

Restriction enzymes and their isoschizomers

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INTRODUCTION

Since the last compilation of restriction enzymes (1), 261 new entries have been added including 16 new specificities. With the growing size of this database and the recognition that the most widespread use of the information is as a database for computer programs predicting restriction enzyme cleavage patterns, the new format has been continued. This format is intended to contain the minimal amount of information required by a computer program. It should be noted that only enzymes for which the recognition sequence is known are included. This new list is shown in the first Table, while an alphabetical listing of all Type II enzymes, including those of unknown recognition sequence, is presented in the second Table. A copy of the restriction enzyme database in its previous format (2), including enzymes of unknown recognition sequence, will be available upon request. It should also be noted that an alternative compilation of these enzymes has recently been produced (3).

The database shown in these Tables is available, in a form that is easily reformatted, online through the GENBANK computer resource. It is also possible to get regular monthly updates, or specialized versions of the database by electronic mail. For instance, a file containing the database in a format that can be used directly by the UWGCG computer software package is available. Files to support other packages are either available or in preparation. Enquiries should be directed to roberts@cshlab.bitnet.

In forming this list, all endonucleases cleaving DNA at a specific sequence have been considered to be restriction enzymes, although in most cases there is no direct genetic evidence for the presence of a restriction-modification system. The endonucleases are named in accordance with the proposal of Smith and Nathans (4).

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Type I enzymes

Enzyme	Recognition sequence	Me site ³	Reference
<i>CfrAI</i>	GCANNNNNNNNGTGG		616,617
<i>EcoAI</i>	GAGNNNNNNNGTCA	2(6) -3(6)	618,619
<i>EcoBI</i>	TGANNNNNNNNNTGCT	3(6) -4(6)	620-624
<i>EcoDI</i>	TTANNNNNNNNGTCY		625
<i>EcoDXXI</i>	TCANNNNNNNNATTC		626,627
<i>EcoEI</i>	GAGNNNNNNNNATGC		628,629
<i>EcoKI</i>	AACNNNNNNNGTGC	2(6) -3(6)	630-633
<i>EcoR124I</i>	GAANNNNNNNRTCG		634
<i>EcoR124/3I</i>	GAANNNNNNNRTCG	-3(6)	634
<i>SrySBI</i>	GAGNNNNNNRTAYG	2(6) -4(6)	635
<i>SrySJI</i>	GAGNNNNNNNGTRC		636
<i>SrySPI</i>	AACNNNNNNNGTRC	2(6) -3(6)	635
<i>SrySQI</i>	AACNNNNNNRTAYG		637

Type II enzymes

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
<i>AarII</i>		GACGTIC		EGMNOPRSUVX	5
<i>AccI</i>		GTIMKAC		ABEGIKMNOPRSUVX	6
	<i>DsaVI</i>	GTMKAC			7
	<i>FblI</i>	GTIMKAC			8
<i>AccI</i>		GRICGYC		EMRV	9
	<i>AhaII</i>	GRICGYC		GN	10
	<i>AosII</i>	GRICGYC			11
	<i>AsiWI</i>	GRICGYC			12
	<i>AsuIII</i>	GRICGYC			12
	<i>BbiII</i>	GRICGYC		AK	13
	<i>HgiI</i>	GRICGYC			14
	<i>HgiDI</i>	GRICGYC			15
	<i>HgiGI</i>	GRICGYC			15
	<i>HgiHIII</i>	GRICGYC			16
	<i>HinII</i>	GRICGYC		OU	17
	<i>Hin8I</i>	GRCGYC			18
	<i>NlaSII</i>	GRCGYC			19
	<i>SspJII</i>	GRCGYC			20
	<i>SspM1II</i>	GRCGYC			20
	<i>SspM2II</i>	GRCGYC			20
<i>AfiIII</i>		CITTAAG		AGKNU	21
	<i>Esp4I</i>	CITTAAG			22
	<i>Vha464I</i>	CITTAAG			8
<i>AfiIII</i>		AICRYGT		GMU	21
<i>AgeI</i>		AICCGGT			23
<i>AhaIII</i>		TTTIAAA		E	24
	<i>DraI</i>	TTTIAAA		ABEGIKMNOPRSUVX	25
<i>AluI</i>		AGICT	3(5)	ABEGIKMNOPRSUVX	26-29
	<i>MliI</i>	AGICT			30,31
	<i>OtuI</i>	AGCT			32
	<i>OtuNI</i>	AGCT			33
	<i>OxaI</i>	AGCT			34
<i>AlwNI</i>		CAGNNNICTG		N	35
<i>ApaI</i>		GGGCCIC	4(5)	BEGIKMNOPRUVX	36,37
	<i>Bsp120I</i>	GIGGCC			38
	<i>EciEI</i>	GGGCC			39
<i>ApaLI</i>		GITGCAC		AEGKNX	40
	<i>Alw44I</i>	GITGCAC		O	41
	<i>AmeI</i>	GTGCAC			32
	<i>SnoI</i>	GITGCAC		MV	42,43
	<i>VneI</i>	GITGCAC			44

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
<i>AsuI</i>		GIGNCC		R	45
	<i>AhaBI</i>	GIGNCC			8
	<i>ApuI</i>	GGNCC			46
	<i>Bac36I</i>	GIGNCC			46
	<i>Bal228I</i>	GIGNCC			14
	<i>BavAII</i>	GIGNCC			47
	<i>Bce22I</i>	GIGNCC			8
	<i>BshKI</i>	GIGNCC			48
	<i>BspBII</i>	GIGNCC			49
	<i>Cfr4I</i>	GGNCC			50,51
	<i>Cfr8I</i>	GGNCC			50,51
	<i>Cfr13I</i>	GIGNCC	4(5)	AKOU	50,52
	<i>Cfr23I</i>	GGNCC			53
	<i>Cfr33I</i>	GGNCC			54
	<i>Cfr45I</i>	GGNCC			54
	<i>Cfr46I</i>	GGNCC			54
	<i>Cfr47I</i>	GGNCC			54
	<i>Cfr52I</i>	GGNCC			55
	<i>Cfr54I</i>	GGNCC			56
	<i>CfrNI</i>	GGNCC			57
	<i>Eco39I</i>	GGNCC			58
	<i>Eco47II</i>	GGNCC			59
	<i>Eco196II</i>	GGNCC			60
	<i>Eco201I</i>	GGNCC			60
	<i>GseI</i>	GGNCC			61
	<i>Hin5II</i>	GGNCC			18
	<i>MjaII</i>	GGNCC			62
	<i>Msp24I</i>	GGNCC			8
	<i>MitTI</i>	GGNCC			63
	<i>NlaDII</i>	GGNCC			64
	<i>NmuEII</i>	GGNCC			65
	<i>NmuSI</i>	GGNCC			66
	<i>NspIV</i>	GIGNCC			67
	<i>NspLII</i>	GGNCC			68,69
	<i>Pde12I</i>	GIGNCC			8
	<i>PseI</i>	GGNCC			70
	<i>PspI</i>	GGNCC			71
	<i>Sau96I</i>	GIGNCC		BEGMNRVX	72
	<i>SauBI</i>	GGNCC			73
	<i>SdyI</i>	GGNCC			74
<i>AsuII</i>		TTTCGAA		G	12,75
	<i>AcaI</i>	TTCGAA			76,77
	<i>AviI</i>	TTCGAA			70
	<i>Bim19I</i>	TTTCGAA			8
	<i>Bpu14I</i>	TTTCGAA			8
	<i>BsiCI</i>	TTCGAA			78
	<i>Bsp82I</i>	TTCGAA			79
	<i>Bsp119I</i>	TTCGAA			55
	<i>BstBI</i>	TTTCGAA		N	80,81
	<i>Csp45I</i>	TTTCGAA		R	82
	<i>FspII</i>	TTTCGAA			83
	<i>LspI</i>	TTTCGAA		V	42,43
	<i>MlaI</i>	TTTCGAA			84
	<i>NspV</i>	TTCGAA		AKOPU	67
	<i>NspBI</i>	TTCGAA			85
	<i>NspFI</i>	TTCGAA			61
	<i>NspJI</i>	TTCGAA			69,77
	<i>SfuI</i>	TTTCGAA		M	86
	<i>Sgr1839I</i>	TTCGAA			87
	<i>SspII</i>	TTTCGAA			88
<i>AvaI</i>	<i>Ssp152I</i>	TTCGAA			87
		CYCGRG		ABEGIKMNOPRSUVX	89,90
	<i>AcrI</i>	CYCGRG			77
	<i>Ama87I</i>	CYCGRG			8

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>AquI</i>	C↓YCGRG	1(5)		91,92
	<i>AspBI</i>	CYCGRG			85
	<i>AspCI</i>	CYCGRG			85
	<i>AspDI</i>	CYCGRG			85
	<i>AvrI</i>	CYCGRG			93
	<i>BcoI</i>	C↓YCGRG			94
	<i>BstSI</i>	C↓YCGRG			80
	<i>Eco88I</i>	CYCGRG			95
	<i>NliI</i>	CYCGRG			85
	<i>NmuAI</i>	CYCGRG			85
	<i>NspIII</i>	C↓YCGRG		PV	67
	<i>NspDI</i>	CYCGRG			85
	<i>NspEI</i>	CYCGRG			96
	<i>NspSAI</i>	C↓YCGRG			97
<i>AvaII</i>		G↓GWCC		ABEGIKMNPRSVX	89,90,98,99
	<i>AflI</i>	G↓GWCC			21
	<i>Asp697I</i>	GGWCC			100
	<i>Asp745I</i>	G↓GWCC			14
	<i>AspBII</i>	GGWCC			85
	<i>AspCII</i>	GGWCC			85
	<i>AspDII</i>	GGWCC			85
	<i>BamNxi</i>	G↓GWCC			101–103
	<i>Bme18I</i>	G↓GWCC			104
	<i>Bme216I</i>	G↓GWCC			105,106
	<i>Bsp100I</i>	GGWCC			38
	<i>BspJ105I</i>	GGWCC			107
	<i>BtrI</i>	GGWCC			108
	<i>CauI</i>	G↓GWCC			109,110
	<i>ClfI</i>	GGWCC			68
	<i>ClnII</i>	GGWCC			111
	<i>DsaIV</i>	G↓GWCC			7
	<i>EagMI</i>	G↓GWCC			112,113
	<i>Eco47I</i>	G↓GWCC		OU	59,114
	<i>ErpI</i>	G↓GWCC			46
	<i>FdiI</i>	G↓GWCC			115,116
	<i>FspMSI</i>	G↓GWCC			46
	<i>GspAI</i>	GGWCC			78
	<i>HgiBI</i>	G↓GWCC			15
	<i>HgiCII</i>	G↓GWCC			15
	<i>HgiEI</i>	G↓GWCC			15
	<i>HgiHIII</i>	G↓GWCC			16
	<i>HgiJI</i>	G↓GWCC			16
	<i>Hsp2I</i>	GGWCC			55
	<i>Kzo49I</i>	G↓GWCC			8
	<i>MfoI</i>	GGWCC			117
	<i>MliI</i>	GGWCC			118
	<i>MspAI</i>	GGWCC			117
	<i>NliII</i>	GGWCC			85
	<i>NmuAII</i>	GGWCC			85
	<i>NspDII</i>	GGWCC			85
	<i>NspGI</i>	GGWCC			69,77
	<i>NspHII</i>	GGWCC			85
	<i>NspKI</i>	GGWCC			68,69
	<i>SfiI</i>	GGWCC			119
	<i>Sgh1835I</i>	GGWCC			87
	<i>SinI</i>	G↓GWCC	4(5)	RSV	70,120,121
	<i>SinAI</i>	GGWCC			122
	<i>SinBI</i>	GGWCC			122
	<i>SinCI</i>	GGWCC			122
	<i>SinDI</i>	GGWCC			122
	<i>SinEI</i>	GGWCC			122
	<i>SinFI</i>	GGWCC			122
	<i>SinGI</i>	GGWCC			122
	<i>SinHI</i>	GGWCC			122

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Sin</i> II	GGWCC			122
	<i>Tru</i> I	GGWCC			123
<i>Ava</i> III		ATGCAT		G	124,125
	<i>Eco</i> T22I	ATGCAIT		KOU	126
	<i>Nsi</i> I	ATGCAIT		BEMNRVX	127
	<i>Uba</i> 1103I	ATGCAIT			128
	<i>Zsp</i> 2I	ATGCAIT			8
<i>Avr</i> II		C!CTAGG		N	93,129
<i>Bal</i> I		TGG!CCA	4(5)	ABEGIKNRSVX	37,130
	<i>Msc</i> I	TGG!CCA		N	32,131
	<i>Msp</i> 20I	TGG!CCA			8
<i>Bam</i> HI		G!GATCC	5(4)	ABEGIKMNOPRSUVX	132–135
	<i>Aac</i> I	GGATCC			136
	<i>Aae</i> I	GGATCC			136
	<i>Aca</i> II	GGATCC			77
	<i>Acc</i> EBI	G!GATCC			137
	<i>Ain</i> II	GGATCC			77
	<i>Al</i> I	G!GATCC			138
	<i>Al</i> i12257I	GGATCC			139
	<i>Al</i> i12258I	GGATCC			139
	<i>Asp</i> TII	GGATCC			77
	<i>Bam</i> FI	GGATCC			140
	<i>Bam</i> KI	GGATCC			140
	<i>Bam</i> NI	GGATCC			102
	<i>Bca</i> 1259I	GGATCC			87
	<i>Bce</i> 751I	G!GATCC			141
	<i>Bco</i> 10278I	GGATCC			142
	<i>Bna</i> I	G!GATCC	5(5)		143,144
	<i>Bsp</i> 30I	GGATCC			38
	<i>Bsp</i> 46I	GGATCC			79
	<i>Bsp</i> 98I	GGATCC			38
	<i>Bsr</i> I	G!GATCC		GPV	145,146
	<i>Bsr</i> 1126I	GGATCC			87
	<i>Bsr</i> 2464I	GGATCC			87
	<i>Bsr</i> 2902I	GGATCC			87
	<i>Bsr</i> QI	GGATCC			80
	<i>Bsu</i> 90I	GG!ATCC			141
	<i>Bsu</i> 8565I	GGATCC			142
	<i>Bsu</i> 8646I	GGATCC			142
	<i>Cel</i> I	GGATCC			77
	<i>Dds</i> I	GGATCC			147
	<i>Gdo</i> I	GGATCC			136
	<i>Gin</i> I	GGATCC			148
	<i>Gox</i> I	GGATCC			136
	<i>Gse</i> III	GGATCC			61
	<i>Mle</i> I	GGATCC			30
	<i>Mlu</i> 23I	G!GATCC			8
	<i>Nas</i> BI	GGATCC			30
	<i>Nsp</i> SAIV	G!GATCC			97
	<i>Pae</i> 177I	GGATCC			60
	<i>Rhs</i> I	GGATCC			149
	<i>Rlu</i> 4I	GGATCC			150
	<i>Sur</i> 2I	G!GATCC			8
	<i>Uba</i> 1110I	GGATCC			151
<i>Bbv</i> I		GCAGC(8/12)	2(5), –2(5)	EGINX	134,152–154
	<i>Alw</i> XI	GCAGC(8/12)			155
	<i>Bsp</i> 423I	GCAGC			156
	<i>Uba</i> 1109I	GCAGC			157
<i>Bbv</i> II		GAAGAC(2/6)			158
	<i>Bbs</i> I	GAAGAC			159
	<i>Bbv</i> 16I	GAAGAC(2/6)			8
	<i>Bsp</i> VI	GAAGAC			156
<i>Bce</i> II		ACGGC(12/13)			160
<i>Bcg</i> I ⁵		GCANNNNNNTCG			161

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
<i>BclI</i>		TIGATCA		BEGINNOPRSUVX	162
	<i>AtuCI</i>	TGATCA			163
	<i>BsiQI</i>	TGATCA			78
	<i>BspXI</i>	TIGATCA			164
	<i>BsrGI</i>	TGATCA			165
	<i>BsrKI</i>	TGATCA			80
	<i>CpeI</i>	TGATCA			166
	<i>CthI</i>	TGATCA			167
	<i>FbaI</i>	TGATCA			119
	<i>Ksp22I</i>	TIGATCA			8
	<i>PovI</i>	TGATCA			112,168
	<i>SseI</i>	TGATCA			61
	<i>SstIV</i>	TGATCA			169
<i>BglII</i>		GCCNNNNINGGC		BEGINNOPRSUVX	170-174
	<i>VanI</i>	GCCNNNNNNGGC			175
<i>BglIII</i>		AIGATCT		ABEGIKMNOPRSUVX	171,172,176
	<i>NspMACI</i>	AIGATCT			177
<i>BinI</i>		GGATC(4/5)			178
	<i>AlwI</i>	GGATC(4/5)	N		159
	<i>BthII</i>	GGATC			179
<i>Bpu10I</i>		CCTNAGC(-5/-2)			180
<i>BsaAI</i>		YACIGTR	N		81
	<i>MspYI</i>	YACIGTR			181
<i>BsaBI</i>		GATNNINNNATC	N		80
	<i>MamI</i>	GATNNINNNATC	M		182
<i>BsePI</i>		GCGCGC			165
	<i>BsoPI</i>	GCGCGC			165
	<i>BsrHI</i>	GCGCGC			165
	<i>BssHII</i>	GICGCGC		EGMNOUVX	165,183
	<i>Eco143I</i>	GCGCGC			150
	<i>Esp7I</i>	GCGCGC			79
	<i>Esp8I</i>	GCGCGC			79
	<i>Kpn30I</i>	GCGCGC			184
<i>BsiI</i>		CTCGTG(-5/-1)			185
<i>BsmI</i>		GAATGC(1/-1)		EGNOUVX	75
<i>BsmAI</i>		GTCTC(1/5)	N		
	<i>Alw26I</i>	GTCTC(1/5)			186
<i>BspCI</i>		GCNNINNGC			156
<i>BspGI</i>		CTGGAC			156
<i>BspHI</i>		TICATGA	N		187
	<i>RspXI</i>	TICATGA	G		188
<i>BspMI</i>		ACCTGC(4/8)	N		159,189
<i>BspMII</i>		TICCGGA	N		159,189
	<i>AccIII</i>	TICCGGA		AEGKR	75,190
	<i>Bbf7411I</i>	TCCGGA			191
	<i>BbvAIII</i>	TICCGGA			192
	<i>Bla7920I</i>	TCCGGA			191
	<i>BseAI</i>	TICCGGA			193
	<i>BsiMI</i>	TCCGGA			78
	<i>BsiOI</i>	TCCGGA			78
	<i>Bsp13I</i>	TICCGGA			8
	<i>BspeI</i>	TICCGGA			161
	<i>Bsu22I</i>	TCCGGA			141
	<i>CauB3I</i>	TICCGGA			194
	<i>Kpn2I</i>	TICCGGA			38,195
	<i>MroI</i>	TICCGGA		MOU	196
<i>BsrI</i>		ACTGG(1/-1)	N		197
<i>BstEII</i>		GIGTNACC		BEGMNOPRSUVX	198,199
	<i>AcrII</i>	GIGTNACC			77
	<i>AspAI</i>	GIGTNACC			43
	<i>BsiKI</i>	GGTNACC			78
	<i>Bst31I</i>	GGTNACC			200
	<i>BstDI</i>	GGTNACC			80
	<i>BstPI</i>	GIGTNACC	K		201

