

Enzyme Information

Enzyme	Recognition Sequence	Sites cut	Sites not cut	Enzyme	Recognition Sequence	Sites cut	Sites not cut
EcoR I	GAATTC	GAATTh ^{m5} C	G ^{m5} AATTC ^b GA ^{m5} ATTC# GAATT ^{m5} C ^b	Hph I	TCACC	TCAC ^{m5} C	T ^{m5} CACC# GGTG ^{m5} A
EcoR II	CCWGG	^{m5} CCWGGb	^{m4} CCWGG C ^{m4} CWGG C ^{m5} CWGG# CC ^{m5} AGG h ^{m5} Ch ^{m5} CWGG	Kpn I	GGTACC ^b	GGTA ^{m5} CC GGTAC ^{m5} C GGTA ^{m5} C ^{m5} C ^b GGT ^{m5} ACC	GGT ^{m5} A ^{m5} CC GGTAC ^{m4} C
EcoR V	GATATC	GATAT ^{m5} C ^b	G ^{m5} ATATC# GAT ^{m5} ATC	Kpn2 I	TCCGGA	TCCGG ^{m5} A	T ^{m5} CCGGA TC ^{m5} CGGA
EcoR 124	GAAN6RTCGB	?	GA ^{m5} AN6RTCGB GAAN6RmTCG	Ksp I	CCGCGG	?	^{m5} CCGCGG C ^{m5} CGCGG
EcoR 124/3	GAAN7RTCGB	?	^{m5} A	Mae II	ACGT	?	A ^{m5} CGTb
Ehe I	GGCGCC	?	GG ^{m5} CGCC	Mam I	GATN4ATC	?	G ^{m5} ATN4 ^{m5} ATC
Esp I	GCTNAGC	GCTNAG ^{m5} C	G ^{m5} CTNAGC	Mbo I	GATC ^b	GAT ^{m4} C GAT ^{m5} C ^b	G ^{m5} ATC# GATH ^{m5} C
Fnu4H I	GCNGC	?	G ^{m5} CNGC GCNG ^{m5} C	Mbo II	GAAGA	T ^{m5} CTT ^{m5} C ^b G ^{m5} AAGA	GAAG ^{m5} A#
FnuD II	CGCG	?	^{m5} CGCG CG ^{m5} CG	Mfi I	REGATCY ^b	?	RG ^{m5} ATCY RGAT ^{m4} CY RGAT ^{m5} CY
Fok I	CATCC	CAT ^{m5} CC CATC ^{m5} C ^b	GG ^{m5} ATG C ^{m5} ATCC CATC ^{m4} C	Mlu I	ACGCGT	^{m5} ACGCGT	A ^{m5} CGCGT
Fse I	GGCCGGCC	?	GG ^{m5} CCGG ^{m5} CC GGC ^{m5} CGGCC GG ^{m5} CCGGCC	Mme II	GATC	?	G ^{m5} ATC
Fsp I	TGCGCA	?	TG ^{m5} CGCA	Mnl I	CCTC ^b	?	^{m5} CCTC ^{m5} C ^{m5} CT ^{m5} C
Hae II	RGCGCY ^b	?	RG ^{m5} CGCY RGh ^{m5} CGh ^{m5} CY	Mro I	TCCGGA	TCCGG ^{m5} A	T ^{m5} CCGGA TC ^{m5} CGGA
Hae III	GGCC	GGC ^{m5} C	GG ^{m5} CC# ^b GGh ^{m5} Ch ^{m5} C	Mse I	TTAA	Tm6AA	?
