCNNNNVGGC	GGCCNNNNVGGCC
Enzyme	Afil	AccB7 I	Dra III	Bg/I	Sfil

4 nucleotide 3' overhangs generated by enzymes

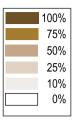
Aatll GACGT^C * * Psp124B1 GAGCT^C * * BstNS1 GCATG^C * * Bmt1 GCATG^C * * Mh1 GDGCH^C * * BstH21 RGCGC^A * * Kpn1 GGGCC^AC * * Kpn1 GGGCC^AC * * Kpn1 GRGCY^AC * * Spt1 CCTGCA^G * * Spt1 CCTGCA^G * * Spt1 GCTGCA^T * * BstX1 CAANNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN	Enzyme	Recognition Sequence 5' — → 3'	ACGT	ACGT AGCT CATG CTAG DGCH GCGC GGC	CATG	CTAG	ресн	2929	99
## GAGCT^C RCATG^Y RCATG^Y GCATG^C GCTAG^C GCTAG^C RGCC^A GGCC^A CTGCA^A GGCC^A CTGCA^A GCCA^A CTGCA^A GCCA^A CTGCA^A	AatII	GACGT^C	*						
RCATG^\(\frac{1}{2}\)	Psp124B I	GAGCT^C		*					
GCATG^C GCTAG^C GCTAG^C GCTAG^C GCTAG^C GCCC^C GGCC^C GGCC	BstNSI	RCATG^Y			*				
GCTAG^C GDGCH^C RGCGC^Y GGGCC^C GGTAC^C GGTAC^C GTGCA^G CTGCA^G CTGCA^G ATGCA^T GWGCW^C ATGCA^T GWGCW^C CAANNNN^NTGG * * * *	Sph I	GCATG^C			*				
GDGCH^C RGCGC^Y GGGCC^C GGTAC^C GGTAC^C GTGCA^G CTGCA^G CTGCA^G ATGCA^T GWGCW^C AANNNN^NTGG * *	Bmt I	GCTAG^C				*			
RGCGC^Y GGGCC^C GGTAC^C GGTAC^C GTGCA^G CTGCA^G CTGCA^G ATGCA^T GWGCW^C CAANNNN^NTGG	Mh/I	GDGCH^C		*			*		*
GGGCC^C GGTAC^C GGTAC^C GRGCY^C CTGCA^G CCTGCA^G ATGCA^T AGGCW^C * * * * * * * * * * * * * * * * * * *	BstH2 I	RGCGC^Y						*	
GGTAC^C GRGCY^C GTGCA^G CCTGCA^G ATGCA^T ATGCA^T GWGCW^C CAANNNN^NTGG * * * *	Apal	0,00999							*
GRGCY^C CTGCA^G CCTGCA^G CCTGCA^G ATGCA^T GWGCW^C * * * * * * * * * * * * * * * * * * *	Kpn1	GGTAC^C							
CTGCA^G CCTGCA^G CCTGCA^T ATGCA^T GWGCW^C * *	Fri0 I	GRGCY^C		*					*
CCTGCA^G ATGCA^T GWGCW^C CAANNNN^NTGG * *	PstI	CTGCA^G							
ATGCA^T GWGCW^C CAANNNN^NTGG * * *	Sbf1	CCTGCA^GG							
	Zsp2 I	ATGCA^T							
CAANNNN^NTGG * * *	Bbv121	GWGCW^C		*					
	BstX I	CAANNNNN^NTGG	*	*	*	*	*	*	*

C/G/ A/G/	4 4	D / A	<i>\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ </i>)

4 nucleotide 5' overhangs generated by enzymes

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Legend:



Indicates activity (%) of restriction endonucleases in different Vivantis buffer systems



Restriction endonuclease that requires specific buffer to perform at 100% activity



High enzyme concentration may result in Star Activity



10mM Tris-HCI (pH 7.5 at 30°C),10mM MgCl₂, and 100µg/ml BSA.



10mM Tris-HCI (pH 7.5 at 30°C),10mM MqCl_o. 50mM NaCl, and 100µg/ml BSA.



50mM Tris-HCI (pH 7.5 at 30°C),10mM MgCl₂, 100mM NaCl, and 100µg/ml BSA.



10mM Tris-HCI (pH 8.5 at 30°C),10mM MgCl₂, 100mM KCl, and 100µg/ml BSA.



30mM Tris-acetate (pH 7.9 at 30°C),10mM Mg-acetate, 60mM K-acetate, and 100μg/ml BSA.



12.5mM Tris-acetate (pH 7.6 at 30°C), 5mM Mg-acetate, 50mM K-acetate, 3.5mM 2-mercaptoethanol and 25µg/ml BSA

1.0X UB

25mM Tris-acetate (pH 7.6 at 30°C), 10mM Mg-acetate, 100mM K-acetate, 7mM 2-mercaptoethanol and 50µg/ml BSA

1.5X UB

37.5mM Tris-acetate (pH 7.6 at 30°C),15mM Mg-acetate, 150mM K-acetate, 10.5mM 2-mercaptoethanol and 75µg/ml BSA

50mM Tris-acetate (pH 7.6 at 30°C),20mM Mg-acetate. 200mM K-acetate, 14mM 2-mercaptoethanol and 100µg/ml BSA



Buffer AccB1 I

10mM Tris-HCl (pH 7.5 at 25°C),10mM MgCl₂, 100mM KCI, and 100µg/ml BSA.

Buffer Ama87I

10mM Tris-HCI (pH8.5), 10mM MgCl₂, 150mM NaCl, 100µg/ml BSA.

Buffer Bgl I

20mM Tris-HCl (pH 8.5),10mM MgCl₂, 200mM NaCl and 1mM DTT.

Buffer Dra III

10mM Tris-HCI (pH 7.6),10mM MgCl₂, 200mM KCI, and 100µg/ml BSA.

Buffer EcoR I

50mM Tris-HCI (pH 7.5 at 30°C),10mM MgCl₂, 100mM NaCl, 0.02% Triton X-100 and 100µg/ml BSA.

Buffer EcoR V.

10mM Tris-HCI (pH 8,5 at 30°C). 10mM MgCl₂, 100mM NaCl and 100μg/ml BSA.

Buffer Mbo II.

33mM Tris-acetate (pH 7.9 at 30°C), 10mM Mg-acetate, 66mM K-acetate, and 1mM DTT.

Buffer Ssp I

10mM Tris-HCl (pH 7.6 at 30°C),10mM MgCl₂, 100mM KCl, and 100µg/ml BSA.

Diluent Buffers:

Viva Buffer A

10mM Tris-HCI (pH 7.4 at 25°C), 50mM KCI, 0.1mM EDTA, 1mM DTT, 200µg/ml BSA and 50% glycerol

NOTE:

DILUTION OF RESTRICTION ENZYMES

Dilution buffer (Diluent) is provided for preparation of diluted enzyme. The diluted enzyme must be stored at -20°C in order to retain its activity. When properly diluted and stored, the diluted enzyme retains 50% to 100% of its activity up to one month of storage at -20°C.

For long term storage and stability, the user is strongly advised to use the storage buffer of the particular enzyme (see "Storage Buffer" for the components) for dilution instead of Dilution Buffer (Diluent), for diluted enzyme preparation.

Example:

Initial concentration of restriction enzyme = 10u/μI

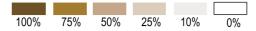
Desired concentration of restriction enzyme = 2u/µI

Therefore, a 5x dilution is to be performed.

Mix $5\mu I$ of restriction enzyme (10u/ $\mu I)$ with $20\mu I$ of Dilution Buffer (Diluent) or Storage Buffer thoroughly with pipette and store diluted enzyme at -20°C. Please avoid repeated freeze-

Note: **DO NOT VORTEX** the enzyme mixture Please perform all dilutions on ice at all times.