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Cfr</i> 7I	GGTNACC			50
	<i>Cfr</i> 19I	GGTNACC			54
	<i>Eca</i> I	G↓GTNACC	5(6)		202
	<i>Eco</i> 9II	G↓GTNACC			203,204
	<i>Eco</i> O65I	G↓GTNACC		G	73,205
	<i>Kox</i> I	GGTNACC			206
	<i>Nsp</i> SAII	G↓GTNACC			97
	<i>Sci</i> AI	GGTNACC			77
<i>Bst</i> XI		CCANNNNN↓NTGG		EGMNORUVX	165,207
	<i>Bss</i> GI	CCANNNNNNTGG			165
	<i>Bst</i> TI	CCANNNNNNTGG			165
<i>Cau</i> II		CC↓SGG			109,110,208
	<i>Aha</i> I	CC↓SGG			10
	<i>Ase</i> II	CC↓SGG			209
	<i>Asp</i> 1I	CCSGG			150
	<i>Bcn</i> I	CC↓SGG	2(4)	AK	210–215
	<i>Bsp</i> 7I	CCSGG			38
	<i>Bsp</i> 8I	CCSGG			38
	<i>Bsp</i> 55I	CCSGG			79
	<i>Bsp</i> J67I	CCSGG			107
	<i>Eci</i> DI	CCSGG			39
	<i>Eco</i> 121I	CCSGG			216
	<i>Eco</i> 179I	CCSGG			60
	<i>Eco</i> 190I	CCSGG			216
	<i>Hgi</i> S21I	CCSGG			217
	<i>Hgi</i> S22I	CC↓SGG			14
	<i>Hin</i> 3I	CCSGG			218
	<i>Nci</i> I	CC↓SGG		BEGMNOUVX	219,220
	<i>Pae</i> 181I	CCSGG			60
	<i>Rsh</i> II	CCSGG			105
	<i>Ssp</i> 2I	CCSGG			150
	<i>Tmu</i> 1I	CCSGG			203
<i>Cfr</i> I		Y↓GGCCR	4(5)		51,215,221,222
	<i>Cfr</i> 14I	YGGCCR			50
	<i>Cfr</i> 38I	YGGCCR			53
	<i>Cfr</i> 39I	YGGCCR			54
	<i>Cfr</i> 40I	YGGCCR			54
	<i>Eae</i> I	Y↓GGCCR	4(5)	EGKMN VX	223,224
	<i>Eci</i> BI	YGGCCR			39
	<i>Eco</i> 90I	YGGCCR			203
	<i>Eco</i> 164I	YGGCCR			150
	<i>Eco</i> HI	YGGCCR			33
<i>Cfr</i> 10I		R↓CCGGY	2(5)	AKMOU	50,51,215,225
	<i>Bsp</i> 21I	RCCGGY			203
<i>Cl</i> I		AT↓CGAT	5(6)	ABEGKMNPRSVX	226,227
	<i>Aag</i> I	AT↓CGAT			228
	<i>Apu</i> 16I	ATCGAT			229
	<i>Asp</i> 707I	ATCGAT			100
	<i>Ban</i> III	ATCGAT		OU	5
	<i>Bbv</i> AII	AT↓CGAT			230
	<i>Bcm</i> I	AT↓CGAT			231
	<i>Bdi</i> I	AT↓CGAT			232
	<i>Bli</i> 41I	AT↓CGAT			8
	<i>Bli</i> 86I	AT↓CGAT			141
	<i>Bli</i> R1	ATCGAT			233
	<i>Bsi</i> I	AT↓CGAT			43,234
	<i>Bsp</i> 2I	ATCGAT			38
	<i>Bsp</i> 4I	ATCGAT			38
	<i>Bsp</i> 84I	ATCGAT			79
	<i>Bsp</i> 106I	AT↓CGAT			235
	<i>Bsp</i> DI	AT↓CGAT			161
	<i>Bsp</i> XI	AT↓CGAT			164
	<i>Bsu</i> 15I	AT↓CGAT			229,236
	<i>Bru</i> I	ATCGAT			237

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Csp4I</i>	ATCGAT			79
	<i>LplI</i>	AT↓CGAT			238
	<i>PglIB4I</i>	AT↓CGAT			8
<i>CviI</i>		RG↓CY	3(5)		107,239
	<i>CviKI</i>	RGCY			240
	<i>CviLI</i>	RGCY			240
	<i>CviMI</i>	RGCY			240
	<i>CviNI</i>	RGCY			240
	<i>CviOI</i>	RGCY			240
<i>DdeI</i>		C↓TNAG	1(5)	BEGINNOPRUVX	241–243
<i>DpnI</i> *		GA↓TC		ABEGINNRSVX	244–246
	<i>CfuI</i> *	GA↓TC			247,248
	<i>NanII</i> *	GATC			249
	<i>NgoDIII</i> *	GATC			250
	<i>NmuDI</i> *	GATC			65
	<i>NmuEI</i> *	GATC			65
	<i>NsuDI</i> *	GATC			65
	<i>Uba1103II</i> *	GATC			128
<i>DraII</i>		RG↓GNCCY		EGM	251–253
	<i>EcoO109I</i>	RG↓GNCCY		AGKNOUVX	254
	<i>PssI</i>	RGGNC↓CY		I	147,255
	<i>VneAI</i>	RGGNCCY			175
<i>DraIII</i>		CACNNN↓GTG		EMNX	251–253
<i>DrdI</i>		GACNNNN↓NNGTC		N	256
<i>DrdII</i>		GAACCA			257
<i>DsaI</i>		C↓CRYGG		M	258
<i>EciI</i>		TCCGCC			39
<i>Eco31I</i>		GGTCTC(1/5)			259
	<i>Bli49I</i>	GGTCTC			141
	<i>BsaI</i>	GGTCTC(1/5)		N	81
	<i>Eco42I</i>	GGTCTC			216
	<i>Eco51I</i>	GGTCTC			150
	<i>Eco95I</i>	GGTCTC			260
	<i>Eco97I</i>	GGTCTC			261
	<i>Eco101I</i>	GGTCTC			216
	<i>Eco120I</i>	GGTCTC			216
	<i>Eco127I</i>	GGTCTC			60
	<i>Eco129I</i>	GGTCTC			60
	<i>Eco155I</i>	GGTCTC			216
	<i>Eco156I</i>	GGTCTC			216
	<i>Eco157I</i>	GGTCTC			216
	<i>Eco162I</i>	GGTCTC			60
	<i>Eco185I</i>	GGTCTC			60
	<i>Eco191I</i>	GGTCTC			216
	<i>Eco203I</i>	GGTCTC			216
	<i>Eco204I</i>	GGTCTC			216
	<i>Eco205I</i>	GGTCTC			216
	<i>Eco217I</i>	GGTCTC			55
	<i>Eco225I</i>	GGTCTC			150
	<i>Eco233I</i>	GGTCTC			38
	<i>Eco239I</i>	GGTCTC			150
	<i>Eco240I</i>	GGTCTC			55
	<i>Eco241I</i>	GGTCTC			55
	<i>Eco246I</i>	GGTCTC			38
	<i>Eco247I</i>	GGTCTC			38
	<i>EcoA4I</i>	GGTCTC(1/5)			8
	<i>PpaI</i>	GGTCTC			159
	<i>Sau12I</i>	GGTCTC			203
<i>Eco47III</i>		AGC↓GCT		AKMOU	59
	<i>AiiI</i>	AGC↓GCT			262
<i>Eco57I</i>		CTGAAG(16/14)			263
	<i>Bsp6II</i>	CTGAAG			203
	<i>Eco125I</i>	CTGAAG			60
	<i>FsfI</i>	CTGAAG			264

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
<i>EcoNI</i>		CCTNN1NNNAGG		N	265
	<i>BstWI</i>	CCTNNNNNAGG			81
<i>EcoRI</i>		G1AATTC	3(6)	ABEGIKMNOPRSUVX	266-268
	<i>Eco82I</i>	GAATTC			260
	<i>Eco159I</i>	GAATTC			60
	<i>Eco228I</i>	GAATTC			55
	<i>Eco237I</i>	GAATTC			150
	<i>Eco252I</i>	GAATTC			150
	<i>HalB6I</i>	G1AATTC			8
	<i>RsrI</i>	G1AATTC			269-271
	<i>SsoI</i>	G1AATTC	2(6)		272,273
<i>EcoRII</i> ⁶		1CCWGG	2(5)	BEGOV	274-276
	<i>Acc38I</i>	CCWGG			8
	<i>AeuI</i>	CC1WGG			277
+	<i>AorI</i>	CC1WGG			136
+	<i>ApyI</i>	CC1WGG		M	278
	<i>AtuII</i>	CCWGG			279
	<i>Atu1I</i>	CCWGG			203
	<i>AtuBI</i>	CCWGG			280
	<i>BinSI</i>	CCWGG			179
+	<i>BshGI</i>	CC1WGG			281
+	<i>BsiLI</i>	CCWGG			78
+	<i>Bsp56I</i>	CCWGG			79
	<i>Bsp103I</i>	CCWGG			38
	<i>BstGII</i>	CCWGG			165
+	<i>BstNI</i>	CC1WGG		ENX	282
	<i>BstOI</i>	CCWGG			80
	<i>Cdi27I</i>	CCWGG			184
	<i>Cfr5I</i>	CCWGG			50,51
	<i>Cfr11I</i>	CCWGG			50,51
	<i>Cfr20I</i>	CCWGG			54
	<i>Cfr22I</i>	CCWGG			54
	<i>Cfr24I</i>	CCWGG			54
	<i>Cfr25I</i>	CCWGG			54
	<i>Cfr27I</i>	CCWGG			54
	<i>Cfr28I</i>	CCWGG			54
	<i>Cfr29I</i>	CCWGG			54
	<i>Cfr30I</i>	CCWGG			54
	<i>Cfr31I</i>	CCWGG			54
	<i>Cfr35I</i>	CCWGG			54
	<i>CfrS37I</i>	CCWGG			217
	<i>CthII</i>	CC1WGG			283
	<i>EagKI</i>	CCWGG			284
	<i>EcalI</i>	CCWGG			75
	<i>EclII</i>	CCWGG			285
	<i>Ecl66I</i>	CCWGG			261
	<i>Ecl136I</i>	CCWGG			60
	<i>Ecl137II</i>	CCWGG			216
	<i>EclS39I</i>	CCWGG			217
	<i>Eco38I</i>	CCWGG			58
	<i>Eco40I</i>	CCWGG			58
	<i>Eco41I</i>	CCWGG			58
	<i>Eco60I</i>	CCWGG			51
	<i>Eco61I</i>	CCWGG			51
	<i>Eco67I</i>	CCWGG			260
	<i>Eco70I</i>	CCWGG			260
	<i>Eco71I</i>	CCWGG			286
	<i>Eco128I</i>	CCWGG			60
	<i>Eco170I</i>	CCWGG			150
	<i>Eco193I</i>	CCWGG			216
	<i>Eco206I</i>	CCWGG			38
	<i>Eco207I</i>	CCWGG			38
	<i>Esp2I</i>	CCWGG			38
	<i>Fsp1604I</i>	CC1WGG			287

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
+	<i>Kax165I</i>	CCWGG			150
	<i>Kpn10I</i>	CCWGG			60
	<i>Kpn13I</i>	CCWGG			60
	<i>Kpn14I</i>	CCWGG			216
	<i>Kpn16I</i>	CCWGG			216
	<i>Mlu2300I</i>	CCWGG			8
	<i>MphI</i>	CCWGG			288
	<i>MvaI</i>	CC1WGG	2(4)	AKOU	289,290
	<i>Scg2I</i>	CCWGG			291
	<i>Sft2aI</i>	CCWGG			292
	<i>Sft2bI</i>	CCWGG			292
	<i>Sgr20I</i>	CCWGG			293
	<i>SleI</i>	CCWGG			294
	<i>SspAI</i>	CCWGG			294
	<i>TaqXI</i>	CC1WGG			295
<i>EcoRV</i>	<i>ZanI</i>	CC1WGG			296
		GAT1ATC	2(6)	ABEGIKMNOPRSUVX	297–299
	<i>Bsp16I</i>	GATATC			38
	<i>BstRI</i>	GATATC			80
	<i>CeqI</i>	GAT1ATC			300
	<i>Eco32I</i>	GAT1ATC			51,301
	<i>Eco178I</i>	GATATC			60
	<i>HjaI</i>	GAT1ATC			302
	<i>NanI</i>	GATATC			249
	<i>NfiAI</i>	GATATC			303
<i>EspI</i>	<i>NsiCI</i>	GAT1ATC			304
		GC1TNAGC		EGU	305
	<i>CeIII</i>	GCTNAGC		M	77
	<i>Uba1102I</i>	GC1TNAGC			306
<i>Esp3I</i>		GAGACG(1/5)			79
<i>FauI</i>		CCCGC(4/6)			307
<i>FinI</i>		GTCCC			159
<i>Fnu4HI</i>		GC1NGC		N	308
	<i>Bsp6I</i>	GC1NGC			309
	<i>FbrI</i>	GC1NGC			119
	<i>IspI</i>	GC1NGC			181
	<i>Uur960I</i>	GC1NGC			310
<i>FnuDII</i>		CG1CG			311
	<i>AccII</i>	CG1CG		AEGKVX	6,312
	<i>BceFI</i>	CGCG			313
	<i>BceRI</i>	CGCG			140
	<i>BepI</i>	CG1CG	1(5)		314,315
	<i>Bpu95I</i>	CG1CG			8
	<i>Bsp50I</i>	CG1CG			79
	<i>Bsp70I</i>	CGCG			79
	<i>BspJ76I</i>	CGCG			107
	<i>BstUI</i>	CG1CG		N	80
	<i>Bsu1192II</i>	CGCG			74
	<i>Bsu1193I</i>	CGCG			74,140
	<i>Bsu1532I</i>	CG1CG			316
	<i>Bsu6633I</i>	CGCG			140,317
	<i>BsuEII</i>	CGCG	1(5)		74,318,319
	<i>Cpa1150I</i>	CGCG			87
	<i>CpaAI</i>	CGCG			320
	<i>FspMI</i>	CGCG			159
	<i>Hin1056I</i>	CGCG			321
	<i>MvaAI</i>	CGCG			117
<i>FokI</i>	<i>MvnI</i>	CG1CG		M	322
	<i>PflAI</i>	CGCG			32
	<i>Scel</i>	CGCG			77
	<i>Thal</i>	CG1CG		BI	323
	<i>Tmal</i>	CGCG			63
		GGATG(9/13)	3(6), –2(6)	AEGIKMNUVX	324 – 327
	<i>HinGUII</i>	GGATG			328 – 330

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
<i>FseI</i>		GGCCGG↓CC			331
<i>GdiII</i>		YGGCCG(−5/−1)			332
<i>GsuI</i>		CTGGAG(16/14)			263,333
	<i>Bco35I</i>	CTGGAG			55
	<i>Bsp22I</i>	CTGGAG			203
	<i>Bsp28I</i>	CTGGAG			38
	<i>Bsp174I</i>	CTGGAG			107
<i>HaeI</i>		WGG↓CCW			334
<i>HaeII</i>		RGCGC↓Y		ABEGIKMNOPRSUX	335,336
	<i>Bme142I</i>	RGC↓GCT			337
	<i>HinHI</i>	RGCGCT			338
	<i>NgoI</i>	RGCGCT			339
<i>HaeIII</i>		GG↓CC	3(5)	ABEGIKMNOPRSUVX	340–342
	<i>AclIV</i>	GGCC			77
	<i>Asp742I</i>	GGCC			217
	<i>AspTIII</i>	GGCC			77
	<i>Bal475I</i>	GGCC			87
	<i>Bal3006</i>	GGCC			87
	<i>Bce71I</i>	GGCC			343
	<i>Bco33I</i>	GGCC			38
	<i>Bfi458I</i>	GGCC			87
	<i>Bim19II</i>	GG↓CC			8
	<i>BlI</i>	GGCC			344
	<i>BluII</i>	GGCC			332
	<i>BseI</i>	GGCC			345
	<i>BshI</i>	GGCC			78
	<i>BshAI</i>	GGCC			78
	<i>BshBI</i>	GGCC			78
	<i>BshCI</i>	GGCC			78
	<i>BshDI</i>	GGCC			78
	<i>BshEI</i>	GGCC			78
	<i>BshFI</i>	GG↓CC			346
	<i>BsiAI</i>	GGCC			78
	<i>BsiDI</i>	GGCC			78
	<i>BsiHI</i>	GGCC			78
	<i>Bsp71I</i>	GGCC			343
	<i>Bsp211I</i>	GG↓CC			343
	<i>Bsp226I</i>	GGCC			343
	<i>BspBRI</i>	GG↓CC			347
	<i>BspRI</i>	GG↓CC			348–350
	<i>BssCI</i>	GGCC			165
	<i>BstCI</i>	GGCC		G	165
	<i>BstJI</i>	GGCC			80
	<i>Bsu1076I</i>	GGCC			140
	<i>Bsu1114I</i>	GGCC			140
	<i>BsuRI</i>	GG↓CC	3(5)	G	341,351,352
	<i>BteI</i>	GGCC			233
	<i>ClmI</i>	GGCC			111
	<i>ClfI</i>	GG↓CC			353
	<i>Csp2I</i>	GGCC			56
	<i>DsaII</i>	GG↓CC			258
	<i>FinSI</i>	GGCC			119
	<i>FnuDI</i>	GG↓CC			311
	<i>HhgI</i>	GGCC			75
	<i>MniI</i>	GGCC			288
	<i>MnnII</i>	GGCC			354
	<i>NgoII</i>	GGCC			355
	<i>NgoPII</i>	GG↓CC			304
	<i>NgoSI</i>	GGCC			356
	<i>NlaI</i>	GGCC			357
	<i>PaiI</i>	GGCC			148
	<i>PaiI</i>	GG↓CC		EPV	358,359
	<i>Pde133I</i>	GG↓CC			8
	<i>PpuI</i>	GGCC			148

