



A typical freezer in a molecular biology laboratory consists of a host of the quintessential restriction enzyme 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 for "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!

10-go, giving scientists time to tango!

Lot #
Store AT -20°C

10-GO RE KIT SET (D) - RK4000

Kit Component : Restriction Endonucleases (10 types)
0.5ml Diluent Viva Buffer A
1ml 10X Buffers (Buffer V1, V2, V3, V4, V5, Universal Buffer, Buffer EcoR I)
(BSA included in all Reaction Buffer)

RESTRICTION ENDONUCLEASES SPECIFICATION:

ENZYME	(u/μl)	TOTAL UNIT	REG.SITE	OPT. BUFFER	OPT.(°C)	THERMA INAC.	ACTIVITY IN REACTION BUFFER				
							V1	V2	V3	V4	V5
BamH I	15	400	5'..G^GATCC..3'	V2	37	65°C, 20min.	75%	100%	75%	75%	75%
Bme18 I (Ava II)	5	50	5'..G^GWCC..3'	V3	37	65°C, 20min.	50%	75%	100%	100%	100%
EcoR I	20	500	5'..G^AATTC..3'	SP	37	65°C, 20min.	50%	50%	100%	100%	50%
FauND I (Nde I)	10	50	5'..CA^TATG..3'	V5	37	65°C, 20min.	50%	75%	50%	75%	100%
Hind III	20	400	5'..A^AGCTT..3'	V2	37	65°C, 20min.	75%	100%	75%	75%	75%
Kpn I	20	200	5'..GGTAC^C..3'	V1	37	Non-Thermal Inac.	100%	25%	25%	25%	75%
Psp124B I (Sac I)	20	100	5'..GAGCT^C..3'	V2	37	65°C, 20min.	100%	100%	75%	100%	100%
Sfr274 I (Xho I)	10	200	5'..C^TCGAG..3'	V1	50	65°C, 20min.	100%	100%	50%	50%	75%
Sma I	20	100	5'..CCC^GGG..3'	V5	25	65°C, 20min.	25%	10%	10%	75%	100%
Taq I	20	300	5'..T^CGA..3'	V5	65	Non-Thermal Inac.	10%	100%	75%	75%	100%

REMARK:

BamH I High enzyme concentration may result in Star Activity.
EcoR I High enzyme concentration may result in Star Activity.
FauND I Sensitive to impurities present in some DNA preparation.
Sma I Blocked by CpG Methylation.
Taq I Blocked by overlapping dam-methylation.

BUFFER COMPOSITION:

	10mM Tris-HCL (pH 7.5 at 30°C), 10mM MgCl ₂ , and 100μg/ml BSA.
	10mM Tris-HCL (pH 7.5 at 30°C), 10mM MgCl ₂ , 50mM NaCl and 100μg/ml BSA.
	50mM Tris-HCL (pH 7.5 at 30°C), 10mM MgCl ₂ , 100mM NaCl and 100μg/ml BSA.
	10mM Tris-HCL (pH 8.5 at 30°C), 10mM MgCl ₂ , 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.
	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.
	Buffer EcoR I 50mM Tris-HCL (pH 7.5 at 30°C), 10mM MgCl ₂ , 100mM NaCl, 0.02% triton X-100 and 0.1mg/ml BSA.

BUFFER COMPATIBILITY CHART

RK6000	BamH I	Bme18 I	EcoR I	FauND I	Hind III	Kpn I	Psp124B I	Sfr274 I	Sma I	Taq I
BamH I		V4/V5	V4	V1/V4/V5	V1/V4/V5	V1	V1/V4/V5	V1	V5	V5
Bme18 I	V4/V5		V3/V4	V4/V5	V2/V5	V5	V4/V5	1.0X UB	V5	V5
EcoR I	V4	V3/V4		V4	V3/V4	V1	V4	all	V4	V3/V4
FauND I	V1/V4/V5	V4/V5	V4		V1/V4/V5	V1	V1/V4/V5	V1	V5	V5
Hind III	V1/V4/V5	V2/V5	V3/V4	V1/V4/V5		V1	V2	V2	V5	V2
Kpn I	V1	V5	V1	V1	V1		V1	V1	V5	V5
Psp124B I	V1/V4/V5	V4/V5	V4	V1/V4/V5	V2	V1		V/V2	V5	V2/V5
Sfr274 I	V1	1.0X UB	all	V1	V2	V1	V1/V2		1.0X UB	V2
Sma I	V5	V5	V4	V5	V5	V5	V5	1.0X UB		V5
Taq I	V5	V5	V3/V4	V5	V2	V5	V2/V5	V2	V5	

Activity:

	100%		Specific buffer
	75%		Not applicable
	50%		

ENZYME	LIGATION / RECURTING ASSAY	OVERDIGESTION ASSAY
BamH I	After 15-fold overdigestion with BamH I , more than 90% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 30 uof BamH I for 16 hours at 37°C (Without BSA).
Bme18 I (Ava II)	After 5-fold overdigestion with Bme18 I , more than 90% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 10u of Bme18 I for 16 hours at 37°C (Without BSA).
EcoR I	After 20-fold overdigestion with EcoR I , more than 95% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of EcoR I for 16 hours at 37°C (Without BSA).
FauND I (Nde I)	After 10-fold overdigestion with FauND I , 60% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of FauND I for 16 hours at 37°C.
Hind III	After 20-fold overdigestion with Hind III , more than 90% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of Hind III for 16 hours at 37°C.
Kpn I	After 20-fold overdigestion with Kpn I , more than 90% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of Kpn I for 16 hours at 37°C (Without BSA).
Psp124B I (Sac I)	After 20-fold overdigestion with Psp124B I , more than 95% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of Psp124B I for 16 hours at 37°C (Without BSA).
Sfr274 I (Xho I)	After 10-fold overdigestion with Sfr274 I , 90% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 20u of Sfr274 I for 16 hours at 50°C.
Sma I	After 20-fold overdigestion with Sma I , more than 90% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of Sma I for 16 hours at 25°C.
Taq I	After 20-fold overdigestion with Taq I , more than 95% of the DNA fragments can be ligated and recut.	An unaltered banding pattern was observed after 1µg of DNA was digested with 40u of Taq I for 16 hours at 65°C.