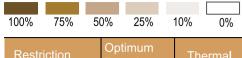
Legend:



Restriction	Optimum	Thermal	Activity (%)
Endonuclease	Reaction	Inactivation	V1 V2 V3 V4 V5 UB
	Temperature		0.5X 1.0X 1.5X 2.0X
Aat II*	37°C	65°C	
Acc16 I	37°C	65°C	
Acc65 I	37°C	65°C	
AccB1 I*	37°C	65°C	
AccB7 I [★]	37°C	65°C	
AccBS I	37°C	65°C	
Acl I	37°C	65°C	
Acs I	50°C	NO	
Afi l	55°C	80°C	
Ahl I	37°C	NO	
Alu I	37°C	65°C	
Ama87 I	37°C	65°C	
Apa I	37°C	65°C	
AsiG I	37°C	65°C	
AspA2 I	37°C	80°C	
AspLE I	37°C	NO	
AspS9 I	37°C	65°C	
AsuHP I	37°C	65°C	
AsuNH I	37°C	65°C	
BamH I	37°C	65°C	
Bbv12 I	37°C	NO	
Bg/ I [★]	37°C	65°C	
Bgl II	37°C	NO	
BmcA I	37°C	65°C	
Bme18 I	37°C	65°C	
BmeR I	37°C	65°C	
Bmi I	37°C	65°C	
BmrF I	37°C	NO	
Bmt I	37°C	65°C	
<i>Bpu</i> 10 I *	37°C	80°C	
Bpu14 I	37°C	65°C	
BpuM I	37°C	65°C	
BpvU I	37°C	80°C	
Bse1 I	65°C	80°C	
Bse118 I	65°C	NO	
Bse21 I	37°C	NO	
Bse3D I	60°C	NO	
Bse8 I*	60°C	NO	
BseP I	50°C	65°C	

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Legend:



100% /5% 5	0% 25% Optimum	10% 0%	Activity (%)
Restriction	Reaction	Thermal	LID.
Endonuclease	Temperature	Inactivation	V1 V2 V3 V4 V5 0.5X 1.5X 2.0X
BseX3 I	50°C	NO	
BshV I	37°C	65°C	
Bsn I	37°C	65°C	
Bso31 I	55°C	NO	
Bsp13 I	50°C	65°C	
Bsp1720 I	37°C	NO	
<i>Bsp</i> 19 I	37°C	65°C	
BssM I	37°C	NO	
BssN I	37°C	NO	
BssNA I	37°C	NO	
BssT1 I	60°C	NO	
Bst2B I	60°C	NO	
Bst2U I	60°C	NO	
Bst4C I	65°C	NO	
Bst6 I	65°C	80°C	
BstAU I	37°C	80°C	
BstBA I	65°C	NO	
BstDE I	60°C	NO	
BstDS I	65°C	NO	
BstEN I	65°C	80°C	
BstF5 I	65°C	NO	
BstFN I	60°C	NO	
BstH2 I	65°C	NO	
BstHH I	50°C	NO	
BstMA I	55°C	NO	
BstMB I	65°C	80°C	
BstMC I	50°C	NO	
BstNS I	37°C	65°C	
BstPA I	65°C	NO	
BstSN I	37°C	NO	
BstV2 I	55°C	NO	
BstX I	37°C	65°C	
BstX2 I	60°C	NO	
Btr I	60°C	80°C	
BtuM I	37°C	NO	
CciN I	37°C	65°C	
Din I	37°C	65°C	
Dra I	37°C	65°C	
Dra III [★]	37°C	65°C	

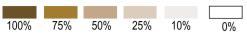
Legend:

100%	75%	50%	25%	10%	0%

Restriction	Optimum	Thormol	Activity (%)
Endonuclease	Reaction	Thermal Inactivation	V1 V2 V3 V4 V5 UB
0.01	Temperature		0.5X 1.0X 1.5X 2.0X
DseD I	37°C	NO	
EcoICR I	37°C	65°C	
EcoR I*	37°C	65°C	
EcoR V*	37°C	NO	
Erh I	37°C	65°C	
FauND I	37°C	65°C	
Fbl I	55°C	NO	
Fokl	37°C	65°C	
FriO I	37°C	65°C	
Hind II	37°C	65°C	
Hind III	37°C	65°C	
Hinf I	37°C	NO	
Hpa I *	37°C	65°C	
Hpa II	37°C	NO	
HspA I	37°C	NO	
Kpn I*	37°C	NO	
Ksp22 I [★]	37°C	65°C	
Mbo II	37°C	65°C	
Mhl I *	37°C	65°C	
Mlu I	37°C	65°C	
Mnl I	37°C	65°C	
MroN I	37°C	NO	
MroX I	37°C	65°C	
Msp I	37°C	65°C	
Msp20 I	37°C	65°C	
MspA1 I	37°C	65°C	
Pce I	50°C	80°C	
Pct I	37°C	65°C	
Psp124B I	37°C	65°C	
PspC I	37°C	65°C	
PspE I [★]	37°C	65°C	
PspOM I	37°C	65°C	
Pst I [★]	37°C	NO	
Pvu II*	37°C	NO	
Rsa I	37°C	NO	
Rsr2 I	37°C	65°C	
Sal I *	37°C	65°C	
Sbf I *	37°C	80°C	
Sfi I	50°C	65°C	

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Legend:



Restriction	Optimum	Thermal	Activity (%)
Endonuclease	Reaction Temperature	Inactivation	V1 V2 V3 V4 V5 UB 0.5X 1.0X 1.5X 2.0X
SfaN I	37°C	NO	
Sfr274	50°C	65°C	
Sfr303	37°C	65°C	
Sma I	25°C	65°C	
Smi I	37°C	65°C	
SmiM I	37°C	65°C	
Sph I	37°C	65°C	
Sse9 I	55°C	65°C	
Ssp I*	37°C	65°C	
Taq I	65°C	NO	
Tru9	65°C	NO	
<i>Tth</i> 111 I [★]	65°C	NO	
Vha464 I	37°C	65°C	
Vne I	37°C	65°C	
Vsp I	37°C	65°C	
Xba I	37°C	65°C	
Xma *	37°C	65°C	
Zra I *	37°C	80°C	
Zsp2 I	37°C	65°C	

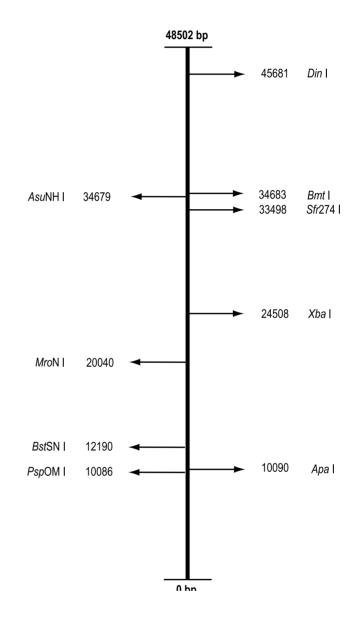
Troubleshooting-Restrictions Endonucleases