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>SfaI</i>	GG CC			360
	<i>SpIII</i>	GGCC			361
	<i>SuaI</i>	GG CC			362
	<i>SuII</i>	GGCC			363
	<i>TspZNI</i>	GGCC			364
	<i>TreAI</i>	GGCC			365
	<i>TniI</i>	GGCC			116
	<i>VhaI</i>	GGCC			149
	<i>VniI</i>	GGCC			175
<i>HgaI</i>		GACGC(5/10)		NX	338,366,367
<i>HgiAI</i>		GWGCW C		NX	368
	<i>Alw21I</i>	GWGCW C			41
	<i>AspHI</i>	GWGCW C		M	369
	<i>Bbv12I</i>	GWGCW C			8
	<i>Pph3215I</i>	GWGCWC			87
<i>HgiCI</i>		G GYRCC			15,370
	<i>AccB1I</i>	G GYRCC			8
	<i>BanI</i>	G GYRCC		EGIMNOPUVX	5,370
	<i>Eco50I</i>	GGYRCC			150
	<i>Eco64I</i>	G GYRCC			260,371
	<i>Eco168I</i>	GGYRCC			216
	<i>Eco169I</i>	GGYRCC			150
	<i>Eco171I</i>	GGYRCC			60
	<i>Eco173I</i>	GGYRCC			60
	<i>Eco195I</i>	GGYRCC			216
	<i>Esp1I</i>	GGYRCC			55
	<i>Esp6I</i>	GGYRCC			79
	<i>Esp9I</i>	GGYRCC			79
	<i>Esp10I</i>	GGYRCC			79
	<i>Esp11I</i>	GGYRCC			79
	<i>Esp12I</i>	GGYRCC			79
	<i>Esp13I</i>	GGYRCC			79
	<i>Esp14I</i>	GGYRCC			79
	<i>Esp15I</i>	GGYRCC			150
	<i>Esp22I</i>	GGYRCC			150
	<i>HgiHI</i>	G GYRCC			16
	<i>MspB4I</i>	G GYRCC			8
	<i>SspM1III</i>	GGYRCC			20
<i>HgiEII</i>		ACCNNNNNNGGT			15
<i>HgiJII</i>		GRG CY C			16
	<i>BanII</i>	GRG CY C		EGIKMNOPRSUVX	5
	<i>BpuI</i>	GRG CYC			161,372
	<i>Bsp117I</i>	GRG CYC			38
	<i>Bsp519I</i>	GRG CY C			8
	<i>Bsu1854I</i>	GRG CY C			316
	<i>BvuI</i>	GRG CY C			373
	<i>Cfr48I</i>	GRG CYC			54
	<i>Eco24I</i>	GRG CY C			184,374
	<i>Eco25I</i>	GRG CYC			184
	<i>Eco26I</i>	GRG CYC			51
	<i>Eco35I</i>	GRG CYC			184
	<i>Eco68I</i>	GRG CYC			261
	<i>Eco113I</i>	GRG CYC			216
	<i>Eco180I</i>	GRG CYC			60
	<i>Eco211I</i>	GRG CYC			38
	<i>Eco215I</i>	GRG CYC			203
	<i>Eco216I</i>	GRG CYC			203
	<i>Eco232I</i>	GRG CYC			55
	<i>EcoT38I</i>	GRG CYC			126
	<i>EcoT88I</i>	GRG CYC			375
	<i>EcoT93I</i>	GRG CYC			375
	<i>EcoT95I</i>	GRG CYC			375
	<i>KoxII</i>	GRG CY C			206
<i>HhaI</i>		GCG C	2(5)	ABEGKNOPRSUX	376,377
	<i>BcaI</i>	GCGC			33

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>CfoI</i>	GCG C		BIMRV	147
	<i>Csp1470I</i>	GCGC			87
	<i>FnuDIII</i>	GCG C			311
	<i>Hin6I</i>	G CGC			378
	<i>Hin7I</i>	GCGC			18
	<i>HinGUI</i>	GCGC			328,379
	<i>HinP1I</i>	G CGC		NX	380
	<i>HinS1I</i>	GCGC			380
	<i>HinS2I</i>	GCGC			380
	<i>MnnIV</i>	GCGC			354
	<i>ScrNI</i>	G CGC			381
<i>HindII</i>		GTY CAC	5(6)	M	382–385
	<i>ChuII</i>	GTYRAC			386
	<i>Hin1160II</i>	GTYRAC			321
	<i>Hin1161II</i>	GTYRAC			321
	<i>HinJC1</i>	GTY CAC			387
	<i>HincII</i>	GTY CAC		ABEGIKNOPRSUVX	388
	<i>MnnI</i>	GTYRAC			354
<i>HindIII</i>		A AGCTT	1(6)	ABEGIKMNOPRSUVX	384,385,389
	<i>Asp52I</i>	AAGCTT			217
	<i>Asp3065I</i>	AAGCTT			390
	<i>BbrI</i>	AAGCTT			75
	<i>BpeI</i>	AAGCTT			391,392
	<i>BstFI</i>	A AGCTT			393
	<i>Cfr32I</i>	AAGCTT			54
	<i>ChuI</i>	AAGCTT			386
	<i>EcoVIII</i>	A AGCTT			394
	<i>Eco65I</i>	AAGCTT			261
	<i>Eco98I</i>	AAGCTT			261
	<i>Eco188I</i>	AAGCTT			216
	<i>Eco231I</i>	AAGCTT			55
	<i>Hin5III</i>	AAGCTT			18
	<i>Hin173I</i>	AAGCTT			328
	<i>Hin1076III</i>	AAGCTT			321
	<i>HinJCII</i>	AAGCTT			387
	<i>HinbIII</i>	AAGCTT			75,395
	<i>HinFI</i>	AAGCTT			396
	<i>HsuI</i>	A AGCTT			75
	<i>MkiI</i>	AAGCTT			288
<i>HinFI</i>		G ANTC		ABEGIKMNOPRSUVX	395,397,398
	<i>CviBI</i>	G ANTC			399
	<i>CviCI</i>	GANTC			399
	<i>CviDI</i>	GANTC			399
	<i>CviEI</i>	GANTC			399
	<i>CviFI</i>	GANTC			399
	<i>CviGI</i>	GANTC			399
	<i>FnuAI</i>	G ANTC			311
	<i>HhaII</i>	G ANTC	2(6)		400–402
	<i>NcaI</i>	GANTC			403
	<i>NovI</i>	GANTC			403
	<i>NsiHI</i>	GANTC			404
<i>HpaI</i>		GTT AAC	5(6)	ABEGIKMNOPRSUVX	405–407
	<i>BseII</i>	GTTAAC			345
	<i>SsrI</i>	GTT AAC			408
	<i>SsrB6I</i>	GTT AAC			8
<i>HpaII</i>		C CGG	2(5)	BEGMNOPRSUVX	342,405,406
	<i>Asp748I</i>	CCGG			217
	<i>BsiSI</i>	CCGG			78
	<i>Bsp5I</i>	CCGG			38
	<i>Bsp47I</i>	CCGG			79
	<i>Bsp48I</i>	CCGG			79
	<i>Bsp116I</i>	CCGG			38
	<i>Bst40I</i>	C CGG			8
	<i>Bsu1192I</i>	CCGG			74,140
	<i>BsuFI</i>	CCGG	1(5)		74,140,319

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>FinII</i>	CCGG			159
	<i>HapII</i>	C↓CGG		AGIK	338,409
	<i>Hin2I</i>	CCGG			218
	<i>Hin5I</i>	CCGG			18
	<i>MniII</i>	CCGG			288
	<i>MnoI</i>	C↓CGG			75,410
	<i>MspI</i>	C↓CGG	1(5)	ABEGIKMNOPRSUVX	154,411,412
	<i>Pde137I</i>	C↓CGG			8
	<i>SecII</i>	CCGG			413
	<i>SfaGII</i>	CCGG			414
<i>HphI</i>		GGTGA(8/7)	-2(5)	NVX	395,415,416
	<i>NgoBI</i>	GGTGA	-2(5)		417,418
<i>KpnI</i>		GGTAC↓C		ABEGIKMNOPRSUVX	419,420
	<i>Acc65I</i>	G↓GTACC			8
	<i>AhaB8I</i>	G↓GTACC			8
	<i>Asp718I</i>	G↓GTACC		M	421
	<i>BspJ106I</i>	GGTACC			107
	<i>Eco149I</i>	GGTACC			60
	<i>Esp19I</i>	GGTACC			55
	<i>KpnK14I</i>	GGTACC			217
	<i>NmiI</i>	GGTACC			422
	<i>Sau10I</i>	GGTACC			203
	<i>StiI</i>	G↓GTACC			423,424
	<i>StiAI</i>	GGTACC			122
	<i>StiBI</i>	GGTACC			122
	<i>StiCI</i>	GGTACC			122
	<i>StiDI</i>	GGTACC			122
	<i>StiEI</i>	GGTACC			122
	<i>StiFI</i>	GGTACC			122
	<i>StiGI</i>	GGTACC			122
	<i>StiHI</i>	GGTACC			122
	<i>StiJI</i>	GGTACC			122
	<i>StiKI</i>	GGTACC			122
	<i>StiLI</i>	GGTACC			122
	<i>StiMI</i>	GGTACC			122
	<i>StiNI</i>	GGTACC			375,424
<i>Ksp632I</i>		CTCTTC(1/4)		M	425
	<i>EarI</i>	CTCTTC(1/4)		N	426
	<i>Uba1104I</i>	CTCTTC(1/5)			427
<i>MaeI</i>		C↓TAG		M	428
	<i>MjaI</i>	CTAG			62
<i>MaeII</i>		A↓CGT		M	428
<i>MaeIII</i>		↓GTNAC		M	428
<i>MboI</i> ⁷		↓GATC		BEGIKNOPRSVX	429
+	<i>Bce243I</i>	↓GATC			430
+	<i>Bme12I</i>	GATC			431
	<i>BsaPI</i>	GATC			165
	<i>Bsp9I</i>	GATC			38
	<i>Bsp18I</i>	GATC			38
+	<i>Bsp49I</i>	GATC			79
+	<i>Bsp51I</i>	GATC			79
+	<i>Bsp52I</i>	GATC			79
+	<i>Bsp54I</i>	GATC			79
+	<i>Bsp57I</i>	GATC			79
+	<i>Bsp58I</i>	GATC			79
+	<i>Bsp59I</i>	GATC			79
+	<i>Bsp60I</i>	GATC			79
+	<i>Bsp61I</i>	GATC			79
+	<i>Bsp64I</i>	GATC			343
+	<i>Bsp65I</i>	GATC			79
+	<i>Bsp66I</i>	GATC			79
+	<i>Bsp67I</i>	↓GATC			343
+	<i>Bsp72I</i>	GATC			79
	<i>Bsp74I</i>	GATC			343

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Bsp76I</i>	GATC			343
+	<i>Bsp91I</i>	GATC			79
	<i>Bsp105I</i>	↓GATC			343
+	<i>Bsp122I</i>	GATC			79
+	<i>Bsp2095I</i>	↓GATC			8
+	<i>BspAI</i>	↓GATC			49
+	<i>BspJ64I</i>	GATC			79
	<i>BsrPII</i>	GATC			165
	<i>BssGII</i>	GATC			165
	<i>BstEIII</i>	GATC			75,198,432
	<i>BstXII</i>	GATC			165
+	<i>Btcl</i>	GATC			433
	<i>CacI</i>	↓GATC			434
+	<i>CcyI</i>	↓GATC			435
	<i>Cin1467I</i>	GATC			87
	<i>CpaI</i>	GATC			349
+	<i>CpfI</i>	↓GATC			251
+	<i>Csp5I</i>	GATC			79
	<i>Cte1179I</i>	GATC			87
	<i>Cte1180</i>	GATC			87
	<i>CryI</i>	GATC			320
	<i>CviAI</i>	↓GATC	2(6)		436,437
	<i>CviHI</i>	GATC			240
	<i>DpnII</i>	GATC			244,246
	<i>FnuAII</i>	GATC			75,311
	<i>FnuCI</i>	↓GATC			311
+	<i>FnuEI</i>	↓GATC			311
	<i>HaeI</i>	↓GATC			438
+	<i>Kzo9I</i>	↓GATC			104
	<i>MeuI</i>	GATC			30
	<i>MkrAI</i>	GATC			233
	<i>MmeII</i>	GATC			439
	<i>MnoIII</i>	GATC			75
	<i>MosI</i>	GATC			429
	<i>Msp67II</i>	GATC			46
+	<i>MspBI</i>	GATC			117
	<i>MthI</i>	GATC			165
	<i>Mth1047I</i>	GATC			87
	<i>MthAI</i>	GATC			117
	<i>NdeII</i>	↓GATC		BGM	403
	<i>NfiI</i>	GATC			403
	<i>NfiAII</i>	GATC			303
	<i>NfiBI</i>	GATC			440
	<i>NlaII</i>	↓GATC			357
	<i>NlaDI</i>	GATC			64
	<i>NmeCI</i>	↓GATC			304
	<i>NphI</i>	↓GATC			304
	<i>NsiAI</i>	GATC			441
	<i>NspAI</i>	GATC			30
	<i>NsuI</i>	GATC			65
+	<i>Pei9403I</i>	GATC			142
	<i>Pfal</i>	GATC			411
	<i>Rlu1I</i>	GATC			203
	<i>SalAI</i>	GATC			265
	<i>SalHI</i>	GATC			265
+	<i>Sau3AI</i>	↓GATC	4(5)	ABEGIKMNOPRSUVX	442,443
	<i>Sau6782I</i>	GATC			444
+	<i>SauCI</i>	GATC			445
+	<i>SauDI</i>	GATC			445
+	<i>SauEI</i>	GATC			445
+	<i>SauFI</i>	GATC			445
+	<i>SauGI</i>	GATC			445
+	<i>SauMI</i>	GATC			233
	<i>SinMI</i>	GATC			65

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>TruII</i>	GATC			123
<i>MboII</i>		GAAGA(8/7)	5(6)	BGIKNPRSUVX	429,446–448
	<i>NcuI</i>	GAAGA			65
	<i>TceI</i>	GAAGA			363
<i>McrI</i>		CIGRYCG			449
<i>MfeI</i>		CAATTG			450
<i>MluI</i>		AICGCGT		ABEGIKMNOPRUVX	324
	<i>ApeI</i>	ACGCGT			265
	<i>Uba6I</i>	ACGCGT			55
<i>MlyI</i>		GASTC			42
<i>MmeI</i>		TCCRAC(20/18)			439
<i>MnII</i>		CCTC(7/7)		EGNX	154,451
<i>MseI</i>		TITAA		N	452
	<i>Tru9I</i>	TITAA			8
<i>MstI</i>		TGCIGCA		X	153,453
	<i>AcaIII</i>	TGCGCA			77
	<i>AosI</i>	TGCIGCA		G	11
	<i>AvII</i>	TGCIGCA		M	70
	<i>CleII</i>	TGCGCA			77
	<i>CluII</i>	TGCGCA			68
	<i>FdiII</i>	TGCIGCA		U	115,116
	<i>FspI</i>	TGCIGCA		EGNS	83,454
	<i>GspAII</i>	TGCGCA			78
	<i>NspHIII</i>	TGCGCA			70
	<i>NspLI</i>	TGCGCA			68,69
	<i>NspMI</i>	TGCGCA			69,77
<i>MwoI</i>		GCNNNNNINNGC			455
<i>NaeI</i>		GCCIGGC		EGKMNOUVX	456
	<i>AmeII</i>	GCCGGC			32
	<i>AniMI</i>	GCCGGC			159
	<i>ApeAI</i>	GCCGGC			32
	<i>AprI</i>	GCCGGC			30
	<i>Eco56I</i>	GICCGGC			51,457
	<i>Esp5I</i>	GCCGGC			203
	<i>MisI</i>	GCCGGC			198
	<i>NasWI</i>	GCCGGC			30
	<i>NbaI</i>	GCCGGC			74
	<i>NbrI</i>	GCCGGC			74
	<i>NgoAIV</i>	GICCGGC			458
	<i>NgoMI</i>	GCCGGC			459
	<i>NmuI</i>	GCCGGC			403
	<i>NmuFI</i>	GCCGGC			65
	<i>NspWI</i>	GCCGGC			30
	<i>NtaSII</i>	GCCGGC			119
	<i>PglI</i>	GCCGGC			460
	<i>Psp61I</i>	GCCGGC			46
	<i>RluI</i>	GCCGGC			461–463
	<i>SacAI</i>	GCCGGC			57
	<i>SalCI</i>	GCCGGC			32
	<i>SaoI</i>	GCCGGC			464
	<i>SauAI</i>	GCCGGC			440
	<i>SauBMKI</i>	GCCIGGC			465
	<i>SkaI</i>	GCCGGC			65
	<i>Slu177II</i>	GCCIGGC			87
<i>NarI</i>		GGICGCC		BEGMNOPVX	466
	<i>BbeI</i>	GGCGCIC		AK	467
	<i>BbeAI</i>	GGCGCC			179
	<i>BinSII</i>	GGCGCC			179
	<i>Eco78I</i>	GGCIGCC			468
	<i>EheI</i>	GGCIGCC		OU	469,470
	<i>MchI</i>	GGICGCC			471
	<i>Mly113I</i>	GGICGCC			8
	<i>NamI</i>	GGCGCC			472
	<i>NdaI</i>	GGICGCC			473