Hap II	CCGG	?	C ^{m5} CGG#	Msp I	CCGG ^b	^{m4} CCGG C ^{m4} CGG C ^{m5} CGG	^{m5} CCGG# h ^{m5} Ch ^{m5} CGG
Hga I	GACGC	?	GA ^{m5} CGC GACG ^{m5} C	Mst II	CCTNAGG	^{m5} CCTNAGG	?
HgiA I	GRGICY	GRGICY ^{m5} C	GRG ^{m5} CYC	Mun I	CAATTG	?	CA ^{m5} ATTG
HgiC I	GGYRCC	?	GGYRC ^{m5} C	Mva I	CCWGG	C ^{m5} CWGG ^b ^{m5} CCWGG CC ^{m5} AGG ^b	C ^{m4} CWGG# ^{m4} CCWGG ^{m5} C ^{m5} CWGG ^b
HgiC II	GGWCC	?	GGWC ^{m5} C	Mvn I	CGCG	?	m5CGCG
HgiE I	GGWCC	?	GGWC ^{m5} C	Nae I	GCCGGC ^b	?	G ^{m5} CCGGC GC ^{m5} CGGC GCCGG ^{m5} C
HgiJ II	GGYRCC	?	GGYRC ^{m5} C	Nar I	GGCGCC	GGCGC ^{m5} C	GG ^{m5} CGCC GGCGC ^{m4} C
Hha I	CGCG	?	G ^{m5} CGC# GCG ^{m5} C Gh ^{m5} CGh ^{m5} C	Nci I	CCSGG	^{m5} CCSGG	C ^{m4} CSGG C ^{m5} CSGG ^b
Hha II	GANTC	?	G ^{m5} ANTC#	Nco I	CCATGG	CC ^{m5} ATGG	^{m4} CCATGG ^b ^{m5} CCATGG
Hinc II	GTYRAC	GTYRA ^{m5} C	GTYR ^{m5} AC GTYRAh ^{m5} C	Nde I	CATATG	^{m5} CATATG ^b	^{m5} A
Hind II	GTYRAC	?	GTYR ^{m5} AC#	Nde II	GATC	GAT ^{m5} C ^b	G ^{m5} ATC
Hind III	AAGCTT	A ^{m5} AGCTT#	^{m5} AAGCTT# AAG ^{m5} CTT AAGh ^{m5} CTT	Nhe I	GCTAGC	?	GCTAG ^{m5} C
Hinf I	GANTC	GANT ^{m5} C ^b	G ^{m5} ANTC GANTh ^{m5} C	Nla III	CATG	?	C ^{m5} ATG#
HinP I	CGCG	?	G ^{m5} CGC	Not I	GCGGCCGC	GCGGCCG ^{m5} C	GCGG ^{m5} CCGC GCGGC ^{m5} CGC
Hpa I	GTTAAC	GTTA ^{m5} C	GTTA ^{m5} AC# GTTAAh ^{m5} C	Nru I	TCGCGA	TCG ^{m5} CGA	T ^{m5} CGCGA TCGCG ^{m5} A
Hpa II	CCGG	?	^{m4} CCGG ^{m5} CCGG ^b C ^{m4} CGG ^b C ^{m5} CGG# h ^{m5} Ch ^{m5} CGG	Nsi I	ATGCAT	?	ATGC ^{m5} AT ATG ^{m5} CAT
				Nsp I	RCATGY	?	RC ^{m5} ATGY
				NSpB II	CMGCKG	C ^{m5} CGCKG	?
				PfiM I	CCAN5TGG	?	C ^{m4} CAN5TGG C ^{m5} CAN5TGG