Problem	Probable reasons and recommended solutions
No cleavage or partial digestion	Non-optimal reaction conditions The activity of an enzyme is governed by the following conditions: reaction temperature, buffer ionic strength, pH and concentration of Mg²+. Optimum reaction conditions for each enzyme are specified in the product datasheet. Use the recommended reaction buffer supplied with the enzyme. In some cases, the presence of BSA greatly enhances the activity of a restriction endonuclease.
	Nature of DNA The amount of enzyme required for complete digestion depends on the nature of the DNA substrate (ie. linear, supercoiled, number of recognition sites in substrate, number of nucleotides flanking the recognition site). Supercoiled plasmid DNA usually requires 5-10u/µg of DNA for complete digestion.
	Enzyme sensitivity to substrate methylation
	Methylation of nucleotides in the recognition sequence may partially or completely block the restriction endonuclease cleavage. Check for the presence of methylation (e.i Dam, Dcm, EcoK I, EcoB I, CpG or CpNG) in the test DNA and whether the enzyme used is sensitive to that methylation.
	Presence of enzyme inhibitors in substrate DNA
	The activity of a restriction endonuclease may be partially or completely inhibited by contaminants such as solvents, detergents and salts present in the DNA preparations. Check the activity of the enzyme on lambda DNA alone and lambda DNA mixed with the test DNA. If the activity on lambda DNA alone is as indicated in the product datasheet, while the mixed DNA is digested poorly, the test DNA should be repurified.
	Lack of recognition sequences
	If the test DNA is completely undigested, ensure that the DNA sequences recognized by the restriction endonucleases are present in the DNA.
	Improper dilution of enzyme
	If dilution of enzyme prior to addition to the reaction mixture is required, the dilution buffer recommended in the product datasheet should be used. Enzyme activity may be lost if inappropriate dilution buffer is used. Mix the enzyme thoroughly but gently, and do not vortex dilutions or reactions containing restriction endonucleases.
	Lost of activity
	The enzyme has most probably been inactivated due to improper handling or storage.
Presence of additional bands	Partial digestion Complete digestion can often be achieved by increasing the incubation time or the amount of enzyme or performing digest in the presence of BSA.

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Problem	Probable reasons and recommended solutions
Presence of additional bands	Enzyme star activity Star activity is characterized by the presence of additional bands below the expected bands. This phenomenon occurs when restriction enzymes are used at low ionic strength or high pH values in the reaction buffer, prolonged incubation or large excess of enzyme with respect to DNA, high glycerol content (>5%) or presence of organic solvents, substitution of cofactor Mg²+ with other divalent cations such as Mn²+. Apart from the normally recognized sequences, the enzymes cleave at other sites which differ in some positions from the canonical site. Star activity can be avoided by using less amount of enzyme and glycerol (no more than one tenth volume) and using the recommended reaction buffer.
	Contamination of substrate DNA or restriction endonuclease
	Check the banding pattern of undigested test substrate (without addition of enzyme) and digestion of the test substrate with other restriction endonucleases. If the problem still occurs, the substrate DNA and enzyme preparation might have become contaminated due to improper handling. Contamination in the reaction buffer may also give rise to such problems.
Inefficient ligation of fragments	Restriction enzyme remains active
	Most restriction endonucleases can be inactivated thermally by incubation at 65°C or 80°C for 20 minutes. Check the mode of thermal inactivation for the enzyme. If the enzyme cannot be inactivated thermally, perform phenol purification and/or ethanol precipitation of the fragments after restriction digestion.
	Low ligase concentration
	Use higher concentration of ligase for ligation of blunt and one nucleotide overhang-DNA fragments. The recommended concentration is at least 20- $40 \text{u}/\mu\text{g}$ of fragments with addition of 10% PEG in ligation mixture.
	Contamination with non-specific nucleases
	Non-specific nucleases may be present in the restriction enzyme, ligase or DNA preparation due to improper handling. Use new set of reagents.
	Deterioration of buffer components
	Use fresh ligation buffer as far as possible as ATP and DTT are easily subjected to degradation. Avoid multiple frozen and thawed cycles of ligation buffer.

Lambda DNA (cl857Ind 1 Sam7) Restriction Map



Enzymes which cut once on Lambda DNA

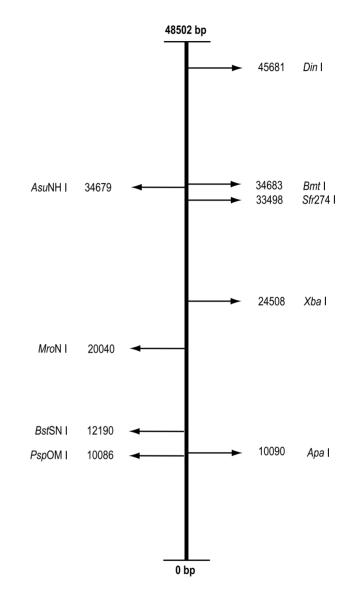
Restriction	Cleave	
Endonucleas	e Position	
Apa I	10090	
AsuNH I	34679	
Bmt I	34683	
BstSNI	12190	
Din I	45681	
MroN I	20040	
PspOM I	10086	
Sfr274 I	33498	
Xba I	24508	
Xba I	24508	

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Restriction Endonucleas	Cleave e Positior	1				Restriction Endonucleas	Cleave e Position					-
Acc65 I	17053	18556				Acll	13530	16291	22581	22596	24643	
AspA2 I	24322	24396				Acri	43489	43393	22301	22000	27070	
Bse21 I	26718	34319				Bpu14 I	18049	25885	27981	29151	30397	
BseX3 I	19944	36654				<i>_</i> ,,	42638	34332			00001	
Bso31 I	11418	42709				BstPA I	8924	9398	13516	15416	36929	
EcoCR I	24774	25879				DSULVI	48156	37893	13310	13410	30323	
Kpn I	17057	18560				FauND I	27631	29884	33680	36113	36669	
Psp124B I	24776	25881				7 dans 1	40132	38358	00000	00110	00000	
Sall	32745	33244				FriO I	585	10090	19767	21574	24776	
Tth111	11205	36123					39457	25881				
		00.20				Hind III	23130	25157	27479	36895	37459	
BpvU I	11936	26257	35790				44141	37584			07 100	
BssNA I	15262	18836	19475			Mlu I	458	5548	15372	17791	19996	
DseD I	5122	9110	11096				22220	20952				
PspC I	26531	41484	42364									
Sma I	19399	31619	39890			Ama87 I	4720	19397	20999	27887	31617	
Vha464 l	6540	12618	42630			/iiiiaor i	33498	38214	39888	21001	01011	
Xma	19397	31617	39888			Bst2B I	20356	25572	27956	29425	34430	
Απα Ι	10001	01017	33000				35219	42416	42737			
						Ksp22 I	8844	9361	13820	32729	37352	
Bsp19 I	19329	23901	27868	44248			43682	46366	47942			
Sfr303 I	20323	20533	21609	40389								
Vne I	5619	21798	27173	40216		BmeR I	6403	11243	12482	12920	16593	
							18549	23465	30472	44674		
BamH I	5505	22346	27972	34499	41732	BstEN I	13513	21296	22381	25178	25227	
BstAU I	5220	6142	15855		32496		35525	38272	41846	47217		
BtuM I	4592	28052	31705	32409		Fbl I	2191	15261	18835	19474	31302	
EcoR I	21226	26104	31747	39168	44972		32746	33245	40202	42922		
Rsr2 I	3801	6042	13984	19289	22243							
Sbfl	2560	2824	11839	19837	37005	Aat II	5109	9398	11247	14978	29040	
Zrm	16423	18686	25687	27265	32804		40810	41117	42251	45567	45596	
21111	10420	10000	20001	21200	02004	BssT1 I	19329	21211	23901	24322	24396 44248	
						Dra III	27868 2959	28793 5618	35016 6640	36050 9004	14482	
D/ III	445	20425	05744	20402	20754	DIA III	30370	31914	41484	47317	48439	
Bgl II	415 38814	22425	35/11	38103	38/54	Erh I	19329	21211	23901	24322	24396	
BseP I	3522	4126	5627	14815	166/0	_,,,,	27868	28793	35016	36050	44248	
<i>D</i> 361 1	28008	7120	0021	17013	10049	Tag II	1310	2388	6690	9566	13135	
Bsp1720 I	10298	10683	11662	16519	20745		16604	17905	18592	22231	45067	
DSP 1120 1	39451	10003	11002	10019	20140	Zra I	5107	9396	11245	14976	29038	
Pce I	12436	31480	32999	39994	40598		40808	41115	42249	45565	45594	
	40616	31100	32000	30004	.0000							
Sph I	2216	12006	23946	24375	24378							
	39422					Thoro are no	rootriotion	citoc for	the follow	ina onzvi	moo:	