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>NunII</i>	GG↓CGCC		G	165
	<i>SfoI</i>	GGCGCC			159
<i>NcoI</i>		C↓CATGG		ABEGIKMNOPRUVX	165
	<i>Bsp19I</i>	C↓CATGG			8
	<i>NspSAIII</i>	CCATGG			97
<i>NdeI</i>		CA↓TATG	4(6)	BEGMNPSVX	474-476
<i>NheI</i>		G↓CTAGC		BEGMNOPRUVX	422
<i>NlaIII</i>		CATG↓		N	357
	<i>Hin1II</i>	CATG↓			477
	<i>Hin8II</i>	CATG			18
<i>NlaIV</i>		GGN↓NCC		N	357
	<i>AspNI</i>	GGN↓NCC			181
	<i>BcrI</i>	GGNNCC			262
	<i>Bsp29I</i>	GGNNCC			38
	<i>BssI</i>	GGNNCC			81
	<i>Rlu3I</i>	GGNNCC			55
<i>NotI</i>		GC↓GGCCGC		ABEGIKMNOPRSUVX	478,479
<i>NruI</i>		TCG↓CGA		ABEGIKMNOPUV	422
	<i>AmaI</i>	TCGCGA			149
	<i>Bsp68I</i>	TCGCGA			79
	<i>MluB2I</i>	TCG↓CGA			8
	<i>SalDI</i>	TCGCGA			32
	<i>Sbo13I</i>	TCG↓CGA			126
	<i>Sna3286I</i>	TCGCGA			87
	<i>SpoI</i>	TCG↓CGA		R	416
<i>NspI</i>		RCATG↓Y		AKMU	67
	<i>NspHI</i>	RCATG↓Y		G	85
<i>NspBII</i>		CMG↓CKG		G	85
	<i>MspAII</i>	CMG↓CKG			8
<i>PflMI</i>		CCANNNN↓NTGG		N	159
	<i>AccB7I</i>	CCANNNN↓NTGG			8
	<i>Van91I</i>	CCANNNN↓NTGG			128
<i>PleI</i>		GAGTC(4/5)		N	480
<i>PmaCI</i>		CAC↓GTG		K	481
	<i>BbrPI</i>	CAC↓GTG		M	482
	<i>BcoAI</i>	CAC↓GTG			483
	<i>Bsp87I</i>	CACGTG			79
	<i>Eco72I</i>	CAC↓GTG			484
	<i>PmlI</i>	CAC↓GTG		N	485
<i>PpuMI</i>		RG↓GWCCY		N	159,486
	<i>Uba1I</i>	RG↓GWCCY			22
	<i>Uba1106I</i>	RGGWCCY			128
<i>PshAI</i>		GACNN↓NNGTC			487
<i>PstI</i>		CTGCA↓G	5(6)	ABEGIKMNOPRSUVX	419,488,489
	<i>AinI</i>	CTGCAG			77
	<i>Ali2882I</i>	CTGCAG			139
	<i>AliAJI</i>	CTGCA↓G			490
	<i>Asp36I</i>	CTGCAG			184
	<i>Asp708I</i>	CTGCAG			100
	<i>Asp713I</i>	CTGCA↓G			14
	<i>AspTI</i>	CTGCAG			77
	<i>BbiI</i>	CTGCAG			13
	<i>Bce170I</i>	CTGCAG			140
	<i>Bsp17I</i>	CTGCAG			38
	<i>Bsp43I</i>	CTGCAG			55
	<i>Bsp63I</i>	CTGCA↓G			343
	<i>Bsp78I</i>	CTGCAG			343
	<i>Bsp81I</i>	CTGCAG			79
	<i>Bsp93I</i>	CTGCAG			79
	<i>Bsp107I</i>	CTGCAG			38
	<i>Bsp108I</i>	CTGCAG			38
	<i>BspBI</i>	CTGCA↓G			49
	<i>BsuBI</i>	CTGCAG			140,491
	<i>CauIII</i>	CTGCAG			492

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>CfiI</i>	CTGCA↓G			438
	<i>CfrA4I</i>	CTGCA↓G			493
	<i>CfuII</i>	CTGCA↓G			494
	<i>CleI</i>	CTGCAG			77
	<i>CstI</i>	CTGCA↓G			495
	<i>EaePI</i>	CTGCAG			313
	<i>Ecl77I</i>	CTGCAG			261
	<i>Ecl133I</i>	CTGCAG			60
	<i>Ecl593I</i>	CTGCAG			217
	<i>Eco48I</i>	CTGCAG			150
	<i>Eco49I</i>	CTGCAG			150
	<i>Eco83I</i>	CTGCAG			261
	<i>Eco161I</i>	CTGCAG			60
	<i>Eco167I</i>	CTGCAG			60
	<i>Esp5II</i>	CTGCAG			203
	<i>Esp141I</i>	CTGCAG			216
	<i>GseII</i>	CTGCAG			61
	<i>HalB6II</i>	CTGCA↓G			8
	<i>Kpn12I</i>	CTGCAG			60
	<i>MauI</i>	CTGCAG			148
	<i>MkrI</i>	CTGCAG			30
	<i>NasI</i>	CTGCAG			119
	<i>NgbI</i>	CTGCAG			30
	<i>NocI</i>	CTGCAG			422
	<i>PmaI</i>	CTGCAG			154
	<i>Pma44I</i>	CTGCAG			496
	<i>PmyI</i>	CTGCAG			497
	<i>SalPI</i>	CTGCA↓G			498,499
	<i>SfiI</i>	CTGCA↓G			288
	<i>SkaII</i>	CTGCAG			65
	<i>XmaII</i>	CTGCAG			500
	<i>XorI</i>	CTGCAG			501
	<i>XphI</i>	CTGCAG			502
	<i>YenI</i>	CTGCA↓G			503
	<i>YenAI</i>	CTGCAG			503
	<i>YenBI</i>	CTGCAG			503
	<i>YenCI</i>	CTGCAG			503
	<i>YenDI</i>	CTGCAG			503
	<i>YenEI</i>	CTGCAG			503
<i>PvuI</i>		CGAT↓CG		ABEGKMNOPRSUVX	504
	<i>BmaI</i>	CGATCG			32,505,506
	<i>BmaAI</i>	CGATCG			33
	<i>BmaBI</i>	CGATCG			32
	<i>BmaCI</i>	CGATCG			32
	<i>BmaDI</i>	CGATCG			32
	<i>Cfr51I</i>	CGATCG			38
	<i>DrdIII</i>	CGATCG			257
	<i>EclII</i>	CGATCG			507
	<i>ErhB9I</i>	CGAT↓CG			8
	<i>NbII</i>	CGAT↓CG			154
	<i>Ple19I</i>	CGAT↓CG			8
	<i>Psu161I</i>	CGAT↓CG			8
	<i>RshI</i>	CGAT↓CG			508
	<i>RspI</i>	CGATCG			509
	<i>SmaAIII</i>	CGATCG			20
	<i>SplAIII</i>	CGATCG			20
	<i>Xgl3216I</i>	CGATCG			87
	<i>Xgl3217I</i>	CGATCG			87
	<i>Xgl3218I</i>	CGATCG			87
	<i>Xgl3219I</i>	CGATCG			87
	<i>Xgl3220I</i>	CGATCG			87
	<i>XmlI</i>	CGATCG			510
	<i>XmlAI</i>	CGATCG			510
	<i>XnuI</i>	CGATCG			354
	<i>XorII</i>	CGAT↓CG		B	501,504

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
<i>PvuII</i>		CAG CTG	4(4)	ABEGIKMNOPRSUVX	504,511
	<i>BavI</i>	CAG CTG			512
	<i>BavAI</i>	CAG CTG			513
	<i>Cfr6I</i>	CAG CTG	4(4)		50,51,511
	<i>EclI</i>	CAG CTG			8,285
	<i>GspI</i>	CAGCTG			77
	<i>MziI</i>	CAGCTG			514
	<i>NmeRI</i>	CAGCTG			233
	<i>PvuHKUI</i>	CAGCTG			142
	<i>SbaI</i>	CAGCTG			375,424
	<i>SciAI</i>	CAGCTG			77
	<i>SmaAIV</i>	CAGCTG			20
	<i>Sol3335I</i>	CAGCTG			142
	<i>SplAIV</i>	CAGCTG			20
<i>RleAI</i>		CCCACA(12/9)			515
<i>RsaI</i>		GT AC		ABEGIMNOPRSUVX	516
	<i>AfaI</i>	GT AC			517
	<i>Csp6I</i>	G TAC			150
	<i>CviQI</i>	G TAC			518
<i>RsrII</i>		CG GWCCG		BEGNX	519
	<i>CpoI</i>	CGGWCCG			32
	<i>CspI</i>	CGGWCCG		R	520
<i>SacI</i>		GAGCT C		AEGIKMNOPRSUVX	521
	<i>Ecl136II</i>	GAG CTC			22
	<i>Ecl137I</i>	GAGCTC			216
	<i>EcoICRI</i>	GAGCTC			148
	<i>NasSI</i>	GAGCTC			119
	<i>ScoI</i>	GAGCTC			464
	<i>SsrI</i>	GAGCT C		B	522,523
<i>SacII</i>		CCGC GG		EINOPRUVX	521
	<i>AosIII</i>	CCGCGG			70
	<i>BacI</i>	CCGCGG			75,353
	<i>Bac465I</i>	CCGCGG			87
	<i>Bsp12I</i>	CCGCGG			38
	<i>Cfr37I</i>	CCGCGG			53
	<i>Cfr41I</i>	CCGCGG			53
	<i>Cfr42I</i>	CCGC GG			17
	<i>Cfr43I</i>	CCGCGG			54
	<i>Cfr45II</i>	CCGCGG			54
	<i>CseI</i>	CCGC GG			524
	<i>DrdAI</i>	CCGCGG			257
	<i>DrdBI</i>	CCGCGG			257
	<i>DrdCI</i>	CCGCGG			257
	<i>DrdEI</i>	CCGCGG			525
	<i>DrdFI</i>	CCGCGG			257
	<i>Dsp1I</i>	CCGCGG			55
	<i>Eae46I</i>	CCGC GG			8
	<i>EccI</i>	CCGCGG			75,526
	<i>Ecl28I</i>	CCGCGG			184
	<i>Ecl37I</i>	CCGCGG			58
	<i>Eco55I</i>	CCGCGG			150
	<i>Eco92I</i>	CCGCGG			261
	<i>Eco96I</i>	CCGCGG			261
	<i>Eco99I</i>	CCGCGG			261
	<i>Eco100I</i>	CCGCGG			216
	<i>Eco104I</i>	CCGCGG			286
	<i>Eco134I</i>	CCGCGG			60
	<i>Eco135I</i>	CCGCGG			60
	<i>Eco158I</i>	CCGCGG			216
	<i>Eco182I</i>	CCGCGG			60
	<i>Eco196I</i>	CCGCGG			60
	<i>Eco208I</i>	CCGCGG			38
	<i>FseI</i>	CCGCGG			527
	<i>Gall</i>	CCGC GG			438
	<i>GceI</i>	CCGC GG			438

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Gce</i> GLI	CCGC1GG			528
	<i>Ksp</i> I	CCGC1GG		EM	529
	<i>Mra</i> I	CCGCGG		G	530
	<i>Ngo</i> III	CCGCGG			531
	<i>Ngo</i> AIII	CCGC1GG			532
	<i>Ngo</i> DI	CCGCGG			250
	<i>Ngo</i> PIII	CCGC1GG			304
	<i>Nla</i> DIII	CCGCGG			64
	<i>Nla</i> SI	CCGCGG			19
	<i>Pae</i> AI	CCGC1GG			113,493
	<i>Saa</i> I	CCGCGG			159
	<i>Sab</i> I	CCGCGG			464
	<i>Sak</i> I	CCGCGG			57
	<i>Sbo</i> I	CCGCGG			533,534
	<i>Sfr</i> I	CCGCGG			533,534
	<i>Sfr</i> 303I	CCGC1GG			8
	<i>Sfr</i> 382I	CCGCGG			87
	<i>Shy</i> I	CCGCGG			535
	<i>Sse</i> II	CCGCGG			61
	<i>Ssp</i> 1725I	CCGCGG			87
	<i>Sst</i> II	CCGC1GG		B	522
	<i>Tgi</i> I	CCGCGG			152
<i>Sal</i> I		G1TCGAC		ABEGIKMNOPRSUVX	536
	<i>Hgi</i> CIII	G1TCGAC			15
	<i>Hgi</i> DII	G1TCGAC			15
	<i>Koy</i> I	GTCGAC			32
	<i>Nop</i> I	G1TCGAC			154
	<i>Rfi</i> FI	GTCGAC			537
	<i>Rhe</i> I	GTCGAC			149
	<i>Rhp</i> I	GTCGAC			149
	<i>Rrh</i> I	GTCGAC			74
	<i>Rro</i> I	GTCGAC			74
	<i>Xam</i> I	GTCGAC			536
	<i>Xci</i> I	G1TCGAC			538
<i>Sau</i> I		CC1TNAGG		M	539
	<i>Aoc</i> I	CC1TNAGG		E	70
	<i>Ary</i> I	CC1TNAGG		GV	540
	<i>Bse</i> 21I	CC1TNAGG			541
	<i>Bsu</i> 36I	CC1TNAGG		N	542
	<i>Cvn</i> I	CC1TNAGG		B	543,544
	<i>Eci</i> CI	CCTNAGG			39
	<i>Eco</i> 76I	CCTNAGG			261
	<i>Eco</i> 81I	CC1TNAGG		AKOU	545
	<i>Eco</i> 115I	CCTNAGG			216
	<i>Eco</i> 118I	CCTNAGG			216
	<i>Lmu</i> 60I	CC1TNAGG			8
	<i>Mst</i> II	CC1TNAGG		EX	154
	<i>Oxa</i> NI	CC1TNAGG			155
	<i>Sec</i> III	CCTNAGG			413
	<i>Ssh</i> AI	CC1TNAGG			546
<i>Sca</i> I		AGT1ACT		ABEGIKMNOPRSUVX	547,548
	<i>Asp</i> 763I	AGTACT			217
	<i>Bst</i> MI	AGTACT			80
<i>Scr</i> FI		CC1NGG		EGMNOSVX	549
	<i>Bsp</i> 53I	CCNGG			79
	<i>Bsp</i> 73I	CCNGG			79
	<i>Dsa</i> V	1CCNGG		M	7
	<i>Eco</i> 43I	CCNGG			550
	<i>Eco</i> 51II	CCNGG			150
	<i>Eco</i> 80I	CCNGG			261
	<i>Eco</i> 85I	CCNGG			550
	<i>Eco</i> 93I	CCNGG			261
	<i>Eco</i> 153I	CCNGG			216
	<i>Eco</i> 200I	CCNGG			60

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Msp67I</i>	CC↓NGG			46
	<i>SsoII</i>	↓CCNGG	2(5)		272,551
<i>SduI</i>		GDGCH↓C			552,553
	<i>AocII</i>	GDGCH↓C			70
	<i>BkaI</i> 125I	GDGCHC			87
	<i>BmyI</i>	GDGCH↓C		M	554
	<i>Bsp</i> 1286I	GDGCH↓C		EGKNRUX	75,140,555
	<i>NspII</i>	GDGCH↓C			67
<i>SecI</i>		C↓CNNGG			413
	<i>BsaII</i>	C↓CNNGG		N	80
<i>SfaNI</i>		GCATC(5/9)		NX	163
	<i>BscAI</i>	GCATC			556
<i>SfeI</i>		C↓TRYAG			557
	<i>BdiSI</i>	C↓TRYAG			558
<i>SfiI</i>		GGCCNNNN↓NGGCC		BEGINOPRSUVX	559
<i>SgrAI</i>		CR↓CCGGYG			560
<i>SmaI</i>		CCC↓GGG	3(5)	ABEGIKMNOPRSUVX	500,561,562
	<i>AhyI</i>	C↓CCGGG			485
	<i>Cfr9I</i>	C↓CCGGG	2(4)	O	50,51,562
	<i>Cfr14I</i>	CCC↓GGG			233
	<i>EclRI</i>	CCC↓GGG			233
	<i>PaeBI</i>	CCC↓GGG			563
	<i>XcyI</i>	C↓CCGGG			564
	<i>XmaI</i>	C↓CCGGG		EINRVX	500
<i>SnaI</i>		GTATAC			565
	<i>UbaI</i> 107I	GTA↓TAC			427
	<i>XcaI</i>	GTA↓TAC			566
<i>SnaBI</i>		TAC↓GTA		EGMNVX	567
	<i>EciAI</i>	TACGTA			39
	<i>Eco</i> 105I	TAC↓GTA		OU	22
	<i>Eco</i> 158II	TACGTA			216
	<i>Ssp</i> JI	TACGTA			20
	<i>Ssp</i> M1I	TACGTA			20
	<i>Ssp</i> M2I	TACGTA			20
<i>SpeI</i>		A↓CTAGT		BEGKMNORSUX	422
<i>SphI</i>		GCATG↓C		ABEGIKMNOPRSUVX	568
	<i>BbuI</i>	GCATG↓C		R	569
	<i>Bsp</i> 121I	GCATGC			38
	<i>PaeI</i>	GCATG↓C			570
	<i>SpaXI</i>	GCATGC			148
<i>SpI</i>		CCGC			32
<i>SpII</i>		C↓GTACG		AK	361
	<i>PfuI</i>	CGTACG			32
	<i>SmaAI</i>	CGTACG			20
	<i>SplAI</i>	CGTACG			20
	<i>SspKI</i>	CGTACG			20
<i>SspI</i>		AAT↓ATT		BEGMNORVX	548
<i>StuI</i>		AGG↓CCT		ABEGIKMNPRVX	571
	<i>AatI</i>	AGG↓CCT		OU	5
	<i>Asp</i> 78I	AGGCCT			217
	<i>ChyI</i>	AGGCCT			33
	<i>Eco</i> 147I	AGG↓CCT			38,572
	<i>GdiI</i>	AGG↓CCT			332
	<i>NtaSI</i>	AGGCCT			119
	<i>Pme</i> 55I	AGG↓CCT			8
	<i>SarI</i>	AGGCCT			556
	<i>SsvI</i>	AGGCCT			573
	<i>Stel</i>	AGGCCT			573
<i>StyI</i>		C↓CWWGG		BEGMNRVX	574
	<i>Bss</i> T1I	C↓CWWGG			8
	<i>Eco</i> 130I	C↓CWWGG		U	38,575
	<i>Eco</i> 208II	CCWWGG			38
	<i>Eco</i> T14I	C↓CWWGG		AK	576
	<i>Eco</i> T104I	CCWWGG			576