LEGEND

a. # denotes canonical modification mTase specificity.
b. See notes section of reference 2.

*Dpn I and its isoschizomers require the presence of 6-methyladenosine within the recognition sequence GATC

Recognition sequences given use the standard abbreviations [*Eur. J. Biochem* (1985) [50:15] to represent ambiguity:

R=G or A
M=A or C
S=G or C
H=A or C or T
V=G or C or A
N=A or C or G or T

Y=C or T
K=G or T
W=A or T
B=G or T or C
D=G or A or T

^{m4}C = N4-methylcytosine
^{m5}C = C5-methylcytosine
^{hm5}C = hydroxymethylcytosine
^mC = methylcytosine
N₄ or C₅-methylcytosine unspecified
^{m6}A = N₆-methyladenine

Sequences are in 5' - 3' order.

Enzyme Information, continued

Enzyme	Recognition Sequence	Sites cut	Sites not cut	Enzyme	Recognition Sequence	Sites cut	Sites not cut
Pfu I	CGTACG	?	CGTA ^{m5} CG	SnaB I	TACGTA	?	TA ^{m5} CGTA T ^{m5} ACGT ^{m5} A
PaeR7 I	CTCGAG	?	CTCG ^{m5} AG# CT ^{m5} CGAG	Sno I	GTGCAC	?	GTG ^{m5} CA ^{m5} C
Pml I	CACGTG	?	CA ^{m5} CGTG	Spe I	ACTAGT	?	^{m5} ACTAGT A ^{m5} CTAGT
PpuAI	CGTACG	?	CGTA ^{m5} CG	Sph I	GCATGC	GCATG ^{m5} C Gh ^{m5} CATGh ^{m5} C	GC ^{m5} ATGC
PspA I	CCCGGG	C ^{m5} CCGGG CC ^{m5} CGGG	^{m5} CCCGGG	Spl I	CGTAGC	CGTm6ACG	?
Pst I	CTGCAG	?	^{m5} CTGCAG CTGC ^{m5} AG#	Spo I	TCGCGA	TCGCG ^{m5} A	T ^{m5} CGCGA TCG ^{m5} CGA
Pvu I	CGATCGb	CG ^{m5} ATCG CGAT ^{m5} CG	CGAT ^{m4} CG	Srf I	GCCC/GGGC	GCCC/GGG ^{m5} C	G ^{m5} CCC/GGGC GC ^{m5} CC/GGGC GCC ^{m5} C/GGGC
Pvu II	CAGCTG	?	CAG ^{m4} CTG# CAG ^{m5} CTG	Ssp I	AATATT	^{m5} AATATT	?
Rsa I	GTACb	GTA ^{m5} C ^b	GT ^{m5} AC	Sst I	GAGCTC	?	GAG ^{m5} CTC GAGh ^{m5} CTh ^{m5} C
Rsr I	GAATTC	?	G ^{m5} AATTC GA ^{m5} ATTC# ^b	Stu I	AGGCCT	?	AGGm5CCT AGGCm5CT AGGCm4CT
Rsr II	CGGWCCG	?	^{m5} CGGWCCG CGGW ^{m5} CCG CGGW ^{m5} CCG	StySP I	AACN6GTRC ^b	?	A ^{m5} ACN6GmTRC ^b
Sac I	GAGCTC	G ^{m5} AGCTC	GAG ^{m5} CTC	Taq I	TCGA	T ^{m5} CGA ^b Th ^{m5} CGA ^b	TCG ^{m5} A#
Sac II	CCGCGG	?	^{m5} CCGCGG	Taq II	GACCGA CACCCA	?	G ^{m5} ACCGA
Sal I	GTCTGAC	GTCGA ^{m5} C	GT ^{m5} CGAC GTCG ^{m5} AC#	Tfi I	GAWTC	GAWT ^{m5} C	?
Sau3A I	GATC ^a	G ^{m5} ATC	GAT ^{m5} C# ^b GAT ^{m4} C GATH ^{m5} C	Tha I	CGCG	^{m5} CGCG	^{m5} CGCG h ^{m5} CGh ^{m5} CG
Sau96 I	GGNCC	?	GGN ^{m5} CC# GGNC ^{m5} C GGNh ^{m5} Ch ^{m5} C	Xba I	TCTAGA	?	TCTAG ^{m5} A# Th ^{m5} CTAGA
Sca I	AGTACT	AGTA ^{m5} CT	?	Xho I	CTCGAG ^b	?	CT ^{m5} CGAG CTCG ^{m5} AG ^{m5} CTCGAG
ScrF I	CCNGG	^{m5} CCNGG	C ^{m5} CNGG C ^{m4} CNGG	Xho II	RGATCY	RG ^{m5} ATCY	RGAT ^{m5} CY ^b
SfaN I	GATGC	GATG ^{m5} C	G ^{m5} ATGC	Xma I	CCCGGG	CC ^{m5} CGGG ^b	^{m4} CCCGGG ^{m5} CCCGGG C ^{m4} CCGGG CC ^{m4} CGGG
Sfi I	GGCCN5GGCC	GG ^{m5} CCN5GG ^{m5} CC ^b	GGC ^{m5} CN5GGCC	Xma III	CGGCCG	?	CGGm5CCG
Sfi I	CTGCAG	?	CTGC ^{m5} AG	Xmn I	GAAN4TTC	GA ^{m5} AN4TTC	G ^{m5} AN4TTC GAAN4TT ^{m5} C ^b
SgrA I	CRCCGGYG	?	CRC ^{m5} CGGYG	Xor II	CGATCG	CG ^{m5} ATCG	CGAT ^{m5} CG h ^{m5} CGATH ^{m5} CG
Sin I	GGWCC	?	GGW ^{m5} CC#				
Sma I	CCCGGG	C ^{m5} CCGGG	^{m4} CCCGGG ^{m5} CCCGGG ^b C ^{m4} CCGGG ^b				

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R=G or A	Y=C or T	^{m4} C = N4-methylcytosine
M=A or C	K=G or T	^{m5} C = C5-methylcytosine
S=G or C	W=A or T	^{hm5} C = hydroxymethylcytosine
H=A or C or T	B=G or T or C	^m C = methylcytosine
V=G or C or A	D=G or A or T	N ₄ or C ₅ -methylcytosine unspecified
N=A or C or G or T		^{m6} A = N ₆ -methyladenine
		Sequences are in 5' - 3' order.

*Dpn I and its isoschizomers require the presence of 6-methyladenosine within the recognition sequence GATC

REFERENCE

1. Roberts, R.J. and Macelis, D. (1991) *Nucleic Acids Res.* 19: 2077-2109.
2. Nelson, M. and McClelland, M. (1991) *Nucleic Acids Res.* 19:2045-2071