There are no restriction sites for the following enzymes: Ahl I, CciN I, Sfi I, Smi I

T7 DNA Restriction Map



Enzymes which cut once on Lambda DNA

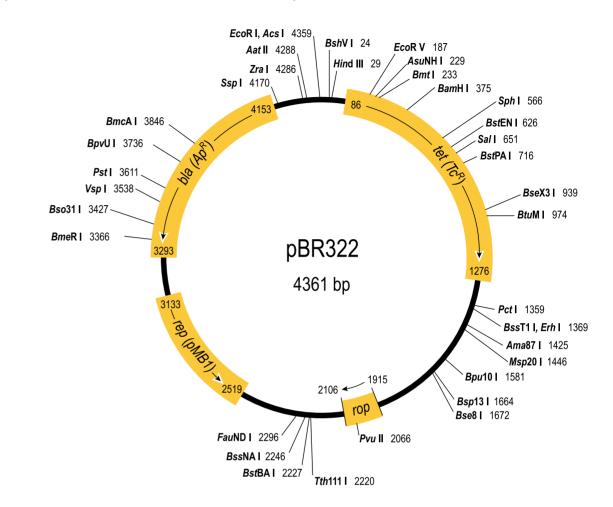
Restriction	Cleave	
Endonuclease	Position	
Apa I	10090	
AsuNH I	34679	
Bmt I	34683	
BstSNI	12190	
Din I	45681	
MroN I	20040	
PspOM I	10086	
Sfr274 I	33498	
Xba I	24508	

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Restriction Endonuclease	Cleave Position					Restriction Endonuclease	Cleave Position				
Ahll	7694	17279				AccB7 I	9316	13374	18766	25857	34825
AsiG I	35710	37465				DNI	36029	36392	38034	11210	24291
Bgl I Bst2U I	13523 2367	35938 8189				BssN I	339 24731	10941 25656	11084 35505	11349	24291
BstX2	11515	36087				BssNA I	7679	19513	23331	24464	27139
Din I	11085	35506				Bookkit	28491	28771	39629		200
Msp20 I	35540	35941				Btr I	1667	2139	3698	20875	23776
							26244	34062	36037		
4 401	00004	05400	22222			Zsp2 I	2610	8753	19990	21979	30001
AspA2 I	20004	35163	33222				38325	39138	39297		
Bse118 I BshV I	23393 22857	35710 27100	37465 37080								
BtuM	3271	27100	28719			BpuM I	537	651	2660	14699	15674
Pvu II	11483	13623	24273				21840	23193	28362	37722	
Xba I	12830	22925	34294			Dra I	276	441	5973	6501	10723
							16871	20298	31371	39748	
Ama87 I	10512	15902	26119	30082							
BmcA I	15517	34102	38852	38994		BmrF I	537	651	2367	2660	8189
							14699 37722	15674	21840	23193	28362
Acc65 I	37	5613	9188	23741	39814	BstX I	3870	9633	10105	13480	13881
Kpn I	41	5617	9192	23745	39818		15397 38965	15736	27959	29443	34663
BssM I	8311	8414	11515	14354	35684	DseD I	538	6209	17203	18055	33010
	36087						2114	6964	17588	32160	34747
BstMB I	8311 36087	8414	11515	14354	35684		34921				
BstPA I	1130 38618	2088	16069	29004	29733						
Acc16 I	4138	11228	15665	25454	28866						
	29804	35315									
Bpu14 I	1121	6017	12160	19215	22862						
D 04	22880	31437	0.400	7450	40004						
Bse8 I	620 19989	3038 37622	3183	7150	18904						
FauND I	5675	6363	14307	19955	22965		e no restrict				
T dunib I	27726	33731	14007	10000	22303	Apa I, B EcoR I.	amH I, Bpv EcoR V, Hir	o i, <i>u</i> sexa id III, <i>Mro</i> N	o i, <i>B</i> sp13 I I , <i>P</i> sp124	i, c <i>ci</i> n i, IB I, <i>Psp</i> O	ECOIUR I, MI, Pst I.
		00101					f Sfr274				, ,

EcoR I, EcoR V, Hind III, MroN I, Psp124B I, PspOM I, Pst I, Sal I, Sbf I, Sfr274 I, Sfr303 I, Sma I, Sph I, Xma I

pBR322 DNA Restriction Map



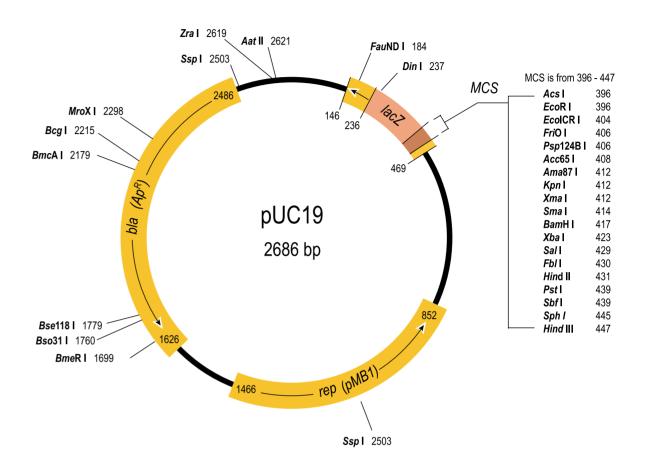
Enzymes which cut once on pBR322 DNA

Restriction Endonuclease	Cleave Position	Restriction Endonuclease	Cleave Position
Aat II	4288	BstEN I	626
Acsl	4359	BstPA I	716
Ama87 I	1425	BtuM I	974
AsuNH I	229	EcoR I	4359
BamH I	375	EcoR V	187
BmcA I	3846	FauND I	2296
BmeR I	3366	Hind III	29
Bmt I	233	Msp20 I	1446
Bpu10 I	1581	Pct I	1359
BpvU I	3736	Pst I	3611
Bse8 I	1672	Pvu II	2066
BseX3 I	939	Sall	651
BshV I	24	Sph I	566
Bso31 I	3427	Ssp I	4170
Bsp13 I	1664	Tth111 I	2220
BssNA I	2246	Vsp I	3538
BssT1 I, Erh I	1369	Zra I	4286
BstBA I	2227		