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>ErhB9II</i>	C1CWWGG			8
	<i>SblAI</i>	CCWWGG			375
	<i>SblBI</i>	CCWWGG			375
	<i>SblCI</i>	CCWWGG			375
<i>TaqI</i>		T1CGA	4(6)	BEGIMNOPRSUVX	577,578
	<i>TfiI</i>	TCGA			578
	<i>TthHB8I</i>	T1CGA	4(6)	AK	578–581
<i>TaqII</i> ⁸		GACCGA(11/9)			75,582
		CACCCA(11/9)			
<i>TfiI</i>		GAWTC			583
<i>Tsp45I</i>		GTSAC			584
<i>TspEI</i>		AATT			585
<i>Tth1111I</i>		GACN1NNGTC		AEGIKNPVX	586
	<i>AspI</i>	GACN1NNGTC		M	587
	<i>FsuI</i>	GACNNNGTC			464
	<i>NatI</i>	GACNNNGTC			119
	<i>SmaAII</i>	GACNNNGTC			20
	<i>SpII</i>	GACNNNGTC			361
	<i>SpIAII</i>	GACNNNGTC			20
	<i>TspI</i>	GACNNNGTC			81
	<i>TieI</i>	GACNNNGTC			586
	<i>TirI</i>	GACNNNGTC			586
<i>Tth1111II</i>		CAARCA(11/9)			588
<i>Uba1105I</i>		GACNNN1NNGTC			589
<i>Uba1108I</i>		TCGTAG			590
<i>VspI</i>		AT1TAAT			591
	<i>AseI</i>	AT1TAAT		N	209
	<i>AsnI</i>	AT1TAAT		M	592
<i>XbaI</i>		T1CTAGA		ABEGIKMNOPRSUVX	593
<i>XcmI</i>		CCANNNNN1NNNTGG		N	32
<i>XhoI</i>		C1TCGAG		ABEGIKMNOPRSUVX	594
	<i>AbrI</i>	C1TCGAG			595
	<i>Asp47I</i>	CTCGAG			217
	<i>Asp703I</i>	CTCGAG			100
	<i>BadI</i>	CTCGAG			191
	<i>BbfI</i>	CTCGAG			191
	<i>BbiIII</i>	CTCGAG			13
	<i>BluI</i>	C1TCGAG			594
	<i>Bsp92I</i>	CTCGAG			79
	<i>BssHI</i>	CTCGAG			165
	<i>BstHI</i>	CTCGAG			165
	<i>BstLI</i>	CTCGAG			80
	<i>BstVI</i>	C1TCGAG	5(6)	G	596–598
	<i>BsuMI</i>	CTCGAG	3(5)		140,319
	<i>BthI</i>	CTCGAG			179
	<i>CcrI</i>	C1TCGAG		X	599
	<i>CjaI</i>	CTCGAG			471
	<i>DdeII</i>	CTCGAG			75
	<i>DrdDI</i>	CTCGAG			525
	<i>MavI</i>	C1TCGAG			600
	<i>McaI</i>	CTCGAG			100
	<i>MecI</i>	CTCGAG			514
	<i>MlaAI</i>	CT1CGAG			471
	<i>MpuI</i>	CTCGAG			514
	<i>MrhI</i>	CTCGAG			471
	<i>MsiI</i>	CTCGAG			75,395
	<i>OcoI</i>	CTCGAG			32
	<i>PaeR7I</i>	C1TCGAG	5(6)	NX	601,602
	<i>PanI</i>	C1TCGAG			148
	<i>PfiNI</i>	CTCGAG			33
	<i>PfiWI</i>	CTCGAG			603
	<i>Sal1974I</i>	CTCGAG			87
	<i>Sau3239I</i>	C1TCGAG			604,605
	<i>Sca1827I</i>	CTCGAG			87

Enzyme ¹	Isoschizomers	Recognition ² Sequence	Me ³ site	Commercial ⁴ source	Reference
	<i>Scil</i>	CTCIGAG			46
	<i>Scil</i> 183II	CTCGAG			87
	<i>Scul</i>	CTCGAG			533
	<i>SerI</i>	CTCGAG			533
	<i>Sff</i> 274I	CITCGAG			8
	<i>Sfu</i> 1762I	CTCGAG			87
	<i>Sgal</i>	CTCGAG			533
	<i>Sgol</i>	CTCGAG			533
	<i>Sgr</i> 184II	CTCGAG			87
	<i>Shy</i> 1766I	CTCGAG			87
	<i>SlaI</i>	CITCGAG			606
	<i>SluI</i>	CTCGAG			534
	<i>SpaI</i>	CTCGAG			533
	<i>Sph</i> 1719I	CTCGAG			87
	<i>Ssp</i> 4I	CTCGAG			55
	<i>Sve</i> 194I	CTCGAG			87
	<i>XpaI</i>	CITCGAG			594
<i>Xho</i> II		RIGATCY		EGMVX	152,321,607
	<i>Aii</i> II	RGATCY			262
	<i>Air</i> AI	RGATCY			32
	<i>Bst</i> YI	RIGATCY		N	608
	<i>Dsa</i> III	RIGATCY			7
	<i>Mfi</i> I	RIGATCY		AK	609
	<i>Tru</i> 201I	RIGATCY			8
<i>Xma</i> III		CIGGCCG	4(5)	B	37,610
	<i>Aa</i> I	CIGGCCG			611
	<i>Bst</i> ZI	CGGCCG			80
	<i>Eag</i> I	CIGGCCG		N	612
	<i>Ecl</i> XI	CIGGCCG		M	613
	<i>Eco</i> 52I	CIGGCCG		AEKOU	51,114
<i>Xmn</i> I		GAANNINNTTC		EGNX	74,614
	<i>Asp</i> 700I	GAANNINNTTC		M	100
	<i>Bbv</i> AI	GAANNINNTTC			615

Type III enzymes

Enzyme	Isoschizomers	Recognition Sequence	Me ³ site	Reference
<i>Eco</i> PI		AGACC	3(6)	638-642
<i>Eco</i> P15I		CAGCAG		643,644
<i>Hin</i> III		CGAAT		645,646
	<i>Hin</i> I	CGAAT		647
<i>Sty</i> LI		CAGAG	4(6)	648

FOOTNOTES

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

2. Recognition sequences are given using the standard abbreviations (Eur. J. Biochem. 150: 1-5, 1985) to represent ambiguity:

R = G or A

Y = C or T

M = A or C

K = G or T

S = G or C

W = A or T

H = A or C or T

B = G or T or C

V = G or C or A

D = G or A or T

N = A or C or G or T

3. The site of methylation by the cognate methylase when known is indicated as follows. The first number shows the base within the recognition sequence that is modified. A negative number indicates the complementary strand, numbered from the 5' base of that strand. The number in parentheses indicates the specific methylation involved. (6) = N6-methyladenosine; (5) = 5-methylcytosine; (4) = N4-methylcytosine.

4. Commercial sources of restriction enzymes are abbreviated as follows:

A	Amersham (2/90)
B	BRL (7/89)
E	Stratagene (2/90)
G	Anglian Biotechnology (10/89)
I	IBI (2/90)
K	Takara (1/88)
M	Boehringer Mannheim (2/90)
N	New England Biolabs (3/90)
O	Toyobo (3/90)
P	PL-Pharmacia-LKB (4/89)
R	Promega Biotec (2/90)
S	Sigma (8/89)
U	USB (3/90)
V	Serva (2/90)
X	New York Biolabs (2/90)

5. *BcgI* cleaves on both sides of the recognition sequence: 10 bases 5' to the recognition sequence and 12 bases 3' to it on both strands. Thus the recognition site is excised in a fragment, 34 base pairs long, with 2-base 3'-extensions at each end.
6. *EcoRII* isoschizomers fall into two classes based upon their sensitivity to methylation. *EcoRII* will not cleave when the second cytosine in the recognition sequence is methylated to 5-methylcytosine whereas *MvaI* will cleave such a sequence. Isoschizomers of *EcoRII* that are like *MvaI* are indicated by +.
7. *MboI* isoschizomers fall into two classes based upon their sensitivity to methylation. *MboI* will not cleave when the recognition sequence contains 6-methyladenosine whereas *Sau3AI* will not cleave when its recognition sequence contains 5-methylcytosine. Isoschizomers of *MboI* that are like *Sau3AI* are indicated by +.
8. *TaqII* differs from other restriction enzymes in recognizing two distinct sequences: GACCGA(11/9) and CACCCA(11/9).

Alphabetical listing of Type II restriction enzymes

known recognition sequences

<i>AaI</i> (<i>XmaIII</i>)	<i>AacI</i> (<i>BamHI</i>)	<i>AaeI</i> (<i>BamHI</i>)	<i>AagI</i> (<i>Clal</i>)
<i>AatI</i> (<i>StuI</i>)	<i>AarII</i>	<i>AbrI</i> (<i>XhoI</i>)	<i>AcaI</i> (<i>AsuII</i>)
<i>AcaII</i> (<i>BamHI</i>)	<i>AcaIII</i> (<i>MstI</i>)	<i>AcaIV</i> (<i>HaeIII</i>)	<i>AccI</i>
<i>AccII</i> (<i>FnuDII</i>)	<i>AccIII</i> (<i>BspMII</i>)	<i>Acc38I</i> (<i>EcoRII</i>)	<i>Acc65I</i> (<i>KpnI</i>)
<i>AccB1I</i> (<i>HgiCI</i>)	<i>AccB7I</i> (<i>PflMI</i>)	<i>AccEBI</i> (<i>BamHI</i>)	<i>AcrI</i> (<i>AvaI</i>)
<i>AcrII</i> (<i>BstEII</i>)	<i>Acyl</i>	<i>AeuI</i> (<i>EcoRII</i>)	<i>AfaI</i> (<i>RsaI</i>)
<i>AffI</i> (<i>AvaII</i>)	<i>AfIII</i>	<i>AfIII</i>	<i>AgeI</i>
<i>AhaI</i> (<i>CauII</i>)	<i>AhaII</i> (<i>AcyI</i>)	<i>AhaIII</i>	<i>AhaB1I</i> (<i>AsuI</i>)
<i>AhaB8I</i> (<i>KpnI</i>)	<i>AhyI</i> (<i>SmaI</i>)	<i>AinI</i> (<i>PstI</i>)	<i>AinII</i> (<i>BamHI</i>)
<i>AitI</i> (<i>Eco47III</i>)	<i>AiiII</i> (<i>XhoII</i>)	<i>AitAI</i> (<i>XhoII</i>)	<i>Alil</i> (<i>BamHI</i>)
<i>Ali2882I</i> (<i>PstI</i>)	<i>Ali12257I</i> (<i>BamHI</i>)	<i>Ali12258I</i> (<i>BamHI</i>)	<i>AliAJI</i> (<i>PstI</i>)
<i>AluI</i>	<i>AlwI</i> (<i>BinI</i>)	<i>Alw21I</i> (<i>HgiAI</i>)	<i>Alw26I</i> (<i>BsmAI</i>)
<i>Alw44I</i> (<i>ApaLI</i>)	<i>AlwNI</i>	<i>AlwXI</i> (<i>BbvI</i>)	<i>Amal</i> (<i>NruI</i>)
<i>Ama87I</i> (<i>AvaI</i>)	<i>AmeI</i> (<i>ApaLI</i>)	<i>AmeII</i> (<i>NaeI</i>)	<i>AniMI</i> (<i>NaeI</i>)
<i>AocI</i> (<i>SauI</i>)	<i>AocII</i> (<i>SduI</i>)	<i>AorI</i> (<i>EcoRII</i>)	<i>AosI</i> (<i>MstI</i>)
<i>AosII</i> (<i>AcyI</i>)	<i>AosIII</i> (<i>SacII</i>)	<i>Apal</i>	<i>ApalI</i>
<i>ApeI</i> (<i>MluI</i>)	<i>ApeAI</i> (<i>NaeI</i>)	<i>AprI</i> (<i>NaeI</i>)	<i>Apul</i> (<i>AsuI</i>)
<i>Apu16I</i> (<i>Clal</i>)	<i>ApyI</i> (<i>EcoRII</i>)	<i>AquI</i> (<i>AvaI</i>)	<i>AseI</i> (<i>VspI</i>)
<i>AseII</i> (<i>CauII</i>)	<i>AsnI</i> (<i>VspI</i>)	<i>AspI</i> (<i>Tih111I</i>)	<i>Asp1I</i> (<i>CauII</i>)
<i>Asp36I</i> (<i>PstI</i>)	<i>Asp47I</i> (<i>XhoI</i>)	<i>Asp52I</i> (<i>HindIII</i>)	<i>Asp78I</i> (<i>StuI</i>)
<i>Asp697I</i> (<i>AvaII</i>)	<i>Asp700I</i> (<i>XmnI</i>)	<i>Asp703I</i> (<i>XhoI</i>)	<i>Asp707I</i> (<i>Clal</i>)
<i>Asp708I</i> (<i>PstI</i>)	<i>Asp713I</i> (<i>PstI</i>)	<i>Asp718I</i> (<i>KpnI</i>)	<i>Asp742I</i> (<i>HaeIII</i>)
<i>Asp745I</i> (<i>AvaII</i>)	<i>Asp748I</i> (<i>HpaII</i>)	<i>Asp763I</i> (<i>ScaI</i>)	<i>Asp3065I</i> (<i>HindIII</i>)
<i>AspAI</i> (<i>BstEII</i>)	<i>AspBI</i> (<i>AvaI</i>)	<i>AspBII</i> (<i>AvaII</i>)	<i>AspCI</i> (<i>AvaI</i>)
<i>AspCII</i> (<i>AvaII</i>)	<i>AspDI</i> (<i>AvaI</i>)	<i>AspDII</i> (<i>AvaII</i>)	<i>AspHI</i> (<i>HgiAI</i>)
<i>AspNI</i> (<i>NlaIV</i>)	<i>AspTI</i> (<i>PstI</i>)	<i>AspTII</i> (<i>BamHI</i>)	<i>AspTIII</i> (<i>HaeIII</i>)
<i>AsrWI</i> (<i>AcyI</i>)	<i>AsuI</i>	<i>AsuII</i>	<i>AsuIII</i> (<i>AcyI</i>)
<i>AtuII</i> (<i>EcoRII</i>)	<i>Atu1I</i> (<i>EcoRII</i>)	<i>AtuBI</i> (<i>EcoRII</i>)	<i>AtuCI</i> (<i>BclI</i>)
<i>AvaI</i>	<i>AvaII</i>	<i>AvaIII</i>	<i>Avil</i> (<i>AsuII</i>)
<i>AviII</i> (<i>MstI</i>)	<i>AvrI</i> (<i>AvaI</i>)	<i>AvrII</i>	<i>Axyl</i> (<i>SauI</i>)
<i>BacI</i> (<i>SacII</i>)	<i>Bac36I</i> (<i>AsuI</i>)	<i>Bac465I</i> (<i>SacII</i>)	<i>BadI</i> (<i>XhoI</i>)
<i>BalI</i>	<i>Bal228I</i> (<i>AsuI</i>)	<i>Bal475I</i> (<i>HaeIII</i>)	<i>Bal3006</i> (<i>HaeIII</i>)
<i>BamFI</i> (<i>BamHI</i>)	<i>BamHI</i>	<i>BamKI</i> (<i>BamHI</i>)	<i>BamNI</i> (<i>BamHI</i>)
<i>BamNxi</i> (<i>AvaII</i>)	<i>BanI</i> (<i>HgiCI</i>)	<i>BanII</i> (<i>HgiII</i>)	<i>BanIII</i> (<i>Clal</i>)
<i>BavI</i> (<i>PvuII</i>)	<i>BavAI</i> (<i>PvuII</i>)	<i>BavAII</i> (<i>AsuI</i>)	<i>Bbel</i> (<i>NarI</i>)
<i>BbeAI</i> (<i>NarI</i>)	<i>BbfI</i> (<i>XhoI</i>)	<i>Bbf7411I</i> (<i>BspMII</i>)	<i>BbII</i> (<i>PstI</i>)
<i>BbiII</i> (<i>AcyI</i>)	<i>BbiIII</i> (<i>XhoI</i>)	<i>Bbri</i> (<i>HindIII</i>)	<i>BbrPI</i> (<i>PmaCI</i>)
<i>BbsI</i> (<i>BbvII</i>)	<i>BbuI</i> (<i>SphI</i>)	<i>BbvI</i>	<i>BbvII</i>
<i>Bbv12I</i> (<i>HgiAI</i>)	<i>Bbv16I</i> (<i>BbvII</i>)	<i>BbvAI</i> (<i>XmnI</i>)	<i>BbvAII</i> (<i>Clal</i>)
<i>BbvAIII</i> (<i>BspMII</i>)	<i>Bcal</i> (<i>HhaI</i>)	<i>Bca1259I</i> (<i>BamHI</i>)	<i>Bce22I</i> (<i>AsuI</i>)
<i>Bce71I</i> (<i>HaeIII</i>)	<i>Bce170I</i> (<i>PstI</i>)	<i>Bce243I</i> (<i>MboI</i>)	<i>Bce751I</i> (<i>BamHI</i>)