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Restriction Endonuclease	Cleave Position						
AccB7 I	1321	1370					
AccBSI	2406	4207					
Bse3D I	3427	3601					
Bst6 I	2357	4161					
BstDS I	528	1447					
DseD I	2168	2581					
FbII	652	2245					
FriO I	475	489					
Hind III	653	3907					
MroX I	2033	3965					
Taq II	4052	4069					
Bg/ I	935	1169	3486				
Bst2B I	2646	4030	4337				
BstMA I	2167	3427	4203				
BstV2 I	730	1593	4344				
Dra	3232	3251	3943				
Rsal	165	2281	3846				
Vne I	2289	2787	4033				
Acc16 I	262	1358	1456	3588			
AcI	901	1800	3592	3965			
BstNS I	566	1820	2112	2477			
Din I	415	436	550	1207			
MroN I	401	769	929	1283			
BssN I	414	435	549	1206	3903	4285	
Bst2U I	131	1060	1443	2501	2622	2635	
MspA1 I	1141	2066	2185	2815	3060	4001	
SmiM I	1031	1462	1657	2048	3618	3777	
Bse118 I	160	401	410	769	929	1283	
	3446						
BstMC I	289	656	942	2389	2813	3736	
	3885	000	050	4407	4000	0570	
Taq I	24	339	652	1127	1268	2573	
	4017						
D/ 40 I	000	504	4470	4.400	0000	0704	
Bbv12 I	280	591	1178	1469	2293	2791	
Dm of 0.1	3952	4037	1120	1420	1404	1760	
Bme18 I	799 3504	887 3726	1136	1439	1481	1760	There are no restriction sites for the following
BstDE I	1581	1743	2283	2748	3157	3323	enzymes:
DSIDE I	3863	4289	2200	2170	3137	0020	Acc65 I, Ahl I, Apa I, AsiG I, AspA2 I, Bgl II, Bpu14 I, Bse21 I, BseP I, Bsp1720 I, Bsp19 I,
BstX2 I	375	1667	3114	3125	3211	3223	BstAU I,BstSN I, BstX I, Btr I, CciN I, Dra III,
	3991	4008					EcolCR I, Hpa I, Kpn I, Ksp22 I, Mlu I, Pce I,
Sse9 I	58	251	1319	1333	3233	3539	Psp124B I, PspC I, PspE I, PspOM I, Rsr2 I, Sfi I, Sbf I, Sfr274 I, Sfr303 I, Sma I, Smi I, Vha464 I,
	3794	4359					Sbi 1, Sir214 1, Sir303 1, Siria 1, Sirii 1, Viia464 1, Xba 1, Xma 1, Zsp2

pUC19 DNA Restriction Map



Enzymes which cut once on pUC19 DNA

Restriction Endonuclease	Cleave Position	Restriction Endonuclease	Cleave Position	
Aat II	2621	Hind II	431	
Acc65 I	408	Hind III	447	
Acs I	396	Kpn I	412	
Ama87 I	412	MroX I	2298	
BamH I	417	Psp124B I	406	
BmcA I	2179	Pstl	439	
BmeR I	1699	Sall	429	
Bse118 I	1779	Sbf I	439	
Bso31 I	1760	Sma I	414	
Din I	237	Sph I	445	
EcolCR I	404	Ssp I	2503	
EcoR I	396	Xba I	423	
FauND I	184	Xma I	412	
FbH	430	Zra I	2619	
FriO I	406			

Restriction

Technical Information

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Cleave

Endonuclease	Position	1				
Acc16 I	258	1921				
Acll	1925	2298				
Bgll	251	1819				
Bme18 I	1837	2059				
BpvU I	279	2069				
Bse3D I	1760	1934				
Pvu II	308	630				
DseD I	97	914				
				_		
AccBS I	498	739	2540			
BssN I	236	2236	2618			
Bst2B I	979	2363	2670			
Bst6 I	296	690	2494			
BstH2 I	239	684	1054			
BstNS I	41	445	810			
Dra I	1565 169	1584 410	2276 2179			
Rsa I	1951	2110	2469			
SmiM I	177	1120	2366			
Vne I	577	636	1871			
Vsp I	311	030	10/1			
AccB1 I	235	408	550	1647		
BstMA I	45	1760	2536	2689		
Taq I	400	430	906	2350		
Bbv12	181	406	1124	2285	2370	
Bst2U I	355	546	834	955	968	
BstF5 I	83	327	1658	1839	2126	
BstMC I	279	722	1146	2069	2218	
Fok I Mhl I	90 181	334	1665 1124	1846	2133	
IVITII 1	101	406	1124	2285	2370	
Afi I	53	654	828	846	1012	1291
AspS9 I	286	1741	1820	1837	2059	2675
BstDE I	171	1081	1490	1656	2196	2622
Hinf I	427	641	706	781	1177	1694
MspA1 I	114	308	630	1148	1393	2334

There are no restriction sites for the following enzymes:

AccB7 I, Ahl I, Apa I, AsiG I, AspA2 I, AsuNH I, Bgl II, Bmt I, Bpu10 I, Bpu14 I, Bse21 I,

Bse8 I, BseP I, BseX3 I, BshV I, Bsp13 I, Bsp1720 I, Bsp19 I, BssNA I, BssT1 I, BstAU
I, BstBA I, BstDS I, BstEN I, BstPA I, BstSN I, BstV2 I, BstX I, Btr I, BtuM I, CciN I, Dra

III, EcoR V, Erh I, Hpa I, Ksp22 I, Mlu I, MroN I, Msp20 I, Pce I, Pct I, PspC I, PspE I,

PspOM I, Rsr2 I, Sfi I, Sfr274 I, Sfr303 I, Smi I, Tth111 I, Vha464 I, Zsp2 I

Amplication of various DNA fragments using Vivantis DNA Amplification Reagents

Taq DNA Polymerase / Chromo *Taq* DNA Polymerase / At*Taq* DNA Polymerase / Chromo At*Taq* DNA Polymerase (PL1202 - PL1206 / PL3201 - PL3206)

Product Size	1.5kb	5kb	8kb	10kb	15 and 20kb
ViBuffer (1X)	Α	Α	S	S	S
MgCI ₂	1.5mM	1.5mM	-	-	-
dNTP mix	0.2mM	0.25mM	0.25mM	0.36mM	0.36mM
Primers	0.2μΜ	0.8μΜ	0.4µM	0.6µM	0.6µM
Lambda DNA	0.03μg	0.03µg	0.02μg	0.06µg	0.15μg
Taq DNA Polymerase	2.0u	2.5u	2.5u	2.5u	2.5u
DMSO / Formamide	-	3%	3%	3%	3%

Top up with sterile dH2O to 25µl.

Product Size	1.5kb	5kb	8kb	10kb	15 and 20kb
Denaturation	94°C, 1min	94°C, 2min	94°C, 2min	94°C, 2min	94°C, 2min
Denaturation	94°C, 30s	94°C, 12s	94°C, 12s	94°C, 12s	94°C, 12s
Annealing	52°C, 30s	60°C, 30s	65°C, 30s	59°C, 30s	56°C, 30s
Extension / 1kb	72°C, 1min	68°C, 4min	68°C, 5min	68°C, 12min	68°C, 15min
Cycles	25	30	30	35	10
Final Extension / Extension	72°C, 5min	68°C, 10min	68°C, 10min	68°C, 7min	-
Denaturation	-	-	-	-	94°C, 12s
Annealing	-	-	-	-	56°C, 30s
Extension / 1kb ¹	-	-	-	-	68°C, 15min
Cycles	-	-	-	-	20
Final Extension	-	-	-	-	68°C, 7min

MaxTaq DNA Polymerase / Chromo MaxTaq DNA Polymerase (PL2201 - PL2206)

Product Size	8kb	10 and 12kb	15 and 20kb	30 and 40kb
ViBuffer (1X)	S	S	S	S
MgCl ₂	-	-	-	-
dNTP mix	0.25mM	0.36mM	0.36mM	0.36mM
Primers	0.5μΜ	0.4µM	0.4μΜ	0.4μΜ
Lambda DNA	0.05μg	0.03μg	0.03μg	0.25μg
MaxTaq DNA Polymerase	2.0u	2.0u	2.0u	2.0u
DMSO / Formamide	3%	3%	3%	3%

Top up with sterile dH $_{\rm 2}\text{O}$ to 25 $\mu\text{I}.$

Product Size	8kb	10 and 12kb	15 and 20kb	30 and 40kb
Denaturation	94°C, 2min	94°C, 2min	94°C, 2min	94°C, 2min
Denaturation	94°C, 12s	94°C, 12s	94°C, 12s	94°C, 12s
Annealing	65°C, 30s		56°C, 30s	58.5°C, 30s
Extension / 1kb	68°C,5min	68°C, 12min	68°C, 15min	68°C, 25min
Cycles	30	35	10	10
Final Extension / Extension	68°C, 10min	68°C, 7min	-	-
Denaturation	-	-	94°C, 12s	94°C, 12s
Annealing	-	-	56°C, 30s	58.5°C, 30s
Extension / 1kb ¹	-	-	68°C, 15min	68°C, 30min
Cycles	-	-	20	20
Final Extension	-	-	68°C, 7min	68°C, 7min

¹Plus additional 20s auto-extension after each cycle.