<i>BceFI</i> (<i>FnuDII</i>)	<i>BceRI</i> (<i>FnuDII</i>)	<i>BceII</i>	<i>BcgI</i>
<i>BclI</i>	<i>BcmI</i> (<i>Clal</i>)	<i>Bcni</i> (<i>CauII</i>)	<i>BcoI</i> (<i>AvaI</i>)
<i>Bco33I</i> (<i>HaeIII</i>)	<i>Bco35I</i> (<i>GsuI</i>)	<i>Bco10278I</i> (<i>BamHI</i>)	<i>BcoAI</i> (<i>PmaCI</i>)
<i>BcrI</i> (<i>NlaIV</i>)	<i>BdiI</i> (<i>Clal</i>)	<i>BdiSI</i> (<i>SfeI</i>)	<i>BepI</i> (<i>FnuDII</i>)
<i>Bfi458I</i> (<i>HaeIII</i>)	<i>BfrI</i> (<i>AflI</i>)	<i>BglI</i>	<i>BglII</i>
<i>Bim19I</i> (<i>AsuII</i>)	<i>Bim19II</i> (<i>HaeIII</i>)	<i>BinI</i>	<i>BinSI</i> (<i>EcoRII</i>)
<i>BinSI</i> (<i>NarI</i>)	<i>Bka1125I</i> (<i>SduI</i>)	<i>Bla7920I</i> (<i>BspMII</i>)	<i>Blil</i> (<i>HaeIII</i>)
<i>Bli41I</i> (<i>Clal</i>)	<i>Bli49I</i> (<i>Eco31I</i>)	<i>Bli86I</i> (<i>Clal</i>)	<i>BluRI</i> (<i>Clal</i>)
<i>Blul</i> (<i>XhoI</i>)	<i>BluII</i> (<i>HaeIII</i>)	<i>Bmal</i> (<i>PvuI</i>)	<i>BmaAI</i> (<i>PvuI</i>)
<i>BmaBI</i> (<i>PvuI</i>)	<i>BmaCI</i> (<i>PvuI</i>)	<i>BmaDI</i> (<i>PvuI</i>)	<i>Bme12I</i> (<i>MboI</i>)
<i>Bme18I</i> (<i>AvaII</i>)	<i>Bme142I</i> (<i>HaeII</i>)	<i>Bme216I</i> (<i>AvaII</i>)	<i>BmyI</i> (<i>SduI</i>)
<i>BnaI</i> (<i>BamHI</i>)	<i>BpeI</i> (<i>HindIII</i>)	<i>BpuI</i> (<i>HgiIII</i>)	<i>Bpu10I</i>
<i>Bpu14I</i> (<i>AsuII</i>)	<i>Bpu95I</i> (<i>FnuDII</i>)	<i>BsaI</i> (<i>Eco31I</i>)	<i>BsaAI</i>
<i>BsaBI</i>	<i>BsaJI</i> (<i>SecI</i>)	<i>BsaPI</i> (<i>MboI</i>)	<i>Bscl</i> (<i>Clal</i>)
<i>BscAI</i> (<i>SfaNI</i>)	<i>BseI</i> (<i>HaeIII</i>)	<i>BseII</i> (<i>HpaI</i>)	<i>Bse21I</i> (<i>SauI</i>)
<i>BseAI</i> (<i>BspMII</i>)	<i>BsePI</i>	<i>BshI</i> (<i>HaeIII</i>)	<i>BshAI</i> (<i>HaeIII</i>)
<i>BshBI</i> (<i>HaeIII</i>)	<i>BshCI</i> (<i>HaeIII</i>)	<i>BshDI</i> (<i>HaeIII</i>)	<i>BshEI</i> (<i>HaeIII</i>)
<i>BshFI</i> (<i>HaeIII</i>)	<i>BshGI</i> (<i>EcoRII</i>)	<i>BshKI</i> (<i>AsuI</i>)	<i>Bsil</i>
<i>BsiAI</i> (<i>HaeIII</i>)	<i>BsiCI</i> (<i>AsuII</i>)	<i>BsiDI</i> (<i>HaeIII</i>)	<i>BsiHI</i> (<i>HaeIII</i>)
<i>BsiKI</i> (<i>BstEII</i>)	<i>BsiLI</i> (<i>EcoRII</i>)	<i>BsiMI</i> (<i>BspMII</i>)	<i>BsiOI</i> (<i>BspMII</i>)
<i>BsiQI</i> (<i>BclI</i>)	<i>BsiSI</i> (<i>HpaII</i>)	<i>Bsml</i>	<i>BsmAI</i>
<i>BsoPI</i> (<i>BsePI</i>)	<i>Bsp2I</i> (<i>Clal</i>)	<i>Bsp4I</i> (<i>Clal</i>)	<i>Bsp5I</i> (<i>HpaII</i>)
<i>Bsp6I</i> (<i>Fnu4HI</i>)	<i>Bsp6II</i> (<i>Eco57I</i>)	<i>Bsp7I</i> (<i>CauII</i>)	<i>Bsp8I</i> (<i>CauII</i>)
<i>Bsp9I</i> (<i>MboI</i>)	<i>Bsp12I</i> (<i>SacII</i>)	<i>Bsp13I</i> (<i>BspMII</i>)	<i>Bsp16I</i> (<i>EcoRV</i>)
<i>Bsp17I</i> (<i>PstI</i>)	<i>Bsp18I</i> (<i>MboI</i>)	<i>Bsp19I</i> (<i>NcoI</i>)	<i>Bsp21I</i> (<i>Cfr10I</i>)
<i>Bsp22I</i> (<i>GsuI</i>)	<i>Bsp28I</i> (<i>GsuI</i>)	<i>Bsp29I</i> (<i>NlaIV</i>)	<i>Bsp30I</i> (<i>BamHI</i>)
<i>Bsp43I</i> (<i>PstI</i>)	<i>Bsp46I</i> (<i>BamHI</i>)	<i>Bsp47I</i> (<i>HpaII</i>)	<i>Bsp48I</i> (<i>HpaII</i>)
<i>Bsp49I</i> (<i>MboI</i>)	<i>Bsp50I</i> (<i>FnuDII</i>)	<i>Bsp51I</i> (<i>MboI</i>)	<i>Bsp52I</i> (<i>MboI</i>)
<i>Bsp53I</i> (<i>ScrFI</i>)	<i>Bsp54I</i> (<i>MboI</i>)	<i>Bsp55I</i> (<i>CauII</i>)	<i>Bsp56I</i> (<i>EcoRII</i>)
<i>Bsp57I</i> (<i>MboI</i>)	<i>Bsp58I</i> (<i>MboI</i>)	<i>Bsp59I</i> (<i>MboI</i>)	<i>Bsp60I</i> (<i>MboI</i>)
<i>Bsp61I</i> (<i>MboI</i>)	<i>Bsp63I</i> (<i>PstI</i>)	<i>Bsp64I</i> (<i>MboI</i>)	<i>Bsp65I</i> (<i>MboI</i>)
<i>Bsp66I</i> (<i>MboI</i>)	<i>Bsp67I</i> (<i>MboI</i>)	<i>Bsp68I</i> (<i>NruI</i>)	<i>Bsp70I</i> (<i>FnuDII</i>)
<i>Bsp71I</i> (<i>HaeIII</i>)	<i>Bsp72I</i> (<i>MboI</i>)	<i>Bsp73I</i> (<i>ScrFI</i>)	<i>Bsp74I</i> (<i>MboI</i>)
<i>Bsp76I</i> (<i>MboI</i>)	<i>Bsp78I</i> (<i>PstI</i>)	<i>Bsp81I</i> (<i>PstI</i>)	<i>Bsp82I</i> (<i>AsuII</i>)
<i>Bsp84I</i> (<i>Clal</i>)	<i>Bsp87I</i> (<i>PmaCI</i>)	<i>Bsp91I</i> (<i>MboI</i>)	<i>Bsp92I</i> (<i>XhoI</i>)
<i>Bsp93I</i> (<i>PstI</i>)	<i>Bsp98I</i> (<i>BamHI</i>)	<i>Bsp100I</i> (<i>AvaII</i>)	<i>Bsp103I</i> (<i>EcoRII</i>)
<i>Bsp105I</i> (<i>MboI</i>)	<i>Bsp106I</i> (<i>Clal</i>)	<i>Bsp107I</i> (<i>PstI</i>)	<i>Bsp108I</i> (<i>PstI</i>)
<i>Bsp116I</i> (<i>HpaII</i>)	<i>Bsp117I</i> (<i>HgiIII</i>)	<i>Bsp119I</i> (<i>AsuII</i>)	<i>Bsp120I</i> (<i>ApaI</i>)
<i>Bsp121I</i> (<i>SphI</i>)	<i>Bsp122I</i> (<i>MboI</i>)	<i>Bsp211I</i> (<i>HaeIII</i>)	<i>Bsp226I</i> (<i>HaeIII</i>)
<i>Bsp423I</i> (<i>BbvI</i>)	<i>Bsp519I</i> (<i>HgiIII</i>)	<i>Bsp1286I</i> (<i>SduI</i>)	<i>Bsp2095I</i> (<i>MboI</i>)
<i>BspAI</i> (<i>MboI</i>)	<i>BspBI</i> (<i>PstI</i>)	<i>BspBII</i> (<i>AsuI</i>)	<i>BspBRI</i> (<i>HaeIII</i>)
<i>BspCI</i>	<i>BspDI</i> (<i>Clal</i>)	<i>BspEI</i> (<i>BspMII</i>)	<i>BspGI</i>
<i>BspHI</i>	<i>BspJ64I</i> (<i>MboI</i>)	<i>BspJ67I</i> (<i>CauII</i>)	<i>BspJ74I</i> (<i>GsuI</i>)
<i>BspJ76I</i> (<i>FnuDII</i>)	<i>BspJ105I</i> (<i>AvaII</i>)	<i>BspJ106I</i> (<i>KpnI</i>)	<i>BspMI</i>
<i>BspMII</i>	<i>BspRI</i> (<i>HaeIII</i>)	<i>BspVI</i> (<i>BbvII</i>)	<i>BspXI</i> (<i>Clal</i>)
<i>BspXII</i> (<i>BclI</i>)	<i>Bsrl</i>	<i>BsrHI</i> (<i>BsePI</i>)	<i>BsrPII</i> (<i>MboI</i>)
<i>BssI</i> (<i>NlaIV</i>)	<i>BssCI</i> (<i>HaeIII</i>)	<i>BssGI</i> (<i>BstXI</i>)	<i>BssGII</i> (<i>MboI</i>)
<i>BssHI</i> (<i>XhoI</i>)	<i>BssHII</i> (<i>BsePI</i>)	<i>BssT1I</i> (<i>Styl</i>)	<i>BstI</i> (<i>BamHI</i>)
<i>Bst31I</i> (<i>BstEII</i>)	<i>Bst40I</i> (<i>HpaII</i>)	<i>Bst1126I</i> (<i>BamHI</i>)	<i>Bst2464I</i> (<i>BamHI</i>)
<i>Bst2902I</i> (<i>BamHI</i>)	<i>BstBI</i> (<i>AsuII</i>)	<i>BstCI</i> (<i>HaeIII</i>)	<i>BstDI</i> (<i>BstEII</i>)
<i>BstEII</i>	<i>BstEIII</i> (<i>MboI</i>)	<i>BstFI</i> (<i>HindIII</i>)	<i>BstGI</i> (<i>BclI</i>)
<i>BstGII</i> (<i>EcoRII</i>)	<i>BstHI</i> (<i>XhoI</i>)	<i>BstJI</i> (<i>HaeIII</i>)	<i>BstKI</i> (<i>BclI</i>)
<i>BstLI</i> (<i>XhoI</i>)	<i>BstMI</i> (<i>ScaI</i>)	<i>BstNI</i> (<i>EcoRII</i>)	<i>BstOI</i> (<i>EcoRII</i>)
<i>BstPI</i> (<i>BstEII</i>)	<i>BstQI</i> (<i>BamHI</i>)	<i>BstRI</i> (<i>EcoRV</i>)	<i>BstSI</i> (<i>AvaI</i>)
<i>BstTI</i> (<i>BstXI</i>)	<i>BstUI</i> (<i>FnuDII</i>)	<i>BstVI</i> (<i>XhoI</i>)	<i>BstWI</i> (<i>EcoNI</i>)
<i>BstXI</i>	<i>BstXII</i> (<i>MboI</i>)	<i>BstYI</i> (<i>XhoII</i>)	<i>BstZI</i> (<i>XmaIII</i>)
<i>Bsu15I</i> (<i>Clal</i>)	<i>Bsu22I</i> (<i>BspMII</i>)	<i>Bsu36I</i> (<i>SauI</i>)	<i>Bsu90I</i> (<i>BamHI</i>)
<i>Bsu1076I</i> (<i>HaeIII</i>)	<i>Bsu1114I</i> (<i>HaeIII</i>)	<i>Bsu1192I</i> (<i>HpaII</i>)	<i>Bsu1192II</i> (<i>FnuDII</i>)
<i>Bsu1193I</i> (<i>FnuDII</i>)	<i>Bsu1532I</i> (<i>FnuDII</i>)	<i>Bsu1854I</i> (<i>HgiIII</i>)	<i>Bsu6633I</i> (<i>FnuDII</i>)
<i>Bsu8565I</i> (<i>BamHI</i>)	<i>Bsu8646I</i> (<i>BamHI</i>)	<i>BsuBI</i> (<i>PstI</i>)	<i>BsuEII</i> (<i>FnuDII</i>)
<i>BsuFI</i> (<i>HpaII</i>)	<i>BsuMI</i> (<i>XhoI</i>)	<i>BsuRI</i> (<i>HaeIII</i>)	<i>BtcI</i> (<i>MboI</i>)
<i>Btel</i> (<i>HaeIII</i>)	<i>BthI</i> (<i>XhoI</i>)	<i>BthII</i> (<i>BinI</i>)	<i>BtlI</i> (<i>AvaII</i>)
<i>Btul</i> (<i>Clal</i>)	<i>Bvul</i> (<i>HgiIII</i>)	<i>CacI</i> (<i>MboI</i>)	<i>Caul</i> (<i>AvaII</i>)
<i>CauII</i>	<i>CaulII</i> (<i>PstI</i>)	<i>CauB3I</i> (<i>BspMII</i>)	<i>CcrI</i> (<i>XhoI</i>)
<i>CcyI</i> (<i>MboI</i>)	<i>Cdi27I</i> (<i>EcoRII</i>)	<i>CelI</i> (<i>BamHI</i>)	<i>CellI</i> (<i>EspI</i>)
<i>CeqI</i> (<i>EcoRV</i>)	<i>CflI</i> (<i>PstI</i>)	<i>Cfol</i> (<i>HhaI</i>)	<i>CfrI</i>
<i>Cfr4I</i> (<i>AsuI</i>)	<i>Cfr5I</i> (<i>EcoRII</i>)	<i>Cfr6I</i> (<i>PvuII</i>)	<i>Cfr7I</i> (<i>BstEII</i>)
<i>Cfr8I</i> (<i>AsuI</i>)	<i>Cfr9I</i> (<i>SmaI</i>)	<i>Cfr10I</i>	<i>Cfr11I</i> (<i>EcoRII</i>)
<i>Cfr13I</i> (<i>AsuI</i>)	<i>Cfr14I</i> (<i>CfrI</i>)	<i>Cfr19I</i> (<i>BstEII</i>)	<i>Cfr20I</i> (<i>EcoRII</i>)

<i>Cfr22I</i> (<i>EcoRII</i>)	<i>Cfr23I</i> (<i>AsuI</i>)	<i>Cfr24I</i> (<i>EcoRII</i>)	<i>Cfr25I</i> (<i>EcoRII</i>)
<i>Cfr27I</i> (<i>EcoRII</i>)	<i>Cfr28I</i> (<i>EcoRII</i>)	<i>Cfr29I</i> (<i>EcoRII</i>)	<i>Cfr30I</i> (<i>EcoRII</i>)
<i>Cfr31I</i> (<i>EcoRII</i>)	<i>Cfr32I</i> (<i>HindIII</i>)	<i>Cfr33I</i> (<i>AsuI</i>)	<i>Cfr35I</i> (<i>EcoRII</i>)
<i>Cfr37I</i> (<i>SacII</i>)	<i>Cfr38I</i> (<i>CfrI</i>)	<i>Cfr39I</i> (<i>CfrI</i>)	<i>Cfr40I</i> (<i>CfrI</i>)
<i>Cfr41I</i> (<i>SacII</i>)	<i>Cfr42I</i> (<i>SacII</i>)	<i>Cfr43I</i> (<i>SacII</i>)	<i>Cfr45I</i> (<i>AsuI</i>)
<i>Cfr45II</i> (<i>SacII</i>)	<i>Cfr46I</i> (<i>AsuI</i>)	<i>Cfr47I</i> (<i>AsuI</i>)	<i>Cfr48I</i> (<i>HgiIII</i>)
<i>Cfr51I</i> (<i>PvuI</i>)	<i>Cfr52I</i> (<i>AsuI</i>)	<i>Cfr54I</i> (<i>AsuI</i>)	<i>CfrA4I</i> (<i>PstI</i>)
<i>CfrJ4I</i> (<i>SmaI</i>)	<i>CfrNI</i> (<i>AsuI</i>)	<i>CfrS37I</i> (<i>EcoRII</i>)	<i>CfuI</i> (<i>DpnI</i>)
<i>Cfull</i> (<i>PstI</i>)	<i>ChuI</i> (<i>HindIII</i>)	<i>ChuII</i> (<i>HindII</i>)	<i>ChyI</i> (<i>StuI</i>)
<i>Cin1467I</i> (<i>MboI</i>)	<i>CjaI</i> (<i>XhoI</i>)	<i>Clal</i>	<i>ClcI</i> (<i>PstI</i>)
<i>ClcII</i> (<i>MstI</i>)	<i>Clil</i> (<i>AvaII</i>)	<i>ClilI</i> (<i>MstI</i>)	<i>ClmI</i> (<i>HaeIII</i>)
<i>ClmII</i> (<i>AvaII</i>)	<i>Clil</i> (<i>HaeIII</i>)	<i>CpaI</i> (<i>MboI</i>)	<i>Cpa1150I</i> (<i>FnuDII</i>)
<i>CpaAI</i> (<i>FnuDII</i>)	<i>CpeI</i> (<i>BclI</i>)	<i>Cpfl</i> (<i>MboI</i>)	<i>CpoI</i> (<i>RsrII</i>)
<i>CscI</i> (<i>SacII</i>)	<i>CspI</i> (<i>RsrII</i>)	<i>Csp2I</i> (<i>HaeIII</i>)	<i>Csp4I</i> (<i>Clal</i>)
<i>Csp5I</i> (<i>MboI</i>)	<i>Csp6I</i> (<i>RsaI</i>)	<i>Csp45I</i> (<i>AsuII</i>)	<i>Csp1470I</i> (<i>HhaI</i>)
<i>CstI</i> (<i>PstI</i>)	<i>Cte1179I</i> (<i>MboI</i>)	<i>Cte1180I</i> (<i>MboI</i>)	<i>CthI</i> (<i>BclI</i>)
<i>CthII</i> (<i>EcoRII</i>)	<i>CryI</i> (<i>MboI</i>)	<i>CviAI</i> (<i>MboI</i>)	<i>CviBI</i> (<i>HinfI</i>)
<i>CviCI</i> (<i>HinfI</i>)	<i>CviDI</i> (<i>HinfI</i>)	<i>CviEI</i> (<i>HinfI</i>)	<i>CviFI</i> (<i>HinfI</i>)
<i>CviGI</i> (<i>HinfI</i>)	<i>CviHI</i> (<i>MboI</i>)	<i>CviJI</i>	<i>CviKI</i> (<i>CviII</i>)
<i>CviLI</i> (<i>CviII</i>)	<i>CviMI</i> (<i>CviII</i>)	<i>CviNI</i> (<i>CviII</i>)	<i>CviOI</i> (<i>CviII</i>)
<i>CviQI</i> (<i>RsaI</i>)	<i>Cvnl</i> (<i>SauI</i>)	<i>DdeI</i>	<i>DdeII</i> (<i>XhoI</i>)
<i>DdsI</i> (<i>BamHI</i>)	<i>DpnI</i>	<i>DpnII</i> (<i>MboI</i>)	<i>DraI</i> (<i>AhaIII</i>)
<i>DraII</i>	<i>DraIII</i>	<i>DrdI</i>	<i>DraII</i>
<i>DrdIII</i> (<i>PvuI</i>)	<i>DrdAI</i> (<i>SacII</i>)	<i>DrdBI</i> (<i>SacII</i>)	<i>DrdCI</i> (<i>SacII</i>)
<i>DrdDI</i> (<i>XhoI</i>)	<i>DrdEI</i> (<i>SacII</i>)	<i>DrdFI</i> (<i>SacII</i>)	<i>DsaI</i>
<i>DsaII</i> (<i>HaeIII</i>)	<i>DsaIII</i> (<i>XhoII</i>)	<i>DsaIV</i> (<i>AvaII</i>)	<i>DsaV</i> (<i>ScrFI</i>)
<i>DsaVI</i> (<i>AccI</i>)	<i>DspII</i> (<i>SacII</i>)	<i>EaeI</i> (<i>CfrI</i>)	<i>Eae46I</i> (<i>SacII</i>)
<i>EaePI</i> (<i>PstI</i>)	<i>EagI</i> (<i>XmaIII</i>)	<i>EagKI</i> (<i>EcoRII</i>)	<i>EagMI</i> (<i>AvaII</i>)
<i>EarI</i> (<i>Ksp632I</i>)	<i>Eal</i> (<i>BstEII</i>)	<i>EaII</i> (<i>EcoRII</i>)	<i>EccI</i> (<i>SacII</i>)
<i>EciI</i>	<i>EciAI</i> (<i>SnaBI</i>)	<i>EciBI</i> (<i>CfrI</i>)	<i>EciCI</i> (<i>SauI</i>)
<i>EciDI</i> (<i>CauII</i>)	<i>EciEI</i> (<i>ApaI</i>)	<i>EclI</i> (<i>PvuII</i>)	<i>EclII</i> (<i>EcoRII</i>)
<i>Ecl28I</i> (<i>SacII</i>)	<i>Ecl37I</i> (<i>SacII</i>)	<i>Ecl66I</i> (<i>EcoRII</i>)	<i>Ecl77I</i> (<i>PstI</i>)
<i>Ecl133I</i> (<i>PstI</i>)	<i>Ecl136I</i> (<i>EcoRII</i>)	<i>Ecl136II</i> (<i>SacI</i>)	<i>Ecl137I</i> (<i>SacI</i>)
<i>Ecl137II</i> (<i>EcoRII</i>)	<i>Ecl593I</i> (<i>PstI</i>)	<i>EclJ1</i> (<i>PvuI</i>)	<i>EclRI</i> (<i>SmaI</i>)
<i>EclS39I</i> (<i>EcoRII</i>)	<i>EclXI</i> (<i>XmaIII</i>)	<i>EcoVIII</i> (<i>HindIII</i>)	<i>Eco24I</i> (<i>HgiIII</i>)
<i>Eco25I</i> (<i>HgiIII</i>)	<i>Eco26I</i> (<i>HgiIII</i>)	<i>Eco31I</i>	<i>Eco32I</i> (<i>EcoRV</i>)
<i>Eco35I</i> (<i>HgiIII</i>)	<i>Eco38I</i> (<i>EcoRII</i>)	<i>Eco39I</i> (<i>AsuI</i>)	<i>Eco40I</i> (<i>EcoRII</i>)
<i>Eco41I</i> (<i>EcoRII</i>)	<i>Eco42I</i> (<i>Eco31I</i>)	<i>Eco43I</i> (<i>ScrFI</i>)	<i>Eco47I</i> (<i>AvaII</i>)
<i>Eco47II</i> (<i>AsuI</i>)	<i>Eco47III</i>	<i>Eco48I</i> (<i>PstI</i>)	<i>Eco49I</i> (<i>PstI</i>)
<i>Eco50I</i> (<i>HgiCI</i>)	<i>Eco51I</i> (<i>Eco31I</i>)	<i>Eco51II</i> (<i>ScrFI</i>)	<i>Eco52I</i> (<i>XmaIII</i>)
<i>Eco55I</i> (<i>SacII</i>)	<i>Eco56I</i> (<i>NaeI</i>)	<i>Eco57I</i>	<i>Eco60I</i> (<i>EcoRII</i>)
<i>Eco61I</i> (<i>EcoRII</i>)	<i>Eco64I</i> (<i>HgiCI</i>)	<i>Eco65I</i> (<i>HindIII</i>)	<i>Eco67I</i> (<i>EcoRII</i>)
<i>Eco68I</i> (<i>HgiIII</i>)	<i>Eco70I</i> (<i>EcoRII</i>)	<i>Eco71I</i> (<i>EcoRII</i>)	<i>Eco72I</i> (<i>PmaCI</i>)
<i>Eco76I</i> (<i>SauI</i>)	<i>Eco78I</i> (<i>NarI</i>)	<i>Eco80I</i> (<i>ScrFI</i>)	<i>Eco81I</i> (<i>SauI</i>)
<i>Eco82I</i> (<i>EcoRI</i>)	<i>Eco83I</i> (<i>PstI</i>)	<i>Eco85I</i> (<i>ScrFI</i>)	<i>Eco88I</i> (<i>AvaI</i>)
<i>Eco90I</i> (<i>CfrI</i>)	<i>Eco91I</i> (<i>BstEII</i>)	<i>Eco92I</i> (<i>SacII</i>)	<i>Eco93I</i> (<i>ScrFI</i>)
<i>Eco95I</i> (<i>Eco31I</i>)	<i>Eco96I</i> (<i>SacII</i>)	<i>Eco97I</i> (<i>Eco31I</i>)	<i>Eco98I</i> (<i>HindIII</i>)
<i>Eco99I</i> (<i>SacII</i>)	<i>Eco100I</i> (<i>SacII</i>)	<i>Eco101I</i> (<i>Eco31I</i>)	<i>Eco104I</i> (<i>SacII</i>)
<i>Eco105I</i> (<i>SnaBI</i>)	<i>Eco113I</i> (<i>HgiIII</i>)	<i>Eco115I</i> (<i>SauI</i>)	<i>Eco118I</i> (<i>SauI</i>)
<i>Eco120I</i> (<i>Eco31I</i>)	<i>Eco121I</i> (<i>CauII</i>)	<i>Eco125I</i> (<i>Eco57I</i>)	<i>Eco127I</i> (<i>Eco31I</i>)
<i>Eco128I</i> (<i>EcoRII</i>)	<i>Eco129I</i> (<i>Eco31I</i>)	<i>Eco130I</i> (<i>SlyI</i>)	<i>Eco134I</i> (<i>SacII</i>)
<i>Eco135I</i> (<i>SacII</i>)	<i>Eco143I</i> (<i>BsePI</i>)	<i>Eco147I</i> (<i>StuI</i>)	<i>Eco149I</i> (<i>KpnI</i>)
<i>Eco153I</i> (<i>ScrFI</i>)	<i>Eco155I</i> (<i>Eco31I</i>)	<i>Eco156I</i> (<i>Eco31I</i>)	<i>Eco157I</i> (<i>Eco31I</i>)
<i>Eco158I</i> (<i>SacII</i>)	<i>Eco158II</i> (<i>SnaBI</i>)	<i>Eco159I</i> (<i>EcoRI</i>)	<i>Eco161I</i> (<i>PstI</i>)
<i>Eco162I</i> (<i>Eco31I</i>)	<i>Eco164I</i> (<i>CfrI</i>)	<i>Eco167I</i> (<i>PstI</i>)	<i>Eco168I</i> (<i>HgiCI</i>)
<i>Eco169I</i> (<i>HgiCI</i>)	<i>Eco170I</i> (<i>EcoRII</i>)	<i>Eco171I</i> (<i>HgiCI</i>)	<i>Eco173I</i> (<i>HgiCI</i>)
<i>Eco178I</i> (<i>EcoRV</i>)	<i>Eco179I</i> (<i>CauII</i>)	<i>Eco180I</i> (<i>HgiIII</i>)	<i>Eco182I</i> (<i>SacII</i>)
<i>Eco185I</i> (<i>Eco31I</i>)	<i>Eco188I</i> (<i>HindIII</i>)	<i>Eco190I</i> (<i>CauII</i>)	<i>Eco191I</i> (<i>Eco31I</i>)
<i>Eco193I</i> (<i>EcoRII</i>)	<i>Eco195I</i> (<i>HgiCI</i>)	<i>Eco196I</i> (<i>SacII</i>)	<i>Eco196II</i> (<i>AsuI</i>)
<i>Eco200I</i> (<i>ScrFI</i>)	<i>Eco201I</i> (<i>AsuI</i>)	<i>Eco203I</i> (<i>Eco31I</i>)	<i>Eco204I</i> (<i>Eco31I</i>)
<i>Eco205I</i> (<i>Eco31I</i>)	<i>Eco206I</i> (<i>EcoRII</i>)	<i>Eco207I</i> (<i>EcoRII</i>)	<i>Eco208I</i> (<i>SacII</i>)
<i>Eco208II</i> (<i>SlyI</i>)	<i>Eco211I</i> (<i>HgiIII</i>)	<i>Eco215I</i> (<i>HgiIII</i>)	<i>Eco216I</i> (<i>HgiIII</i>)
<i>Eco217I</i> (<i>Eco31I</i>)	<i>Eco225I</i> (<i>Eco31I</i>)	<i>Eco228I</i> (<i>EcoRI</i>)	<i>Eco231I</i> (<i>HindIII</i>)
<i>Eco232I</i> (<i>HgiIII</i>)	<i>Eco233I</i> (<i>Eco31I</i>)	<i>Eco237I</i> (<i>EcoRI</i>)	<i>Eco239I</i> (<i>Eco31I</i>)
<i>Eco240I</i> (<i>Eco31I</i>)	<i>Eco241I</i> (<i>Eco31I</i>)	<i>Eco246I</i> (<i>Eco31I</i>)	<i>Eco247I</i> (<i>Eco31I</i>)
<i>Eco252I</i> (<i>EcoRI</i>)	<i>EcoA4I</i> (<i>Eco31I</i>)	<i>EcoHI</i> (<i>CfrI</i>)	<i>EcoICRI</i> (<i>SacI</i>)
<i>EcoNI</i>	<i>EcoO65I</i> (<i>BstEII</i>)	<i>EcoO109I</i> (<i>DraII</i>)	<i>EcoRI</i>
<i>EcoRII</i>	<i>EcoRV</i>	<i>EcoT14I</i> (<i>SlyI</i>)	<i>EcoT22I</i> (<i>AvaIII</i>)
<i>EcoT38I</i> (<i>HgiIII</i>)	<i>EcoT88I</i> (<i>HgiIII</i>)	<i>EcoT93I</i> (<i>HgiIII</i>)	<i>EcoT95I</i> (<i>HgiIII</i>)

<i>Eco</i> T104I (SryI)	<i>Ehe</i> I (NarI)	<i>Erh</i> B9I (PvuI)	<i>Erh</i> B9II (SryI)
<i>Erp</i> I (AvaII)	<i>Esp</i> I	<i>Esp</i> 11 (HgiCI)	<i>Esp</i> 21 (EcoRII)
<i>Esp</i> 3I	<i>Esp</i> 4I (AffII)	<i>Esp</i> 5I (NaeI)	<i>Esp</i> 5II (PstI)
<i>Esp</i> 6I (HgiCI)	<i>Esp</i> 7I (BsePI)	<i>Esp</i> 8I (BsePI)	<i>Esp</i> 9I (HgiCI)
<i>Esp</i> 10I (HgiCI)	<i>Esp</i> 11I (HgiCI)	<i>Esp</i> 12I (HgiCI)	<i>Esp</i> 13I (HgiCI)
<i>Esp</i> 14I (HgiCI)	<i>Esp</i> 15I (HgiCI)	<i>Esp</i> 19I (KpnI)	<i>Esp</i> 22I (HgiCI)
<i>Esp</i> 141I (PstI)	<i>Fau</i> I	<i>Fba</i> I (BclI)	<i>Fbl</i> I (AccI)
<i>Fbr</i> I (Fnu4HI)	<i>Fdi</i> I (AvaII)	<i>Fdi</i> II (MstI)	<i>Fin</i> I
<i>Fin</i> II (HpaII)	<i>Fin</i> SI (HaeIII)	<i>Fnu</i> 4HI	<i>Fnu</i> AI (HinfI)
<i>Fnu</i> AI (MboI)	<i>Fnu</i> CI (MboI)	<i>Fnu</i> DI (HaeIII)	<i>Fnu</i> DII
<i>Fnu</i> DIII (HhaI)	<i>Fnu</i> EI (MboI)	<i>Fok</i> I	<i>Fsc</i> I (SacII)
<i>Fse</i> I	<i>Fsf</i> I (Eco57I)	<i>Fsp</i> I (MstI)	<i>Fsp</i> II (AsuII)
<i>Fsp</i> 1604I (EcoRII)	<i>Fsp</i> MI (FnuDII)	<i>Fsp</i> MSI (AvaII)	<i>Fsu</i> I (Tth111I)
<i>Gall</i> (SacII)	<i>Gce</i> I (SacII)	<i>Gce</i> GLI (SacII)	<i>Gdl</i> I (StuI)
<i>Gdi</i> II	<i>Gdo</i> I (BamHI)	<i>Gin</i> I (BamHI)	<i>Gox</i> I (BamHI)
<i>Gse</i> I (AsuI)	<i>Gse</i> II (PstI)	<i>Gse</i> III (BamHI)	<i>Gsp</i> I (PvuII)
<i>Gsp</i> AI (AvaII)	<i>Gsp</i> AI (MstI)	<i>Gsu</i> I	<i>Hac</i> I (MboI)
<i>Hae</i> I	<i>Hae</i> II	<i>Hae</i> III	<i>Hal</i> B6I (EcoRI)
<i>Hal</i> B6II (PstI)	<i>Hap</i> II (HpaII)	<i>Hga</i> I	<i>Hgi</i> I (AcyI)
<i>Hgi</i> AI	<i>Hgi</i> BI (AvaII)	<i>Hgi</i> CI	<i>Hgi</i> CII (AvaII)
<i>Hgi</i> CIII (SalI)	<i>Hgi</i> DI (AcyI)	<i>Hgi</i> DII (SalI)	<i>Hgi</i> EI (AvaII)
<i>Hgi</i> EII	<i>Hgi</i> GI (AcyI)	<i>Hgi</i> HI (HgiCI)	<i>Hgi</i> HII (AcyI)
<i>Hgi</i> HIII (AvaII)	<i>Hgi</i> I (AvaII)	<i>Hgi</i> II	<i>Hgi</i> S21I (CauII)
<i>Hgi</i> S22I (CauII)	<i>Hha</i> I	<i>Hha</i> II (HinfI)	<i>Hhg</i> I (HaeIII)
<i>Hin</i> 1I (AcyI)	<i>Hin</i> 1II (NlaIII)	<i>Hin</i> 2I (HpaII)	<i>Hin</i> 3I (CauII)
<i>Hin</i> 5I (HpaII)	<i>Hin</i> 5II (AsuI)	<i>Hin</i> 5III (HindIII)	<i>Hin</i> 6I (HhaI)
<i>Hin</i> 7I (HhaI)	<i>Hin</i> 8I (AcyI)	<i>Hin</i> 8II (NlaIII)	<i>Hin</i> 173I (HindIII)
<i>Hin</i> 1056I (FnuDII)	<i>Hin</i> 1076III (HindIII)	<i>Hin</i> 1160II (HindII)	<i>Hin</i> 1161II (HindII)
<i>Hin</i> GUI (HhaI)	<i>Hin</i> GUII (FokI)	<i>Hin</i> HI (HaeII)	<i>Hin</i> JCI (HindII)
<i>Hin</i> JCII (HindIII)	<i>Hin</i> PII (HhaI)	<i>Hin</i> SI (HhaI)	<i>Hin</i> S2I (HhaI)
<i>Hin</i> bIII (HindIII)	<i>Hinc</i> II (HindII)	<i>Hind</i> II	<i>Hind</i> III
<i>Hinf</i> I	<i>Hinf</i> II (HindIII)	<i>Hjal</i> (EcoRV)	<i>Hpa</i> I
<i>Hpa</i> II	<i>Hph</i> I	<i>Hsp</i> 2I (AvaII)	<i>Hsu</i> I (HindIII)
<i>Isp</i> I (Fnu4HI)	<i>Kox</i> I (BstEII)	<i>Kox</i> II (HgiJII)	<i>Kox</i> 165I (EcoRII)
<i>Koy</i> I (SalI)	<i>Kpn</i> I	<i>Kpn</i> 2I (BspMII)	<i>Kpn</i> 10I (EcoRII)
<i>Kpn</i> 12I (PstI)	<i>Kpn</i> 13I (EcoRII)	<i>Kpn</i> 14I (EcoRII)	<i>Kpn</i> 16I (EcoRII)
<i>Kpn</i> 30I (BsePI)	<i>Kpn</i> K14I (KpnI)	<i>Ksp</i> I (SacII)	<i>Ksp</i> 22I (BclI)
<i>Ksp</i> 632I	<i>Kzo</i> 9I (MboI)	<i>Kzo</i> 49I (AvaII)	<i>Lnu</i> 60I (SauI)
<i>Lpl</i> I (ClaI)	<i>Lsp</i> I (AsuII)	<i>Mae</i> I	<i>Mae</i> II
<i>Mae</i> III	<i>Mam</i> I (BsaBI)	<i>Mau</i> I (PstI)	<i>Mav</i> I (XhoI)
<i>Mbo</i> I	<i>Mbo</i> II	<i>Mca</i> I (XhoI)	<i>Mch</i> I (NarI)
<i>Mcr</i> I	<i>Mec</i> I (XhoI)	<i>Meu</i> I (MboI)	<i>Mfe</i> I
<i>Mff</i> I (XhoII)	<i>Mfo</i> I (AvaII)	<i>Mis</i> I (NaeI)	<i>Mjal</i> (MaeI)
<i>Mjai</i> I (AsuI)	<i>Mki</i> I (HindIII)	<i>Mkr</i> I (PstI)	<i>Mkr</i> AI (MboI)
<i>Mla</i> I (AsuII)	<i>Mla</i> AI (XhoI)	<i>Mle</i> I (BamHI)	<i>Mli</i> I (AvaII)
<i>Mli</i> I (AluI)	<i>Mlu</i> I	<i>Mlu</i> 23I (BamHI)	<i>Mlu</i> 2300I (EcoRII)
<i>Mlu</i> B2I (NruI)	<i>Mly</i> I	<i>Mly</i> 113I (NarI)	<i>Mme</i> I
<i>Mme</i> II (MboI)	<i>Mni</i> I (HaeIII)	<i>Mni</i> II (HpaII)	<i>Mni</i> II
<i>Mnn</i> I (HindII)	<i>Mnn</i> II (HaeIII)	<i>Mnn</i> IV (HhaI)	<i>Mno</i> I (HpaII)
<i>Mno</i> III (MboI)	<i>Mos</i> I (MboI)	<i>Mph</i> I (EcoRII)	<i>Mpu</i> I (XhoI)
<i>Mra</i> I (SacII)	<i>Mrh</i> I (XhoI)	<i>Mro</i> I (BspMII)	<i>Msc</i> I (BalI)
<i>Mse</i> I	<i>Msi</i> I (XhoI)	<i>Msp</i> I (HpaII)	<i>Msp</i> 20I (BalI)
<i>Msp</i> 24I (AsuI)	<i>Msp</i> 67I (ScrFI)	<i>Msp</i> 67II (MboI)	<i>Msp</i> AI (AvaII)
<i>Msp</i> A1I (NspBII)	<i>Msp</i> BI (MboI)	<i>Msp</i> B4I (HgiCI)	<i>Msp</i> YI (BsaAI)
<i>Msi</i> I	<i>Msi</i> II (SauI)	<i>Mth</i> I (MboI)	<i>Mth</i> 1047I (MboI)
<i>Mth</i> AI (MboI)	<i>Mth</i> TI (AsuI)	<i>Mva</i> I (EcoRII)	<i>Mva</i> AI (FnuDII)
<i>Mvn</i> I (FnuDII)	<i>Mwo</i> I	<i>Mzi</i> I (PvuII)	<i>Nae</i> I
<i>Nam</i> I (NarI)	<i>Nan</i> I (EcoRV)	<i>Nan</i> II (DpnI)	<i>Nar</i> I
<i>Nas</i> I (PstI)	<i>Nas</i> BI (BamHI)	<i>Nas</i> SI (SacI)	<i>Nas</i> WI (NaeI)
<i>Nba</i> I (NaeI)	<i>Nbl</i> I (PvuI)	<i>Nbr</i> I (NaeI)	<i>Ncal</i> (HinfI)
<i>Nci</i> I (CauII)	<i>Nco</i> I	<i>Ncu</i> I (MboII)	<i>Nda</i> I (NarI)
<i>Nde</i> I	<i>Nde</i> II (MboI)	<i>Nfi</i> I (MboI)	<i>Nfi</i> AI (EcoRV)
<i>Nfi</i> AI (MboI)	<i>Nfi</i> BI (MboI)	<i>Ngb</i> I (PstI)	<i>Ngol</i> (HaeII)
<i>Ngo</i> II (HaeIII)	<i>Ngo</i> III (SacII)	<i>Ngo</i> AIII (SacII)	<i>Ngo</i> AIV (NaeI)
<i>Ngo</i> BI (HphI)	<i>Ngo</i> DI (SacII)	<i>Ngo</i> DIII (DpnI)	<i>Ngo</i> MI (NaeI)
<i>Ngo</i> PII (HaeIII)	<i>Ngo</i> PIII (SacII)	<i>Ngo</i> SI (HaeIII)	<i>Nhe</i> I
<i>Nla</i> I (HaeIII)	<i>Nla</i> II (MboI)	<i>Nla</i> III	<i>Nla</i> IV
<i>Nla</i> DI (MboI)	<i>Nla</i> DII (AsuI)	<i>Nla</i> DIII (SacII)	<i>Nla</i> SI (SacII)
<i>Nla</i> SII (AcyI)	<i>Nli</i> I (AvaI)	<i>Nli</i> II (AvaII)	<i>Nme</i> CI (MboI)