AtMax *Taq* DNA Polymerase (PL4201 - PL4202)

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Product Size	5kb	5 and 8kb	10, 15 and 20kb
ViBuffer (1X)	Α	S	S
MgCI ₂	1.5mM	-	-
dNTP mix	0.25mM	0.25mM	0.36mM
Primers	0.4μM	0.4μM	0.6μΜ
Lambda DNA	0.15µg	0.15µg	0.15μg
AtMax <i>Taq</i> DNA Polymerase	2.0u	2.0u	2.0u
DMSO / Formamide	3%	3%	3%

Top up with sterile dH,0 to 25µl.

Product Size	5kb	8kb	10kb	15 and 20kb
Denaturation	94°C, 2min	94°C, 2min	94°C, 2min	94°C, 2min
Denaturation	94°C, 12s	94°C, 12s	94°C, 12s	94°C, 12s
Annealing	60°C, 30s	65°C, 30s	59°C, 30s	56°C, 30s
Extension / 1kb	68°C, 4min	68°C, 5min	68°C,12min	68°C, 15min
Cycles	30	30	35	10
Final Extension / Extension	68°C, 10min	68°C, 10min	68°C, 7min	-
Denaturation	-	-	-	94°C, 12s
Annealing	-	-	-	56°C, 30s
Extension / 1kb ¹	-	=	-	68°C, 15min
Cycles	-	-	-	20
Final Extension	-	-	-	68°C, 7min

Pfu DNA Polymerase / Chromo Pfu DNA Polymerase (PL5201 - PL5206)

Product Size	0.5kb	1.5kb	5kb	8kb
ViBuffer (1X)	Α	Α	S	S
MgCl ₂	2.5mM	1.5mM	-	-
dNTP mix	0.2mM	0.2mM	0.25mM	0.25mM
Primers	0.4μM	0.2μM	0.2μM	0.8μΜ
Plasmid DNA	0.03μg	0.03μg	0.03μg	0.03μg
Pfu DNA Polymerase	2.0u	2.0u	2.0u	2.0u
DMSO / Formamide	-	-	3%	3%

Top up with sterile dH₂O to 25μl.

Product Size	0.5kb	1.5kb	5kb	8kb
Denaturation	94°C, 2min	94°C, 1min	94° C, 2min	94°C, 2min
Denaturation	94°C, 30s	94°C, 30s	94°C, 12s	94°C, 12s
Annealing	59°C, 30s	52°C, 30s	60°C, 30s	65°C, 30s
Extension / 1kb	72°C, 90s	72°C, 3min 30s	68°C, 10min	68°C, 16min
Cycles	25	25	30	30
Final Extension / Extension	72°C, 5min	72°C, 5min	68°C, 10min	68°C, 10min
Denaturation	-	-	-	-
Annealing	-	-	-	-
Extension / 1kb ¹	-	-	-	-
Cycles	-	-	-	-
Final Extension	=	-	-	-

RECOMMENDED PROTOCOL FOR PLMM01:

Gently mix all solutions after thawing. Spin down briefly and keep on ice. Add the following components in a 0.2ml thin walled PCR tube on ice:

For 50µl reaction volume:

Reagent:	Volume	Final Concentration	
2X Taq Master Mix	25 μ Ι	*1X	
MgCl ₂ (50mM)	Refer for the TABLE (A)	**For more than 1.5mM MgCl ₂	
Primers (Fwd' / Rev')	Variable	0.1 - 1μM each	
DNA Template	Variable	0.02 - 5 μg	
de-Ionized distilled H ₂ O	Adjust final volume to 50μl		

^{* 1.25} unit Taq DNA polymerase, 1X ViBuffer A, 0.2mM dNTPs and 1.5mM MgCl₂.

^{**2}X Taq Master Mix contains a fixed final MgCl₂ concentration of 1.5mM. However, higher concentration may be achieved by adding additional MgCl₂. Please refer to Table(A) if higher MgCl₂ is preferred.

CYCLING COND		
Denaturation	94°C for 2 minutes	
Denaturation	94°C for 20 seconds	ר
Annealing	50 - 68°C for 30 seconds	25 - 35 cycles
Extension / 1kb	72°C for 30 seconds	J
Final Extension	72°C for 7 minutes	

The protocol may change depending on the template DNA and primers used.

TABLE (A): For more than 1.5mM final MgCl₂ concentration

Volume of MgCl ₂ (50mM) stock to add into reaction mixture (µI)	Final MgCl ₂ concentration (mM)
0.5	2.0
1.0	2.5
1.5	3.0
2.0	3.5
2.5	4.0

Note: Smaller reaction volume may be achieved provided that the same final concentration of each reaction component is maintained.

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Optimization of MgCl₂ concentrations

- 1.Gently vortex and briefly centrifuge all solutions after thawing.
- 2.Dilute your forward and reverse primer to 10μM working stock solutions.

Amplification protocol for 2X Ampli-Optimization Kit (PLAO01)

- 3.Label a set of six 0.2ml microcentrifuge tubes as 1-6.
- 4.Use a sterile microcentrifuge tubes to prepare a **Template-Primer Master Mix** for 6 reactions as follows:

Reagent	Volume (For 7 reactions)	Final Concentration (in each of 50µl reaction)
Forward Primer (10µM) Reverse Primer (10µM) Template DNA	Variable (3.5-35μl) Variable (3.5-35μl) variable	0.1-1μM (recommended 0.2μM) 0.1-1μM (recommended 0.2μM) 0.01-10ng for plasmid 0.05-1μg for genomic DNA
Water, nuclease-free	to 175μΙ	

- 5.Mix thoroughly, centrifuge briefly and aliquot 25µl of **Template-Primer Master Mix** to tube 1-6.
- 6.Add 25μl of **2X** Taq Master Mix (A-F) to tubes 1-6 respectively.
- 7. Mix thoroughly and centrifuge briefly. Place the reactions tubes in a thermocycler and start the amplification program.

Recommended program for most common amplifications:

Step	Temperature	Time	Number of Cycles
Initial Denaturation	94℃	2 min	1
Denaturation	94℃	30 sec	
*Annealing	50 – 68℃	30 sec	25-35
Extension	72℃	30 sec/kb	
Final Extension	72℃	5 min	1

^{*} Recommended annealing temperature is 1-5°C below the lowest melting temperature of the primers in the reaction mixture.

General Technotes

Separation of double-stranded linear DNA fragments on agarose gel¹

% Agarose	Size Range (bp)
0.5	2000-30000
0.7	700-20000
1.0	500-10000
1.2	400-7000
1.5	200-3000
2.0	100-2000

¹Leonard D, Michael K, James B. *Basic Methods in Molecular Biology*. 2nd. Appleton & Lange, CT: Paramount Publishing Business and Professional Group; 1994. Chapter 5.

Separation of double-stranded linear DNA fragments relative to marker dyes on native polyacrylamide gel¹

% Polyacrylamide	Size Range (bp)	Xylene Cyanol FF*	Bromophenol Blue*
3.5	500-2000	460 bp	100 bp
5.0	100-500	260 bp	65 bp
8.0	60-400	160 bp	45 bp
12.0	40-200	70 bp	20 bp
15.0	25-150	60 bp	15 bp
20.0	6-100	45 bp	12 bp

^{*} Dyes will co-migrate with DNA fragments of approximately this size (bp)

¹Sambrook J, Fristch EF, Maniatis T, Molecular Cloning: A Laboratory Manual. 2nd ed. Cold Spring Harbor, NY: Cold Spring Harbor Press; 1989. Chapter 6.

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General Conversions

Molar Conversions

1 μg of 1000bp DNA	= 1.52 pmol	= 9.1 X 10 molecules
1 μg of pUC18/19 DNA (2686 bp)	= 0.57 pmol	= 3.4 X 10 molecules
1 μg of pBR322 DNA (4361 bp)	= 0.35 pmol	= 2.1 X 10 molecules
1 μg of M13mp18 /19 DNA (7249 bp)	= 0.21 pmol	= 1.3 X 10 molecules
1 μg of λ DNA (48502 bp)	= 0.03 pmol	= 1.8 X 10 molecules
1 pmol of 1000 bp DNA	= 0.66 μg	
1 pmol of pUC18/19 DNA (2686 bp)	= 1.77 μg	
1 pmol of pBR322 DNA (4361 bp)	= 2.88 μg	
1 pmol of M13mp18/19 DNA (7249 bp)	= 4.78 μg	
1 pmol of λ DNA (48502 bp)	= 32.01 μg	

Spectrophotometric Conversions

```
\begin{array}{lll} 1 \ A_{260} \ \text{of dsDNA} &= 50 \ \mu\text{g/ml} \\ 1 \ A_{260} \ \text{of ssDNA} &= 33 \ \mu\text{g/ml} \\ 1 \ A_{260} \ \text{of ssRNA} &= 40 \ \mu\text{g/ml} \\ 1 \ \text{mM} \ \text{(in nucleotides)} \ \text{of dsDNA} \\ 1 \ \text{mM} \ \text{(in nucleotides)} \ \text{of ssDNA} \\ 1 \ \text{mM} \ \text{(in nucleotides)} \ \text{of ssDNA} \\ 1 \ \text{mM} \ \text{(in nucleotides)} \ \text{of ssRNA} \\ \end{array} \begin{array}{ll} = 0.15 \ \text{mM} \ \text{(in nucleotides)} \\ = 0.12 \ \text{(in nucleotide
```

High purity DNA has an A₂₆₀ / A₂₈₀ ratio: 1.8 - 2.0 High purity RNA has an A₂₆₀ / A₂₈₀ ratio: 1.9 - 2.1

Conversions of Oligonucleotides

Molecular Weight MW = 333 X N

Concentration

 $C(\mu M \text{ or pmol/}\mu I)$ = $A_{260} / (0.01 \text{ X N})$ C(ng/mI) = $(A_{260} \text{ X MW}) / (0.01 \text{ X N})$

MW = molecular weight
A₂₆₀ = absorbance at 260nm
N = number of bases

DNA/Protein Conversions

1kb of DNA = 333 amino acids ~ 3.7 X 10⁴ Da 10,000 Da protein = 270 bp DNA 30,000 Da protein = 810 bp DNA 50,000 Da protein = 1320 bp DNA 100,000 Da protein = 2700 bp DNA

General PCR Protocol

- 1. Gently vortex and briefly centrifuge all solutions after thawing.
- 2. Dilute forward and reverse primer to 10µM working stock solutions.
- 3. Add the following components to a 0.2ml PCR tube:

Reagent	Final Concentration	Volume per 25µl reaction
Water, nuclease-free	-	Top up to 25µl
10X ViBuffer A	1X	2.5µl
50mM MgCl ₂	1 – 4mM*	Variable
10mM dNTP mix	0.1mM	0.25µl
Forward Primer (10µM)	0.1-1µM**	Variable
Reverse Primer (10µM)	0.1-1µM**	Variable
Polymerase (5u/µl)	0.08u/ul	0.4µl
Template DNA	variable	0.01-10ng for plasmid
		0.05-1µg for genomic DNA
		10 ³ to 10 ⁶ copies

- * Specific MgCl₂ concentration depends on the primers and template combination. For optimization of MgCl₂ concentration please refers to the table below.
- ** Recommended to use 0.2μM final concentration. For degenerate primers the concentration may increase to 0.5 1μM in order to increase the amount of "correct" primer; therefore, increase the yield of the expected product.

Final Concentration	1.0mM	1.5mM	2.0mM	2.5mM	3.0mM	3.5m M	4.0mM
Volume of 50mM MgCl ₂	0.5µl	0.75µl	1µl	1.25µl	1.5µl	1.75µl	2µl

- 1. Mix thoroughly, centrifuge briefly and aliquot the master mix to all PCR tubes.
- 2. Gently spin down to collect the PCR reaction mix in the bottom of the tubes.
- 3. Place the reactions tubes in a thermocycler and start the amplification program.

Recommended program for most common amplifications:

St	tep	Temperature	Time	Number of Cycles
1	Initial Denaturation	94°C	2 min	1
2	Denaturation	94°C	30 sec	
	*Annealing	40 – 68°C	30 sec	25 -35
	Extension	72°C	30 sec/kb	
3	Final Extension	72°C	5 min	1

^{*} Recommended annealing temperature is 1-5°C below the lowest melting temperature of the primers in the reaction mixture.



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General Guideline for PCR Optimization

Primers Guideline on designing primers

- Usually the lengths of primers are 18 28 nucleotides long.
- GC content should be 45 60%. The G and C nucleotides should have a balance distribution within the full length of the primer. Avoid more than three G or C nucleotides at the 3' end to lower the risk of non-specific priming.
- The primer is not self-complement or complementary to other primer in the reaction mixture to avoid internal secondary structure or the formation of primer dimer.
- Melting temperature of the primer should be 50 70°C. Melting temperature of flanking primer should not differ more than 5°C.
- If using degenerate primer, avoid degeneracy for the last 3 nucleotides at the 3'end.
- The designed primers should be checked for possible complementary sites within the template DNA. Primer with more than one complementary site within the template DNA should be avoided, as non-specific priming will occur.
- Standard concentration of primer in the reaction mixture is 0.2µM. Increase the primer concentration may increases the possibility of non-specific priming. However for degenerate primer, the concentration may increase to 0.5-1µM in order to increase the amount of "correct" primer; therefore, increase the yield of the expected product.

Template DNA

The template DNA is likely the largest variable in PCR. This is because the same amount of DNA from different organism represents different copies number. In general, optimal amount for plasmid, cDNA and phage DNA is in the range of 0.01 – 10ng and for genomic DNA is in the range of 0.05 – 1µg per 25µl reaction volume.

MgCl₂ Concentration

Specific MgCl₂ concentration is essential for each amplification reaction. MgCl₂ form complex with dNTPs, primers and template DNA. In addition, Taq require free MgCl₂ as cofactor, therefore MgCl₂ affecting strand dissociation, primer annealing, enzyme activity and fidelity. Thus the optimal concentration of MgCl₂ is very crucial for each amplification reaction. The recommended concentration range is 1 – 4mM MgCl₂. Too low MgCl₂ concentration result in low yield of desire product, while too high MgCl₂ concentration increases the yield of non-specific product and decreases the fidelity of reaction. Lower MgCl₂ concentration is preferred when fidelity of DNA synthesis is critical.

Tag DNA Polymerase

Most amplification reactions use 1 to 2.5 unit of *Taq* DNA Polymerase for 25µl total reaction volume. Generally, high concentration of *Taq* DNA Polymerase produces more products as the efficiency is better. However, non-specific products may appear at higher concentration of *Taq* DNA Polymerase. It is recommended to start with 2 unit of *Taq* DNA Polymerase for 25µl total reaction volume. The unit of *Taq* DNA Polymerase could be decrease after the specific MgCl₂ concentration has been determined in order to avoid non-specific amplification. In addition, *Taq* DNA Polymerase from different manufacturer may have different amplification efficiency and unit definition. Therefore, user may needs to adjust the amount experimentally when using Taq DNA Polymerase from different manufacturer.

Reaction Buffer

10X ViBuffer A is designed for amplification of short PCR products (up to 5kb) and 10X ViBuffer S is designed for amplification of long PCR products (> 5kb).

Initial Denaturation

A complete denaturation of DNA template at the beginning of the amplification reaction is essential as incomplete denaturation of DNA will result in low efficiency in the first amplification cycle, leading to poor yield of the amplified product. For most applications, the initial denaturation of 2 to 5 minutes at 94°C is usually sufficient. For GC rich template, the addition of DMSO between 3 – 10% (5% is recommended as a starting point) may improve the amplification efficiency. With the addition of DMSO, the amount of *Taq* should increase to 2 units per reaction.

Denaturation

Since the amplified product in the first cycle is significantly shorter than the template DNA, denaturation at 94° C for 30 seconds is usually sufficient. For amplicon with high GC content the denaturation time may be increased to 2-4 minutes

Annealing

Annealing temperature is a vital parameter of an amplification reaction. The annealing temperature is usually chosen based on the length and GC content of the oligonucleotide primers. Annealing temperature is often $1-5^{\circ}$ C below the primer melting temperature (Tm). It is the best to choose the sequence of both primers such that the melting temperature (Tm) of both primers does not differ more than 5° C. If the Tm of the flanking primers is different, use $1-5^{\circ}$ C of annealing temperature below the lowest Tm of primers in the reaction mixture. Annealing time for 30 seconds to 2 minutes is usually sufficient. In general, higher annealing temperature will increase primer-template specificity and result in less non-specific amplification.

Elongation

Usually the extending step is performed at 72°C, which is the optimal temperature for primer extension by *Taq* DNA Polymerase. Recommended elongation time is 30 seconds for each kilobase-pairs of product to be amplified.

Number of Cycles

The choice for the number of amplification cycles depends on the amount of template DNA in the reaction mix and on the expected yield of the amplification. If quantity of DNA template is higher than 100 copies, 25 to 35 cycles are usually sufficient. If fidelity of synthesis is crucial, maximum allowable amount of template with minimum amplification cycles is recommended.

Final Elongation

Upon the end of the last cycle, samples are usually incubated at 72°C for 2 to 5 minutes in order to let the *Taq* DNA polymerase to finish all primer extension process and for the terminal transferase activity of the *Taq* DNA polymerase to add extra adenine nucleotides to the 3'-ends of the amplified products. If amplified fragments are to be used for TA cloning purposes, this step can be extended up to 30 minutes.

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Troubleshooting

Problem	Possibility	Recommended Solution
Low yield or no PCR product	Missing component in reaction mixture	Check the reaction components and repeat the reaction.
	Not enough template	Use more template or increase cycle number.
	Not enough cycles	Increase the number of cycles by 3 to 5 increment.
	Annealing temperature too high	Decrease annealing temperature by 2°C at a time.
	Annealing time too short	Increase annealing time to 1-2 minutes.
	GC-rich template	Increase denaturing temperature to 97-99°C and denaturing time to 5-10 minutes. Add GC destabilizing co-solvent such as DMSO (3-10%) to the reaction mixture.
	Elongation time too short	Increase the elongation time by 1 minute at a time.
Smearing	Agarose gel not fresh	Repeat electrophoresis with fresh agarose gel.
	Too many cycles	Reduce the number of cycles by 3 to 5 cycles at a time.
	Too much template	Reduce amount of template by 10 to 1000 times dilution.
	Degraded template	Check and confirm template integrity by agarose gel electrophoresis. If necessary, repurify template using methods that minimizes shearing and nicking.
	Denaturation temperature too high	Decrease denaturing temperature to 94°C.
	Elongation time too short	Increase elongation time by 1 to 2 minutes increments.
Non-specific band	Too many cycles	Reduce the number of cycles by 3 to 5 cycles at a time.
	Too much template	Reduce amount of template by 10-1000 times dilution.
	Annealing temperature too low	Increase annealing temperature by 2 to 3°C at a time.
	Elongation time too long	Reduce elongation time.
	Cross contamination	Use a separate workplace, pipettes and filter tips. Wear glove at all times.

Recommended protocol for first strand cDNA synthesis

1. Prepare the following RNA-primer mixture in a tube on ice:

Component	Amount	Volume
Template: total RNA	0.1 – 10μg (Recommended 5μg)	Variable
or poly A(+) mRNA	0.01 – 1µg (Recommended 0.5µg)	Variable
		Variable
Primer: Oligo(dT) ₁₈	100pmol	Variable
or random Hexamers*	100pmol	Variable
or gene-specific primer	20pmol	Variable
Nuclease-free water		Top up to 13µl

- * The use of random hexamer is not recommended when total RNA is used as template, as rRNA and tRNA may also be primed and copied, resulting in a lower efficiency of cDNA transcribed from mRNA.
- 2. Incubate the mixture at 65°C for 5 minutes and chill on ice for 2 minutes.
- 3. Briefly spin down the mixture.
- 4. Add the following in the order indicated:

Component	Amount	Volume
10X Reaction Buffer	1X	2μΙ
10mM dNTP mix	1mM	2μΙ
Ribonuclease Inhibitor (40u/μl)	20 units	0.5μΙ
Reverse Transcriptase*	Variable	Variable
Nuclease-free water		Top up to 7µl

* Different reverse transcriptase required different unit of enzyme for reverse transcription. Please refer to the table below for the amount of reverse transcriptase.

Reverse Transcriptase	Amount/ reaction
M-MuLV Reverse Transcriptase	200 units
AMV Reverse Transcriptase	10 units

- 5. Mix gently and incubate at 42°C for 60 minutes.
- 6. Stop the reaction by placing the tube on ice.
- 7. The synthesized cDNA can be directly used in PCR, by addition of 1 2µl of the cDNA reaction mixture to a 25µl PCR reaction.

Note: If the cDNA is to be used for cloning, it should first be dephosphorylated as the primers have 5'-PO₄ ends.

Technical Information

Technical Information

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Agarose Gel Preparation

Preparation of 40ml 1% TAE agarose gel

- 1. Weight 0.4g of agarose into a conical flask (the size of the flask should be three times more than the volume of the agarose gel you want to prepare to avoid spill out and over evaporation).
- 2. Add 1X TAE buffer into the conical flask until the weighing machine display 40g.
- 3. With the conical flask still on top of the balance, zero the balance.
- 4. Boil the agarose solution in microwave for about 2 minutes.

Caution: **DO NOT** cover the conical flask with any tissue or stopper. This will cause pressure build up and **explosion**.

5. Make sure there is no undissolved agarose.

Note: For high percentage gel (> 2%) you need to swirl the solution with spatula and reboil to dissolve all the agarose.

- 6. Check the volume of the solution by weighing the conical flask with the solution. Add in distilled water until the display is 0. Boil for a few seconds to mix the solution.
- 7. Cool the agarose to about 55°C by using running tap water around the outside of the conical flask.
- 8. Pour the agarose into the casting tray.
- 9. Insert comb and let it polymerize properly for about 30 minutes at room temperature.



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Vivantis Technologies Sdn Bhd 587389-D

Revongen Corporation Center No12A, Jalan TP5, Taman Perindustrian UEP, 47600 Subang Jaya, Selangor Darul Ehsan, Malaysia.

www.vivantechnologies.com

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