<i>NmeRI</i> (<i>PvuII</i>)	<i>NmiI</i> (<i>KpnI</i>)	<i>NmuI</i> (<i>NaeI</i>)	<i>NmuAI</i> (<i>AvaI</i>)
<i>NmuAII</i> (<i>AvaII</i>)	<i>NmuDI</i> (<i>DpnI</i>)	<i>NmuEI</i> (<i>DpnI</i>)	<i>NmuEII</i> (<i>AsuI</i>)
<i>NmuFI</i> (<i>NaeI</i>)	<i>NmuSI</i> (<i>AsuI</i>)	<i>NocI</i> (<i>PstI</i>)	<i>NopI</i> (<i>SalI</i>)
<i>NotI</i>	<i>NovII</i> (<i>HinfI</i>)	<i>NphI</i> (<i>MboI</i>)	<i>NruI</i>
<i>NsiI</i> (<i>AvaIII</i>)	<i>NsiAI</i> (<i>MboI</i>)	<i>NsiCI</i> (<i>EcoRV</i>)	<i>NsiHI</i> (<i>HinfI</i>)
<i>NspI</i>	<i>NspII</i> (<i>SduI</i>)	<i>NspIII</i> (<i>AvaI</i>)	<i>NspIV</i> (<i>AsuI</i>)
<i>NspV</i> (<i>AsuII</i>)	<i>NspAI</i> (<i>MboI</i>)	<i>NspBI</i> (<i>AsuII</i>)	<i>NspBII</i>
<i>NspDI</i> (<i>AvaI</i>)	<i>NspDII</i> (<i>AvaII</i>)	<i>NspEI</i> (<i>AvaI</i>)	<i>NspFI</i> (<i>AsuII</i>)
<i>NspGI</i> (<i>AvaII</i>)	<i>NspHI</i> (<i>NspI</i>)	<i>NspHII</i> (<i>AvaII</i>)	<i>NspHIII</i> (<i>MsrI</i>)
<i>NspJI</i> (<i>AsuII</i>)	<i>NspKI</i> (<i>AvaII</i>)	<i>NspLI</i> (<i>MstI</i>)	<i>NspLII</i> (<i>AsuI</i>)
<i>NspMI</i> (<i>MsrI</i>)	<i>NspMACI</i> (<i>BglII</i>)	<i>NspSAI</i> (<i>AvaI</i>)	<i>NspSAII</i> (<i>BstEII</i>)
<i>NspSAIII</i> (<i>NcoI</i>)	<i>NspSAIV</i> (<i>BamHI</i>)	<i>NspWI</i> (<i>NaeI</i>)	<i>NsuI</i> (<i>MboI</i>)
<i>NsuDI</i> (<i>DpnI</i>)	<i>NtaI</i> (<i>Tth111I</i>)	<i>NtaSI</i> (<i>SruI</i>)	<i>NtaSII</i> (<i>NaeI</i>)
<i>NunII</i> (<i>NarI</i>)	<i>OcoI</i> (<i>XhoI</i>)	<i>OtuI</i> (<i>AluI</i>)	<i>OtuNI</i> (<i>AluI</i>)
<i>OxaI</i> (<i>AluI</i>)	<i>OxaNI</i> (<i>SauI</i>)	<i>PaeI</i> (<i>SphI</i>)	<i>Pae177I</i> (<i>BamHI</i>)
<i>Pae181I</i> (<i>CauII</i>)	<i>PaeAI</i> (<i>SacII</i>)	<i>PaeBI</i> (<i>SmaI</i>)	<i>PaeR7I</i> (<i>XhoI</i>)
<i>PaII</i> (<i>HaeIII</i>)	<i>PaII</i> (<i>HaeIII</i>)	<i>PanI</i> (<i>XhoI</i>)	<i>Pde12I</i> (<i>AsuI</i>)
<i>Pde133I</i> (<i>HaeIII</i>)	<i>Pde137I</i> (<i>HpaII</i>)	<i>Pei9403I</i> (<i>MboI</i>)	<i>PfaI</i> (<i>MboI</i>)
<i>PflAI</i> (<i>FnuDII</i>)	<i>PfIM1</i>	<i>PfIN1</i> (<i>XhoI</i>)	<i>PfFWI</i> (<i>XhoI</i>)
<i>PfuI</i> (<i>SplI</i>)	<i>PgII</i> (<i>NaeI</i>)	<i>PgIB4I</i> (<i>Clal</i>)	<i>PleI</i>
<i>Ple19I</i> (<i>PvuI</i>)	<i>PmaI</i> (<i>PstI</i>)	<i>Pma44I</i> (<i>PstI</i>)	<i>PmaCI</i>
<i>Pme55I</i> (<i>StuI</i>)	<i>PmlI</i> (<i>PmaCI</i>)	<i>PmyI</i> (<i>PstI</i>)	<i>PovI</i> (<i>BclI</i>)
<i>PpaI</i> (<i>Eco31I</i>)	<i>Pph3215I</i> (<i>HgiAI</i>)	<i>PpuI</i> (<i>HaeIII</i>)	<i>PpuMI</i>
<i>PseI</i> (<i>AsuI</i>)	<i>PshAI</i>	<i>PspI</i> (<i>AsuI</i>)	<i>Psp61I</i> (<i>NaeI</i>)
<i>PssI</i> (<i>DraII</i>)	<i>PstI</i>	<i>Psu161I</i> (<i>PvuI</i>)	<i>PvuI</i>
<i>PvuII</i>	<i>PvuHKUI</i> (<i>PvuII</i>)	<i>RffFI</i> (<i>SalI</i>)	<i>RheI</i> (<i>SalI</i>)
<i>RhpI</i> (<i>SalI</i>)	<i>RhsI</i> (<i>BamHI</i>)	<i>RleAI</i>	<i>RluI</i> (<i>NaeI</i>)
<i>Rlu1I</i> (<i>MboI</i>)	<i>Rlu3I</i> (<i>NlaIV</i>)	<i>Rlu4I</i> (<i>BamHI</i>)	<i>RrhI</i> (<i>SalI</i>)
<i>RroI</i> (<i>SalI</i>)	<i>RsaI</i>	<i>RshI</i> (<i>PvuI</i>)	<i>RshII</i> (<i>CauII</i>)
<i>RspI</i> (<i>PvuI</i>)	<i>RspXI</i> (<i>BspHI</i>)	<i>RsrI</i> (<i>EcoRI</i>)	<i>RsrII</i>
<i>SaaI</i> (<i>SacII</i>)	<i>SabI</i> (<i>SacII</i>)	<i>SacI</i>	<i>SacII</i>
<i>SacAI</i> (<i>NaeI</i>)	<i>SakI</i> (<i>SacII</i>)	<i>SalI</i>	<i>Sal1974I</i> (<i>XhoI</i>)
<i>SalAI</i> (<i>MboI</i>)	<i>SalCI</i> (<i>NaeI</i>)	<i>SalDI</i> (<i>NruI</i>)	<i>SalHI</i> (<i>MboI</i>)
<i>SalPI</i> (<i>PstI</i>)	<i>SaoI</i> (<i>NaeI</i>)	<i>SarI</i> (<i>StuI</i>)	<i>Saul</i>
<i>Sau10I</i> (<i>KpnI</i>)	<i>Sau12I</i> (<i>Eco31I</i>)	<i>Sau96I</i> (<i>AsuI</i>)	<i>Sau3239I</i> (<i>XhoI</i>)
<i>Sau6782I</i> (<i>MboI</i>)	<i>SauAI</i> (<i>NaeI</i>)	<i>Sau3AI</i> (<i>MboI</i>)	<i>SauBI</i> (<i>AsuI</i>)
<i>SauBMKI</i> (<i>NaeI</i>)	<i>SauCI</i> (<i>MboI</i>)	<i>SauDI</i> (<i>MboI</i>)	<i>SauEI</i> (<i>MboI</i>)
<i>SauFI</i> (<i>MboI</i>)	<i>SauGI</i> (<i>MboI</i>)	<i>SauMI</i> (<i>MboI</i>)	<i>SbaI</i> (<i>PvuII</i>)
<i>SblAI</i> (<i>SryI</i>)	<i>SbIBI</i> (<i>SryI</i>)	<i>SblCI</i> (<i>SryI</i>)	<i>SboI</i> (<i>SacII</i>)
<i>Sbo13I</i> (<i>NruI</i>)	<i>ScaI</i>	<i>Sca1827I</i> (<i>XhoI</i>)	<i>SceI</i> (<i>FnuDII</i>)
<i>Scg2I</i> (<i>EcoRII</i>)	<i>ScII</i> (<i>XhoI</i>)	<i>Sci1831I</i> (<i>XhoI</i>)	<i>SciAI</i> (<i>BstEII</i>)
<i>SciAII</i> (<i>PvuII</i>)	<i>SciNI</i> (<i>HhaI</i>)	<i>ScoI</i> (<i>SacI</i>)	<i>ScrFI</i>
<i>Scul</i> (<i>XhoI</i>)	<i>SduI</i>	<i>Sdyl</i> (<i>AsuI</i>)	<i>SecI</i>
<i>SecII</i> (<i>HpaII</i>)	<i>SecIII</i> (<i>SauI</i>)	<i>SexI</i> (<i>XhoI</i>)	<i>Sfal</i> (<i>HaeIII</i>)
<i>SfaGUI</i> (<i>HpaII</i>)	<i>SfaNI</i>	<i>SfeI</i>	<i>Sfil</i>
<i>Sfil</i> (<i>PstI</i>)	<i>Sft2aI</i> (<i>EcoRII</i>)	<i>Sft2bI</i> (<i>EcoRII</i>)	<i>SfnI</i> (<i>AvaII</i>)
<i>SfoI</i> (<i>NarI</i>)	<i>SfH</i> (<i>SacII</i>)	<i>Sfr274I</i> (<i>XhoI</i>)	<i>Sfr303I</i> (<i>SacII</i>)
<i>Sfr382I</i> (<i>SacII</i>)	<i>Sful</i> (<i>AsuII</i>)	<i>Sfu1762I</i> (<i>XhoI</i>)	<i>SgaI</i> (<i>XhoI</i>)
<i>Sgh1835I</i> (<i>AvaII</i>)	<i>SgoI</i> (<i>XhoI</i>)	<i>Sgr20I</i> (<i>EcoRII</i>)	<i>Sgr1839I</i> (<i>AsuII</i>)
<i>Sgr1841I</i> (<i>XhoI</i>)	<i>SgrAI</i>	<i>Shyl</i> (<i>SacII</i>)	<i>Shy1766I</i> (<i>XhoI</i>)
<i>SinI</i> (<i>AvaII</i>)	<i>SinAI</i> (<i>AvaII</i>)	<i>SinBI</i> (<i>AvaII</i>)	<i>SinCI</i> (<i>AvaII</i>)
<i>SinDI</i> (<i>AvaII</i>)	<i>SinEI</i> (<i>AvaII</i>)	<i>SinFI</i> (<i>AvaII</i>)	<i>SinGI</i> (<i>AvaII</i>)
<i>SinHI</i> (<i>AvaII</i>)	<i>SinJI</i> (<i>AvaII</i>)	<i>SinMI</i> (<i>MboI</i>)	<i>SkaI</i> (<i>NaeI</i>)
<i>SkaII</i> (<i>PstI</i>)	<i>Slal</i> (<i>XhoI</i>)	<i>SleI</i> (<i>EcoRII</i>)	<i>SluI</i> (<i>XhoI</i>)
<i>Slu1777I</i> (<i>NaeI</i>)	<i>SmaI</i>	<i>SmaAI</i> (<i>SpII</i>)	<i>SmaAII</i> (<i>Tth111I</i>)
<i>SmaAIII</i> (<i>PvuI</i>)	<i>SmaAIV</i> (<i>PvuII</i>)	<i>SnaI</i>	<i>Sna3286I</i> (<i>NruI</i>)
<i>SnaBI</i>	<i>SnoI</i> (<i>ApaLI</i>)	<i>Sol3335I</i> (<i>PvuII</i>)	<i>SpaI</i> (<i>XhoI</i>)
<i>SpaXI</i> (<i>SphI</i>)	<i>SpeI</i>	<i>SphI</i>	<i>Sph1719I</i> (<i>XhoI</i>)
<i>Spil</i>	<i>SpII</i>	<i>SpIII</i> (<i>Tth111I</i>)	<i>SpIII</i> (<i>HaeIII</i>)
<i>SplAI</i> (<i>SpII</i>)	<i>SplAII</i> (<i>Tth111I</i>)	<i>SplAIII</i> (<i>PvuI</i>)	<i>SplAIV</i> (<i>PvuII</i>)
<i>SpoI</i> (<i>NruI</i>)	<i>SseI</i> (<i>BclI</i>)	<i>SseII</i> (<i>SacII</i>)	<i>SshAI</i> (<i>SauI</i>)
<i>SsoI</i> (<i>EcoRI</i>)	<i>SsoII</i> (<i>ScrFI</i>)	<i>SspI</i>	<i>SspII</i> (<i>AsuII</i>)
<i>Ssp2I</i> (<i>CauII</i>)	<i>Ssp4I</i> (<i>XhoI</i>)	<i>Ssp152I</i> (<i>AsuII</i>)	<i>Ssp1725I</i> (<i>SacII</i>)
<i>SspAI</i> (<i>EcoRII</i>)	<i>SspJI</i> (<i>SnaBI</i>)	<i>SspJII</i> (<i>Acyl</i>)	<i>SspKI</i> (<i>SpII</i>)
<i>SspM1I</i> (<i>SnaBI</i>)	<i>SspM1II</i> (<i>Acyl</i>)	<i>SspM1III</i> (<i>HgiCI</i>)	<i>SspM2I</i> (<i>SnaBI</i>)
<i>SspM2II</i> (<i>Acyl</i>)	<i>SsrI</i> (<i>HpaI</i>)	<i>SsrB6I</i> (<i>HpaI</i>)	<i>SstI</i> (<i>SacI</i>)
<i>SstII</i> (<i>SacII</i>)	<i>SstIV</i> (<i>BclI</i>)	<i>SsvI</i> (<i>StuI</i>)	<i>Stel</i> (<i>StuI</i>)
<i>StiI</i> (<i>KpnI</i>)	<i>StiAI</i> (<i>KpnI</i>)	<i>StiBI</i> (<i>KpnI</i>)	<i>StiCI</i> (<i>KpnI</i>)
<i>StiDI</i> (<i>KpnI</i>)	<i>StiEI</i> (<i>KpnI</i>)	<i>StiFI</i> (<i>KpnI</i>)	<i>StiGI</i> (<i>KpnI</i>)

<i>Sth</i> HI (<i>Kpn</i> I)	<i>Sth</i> II (<i>Kpn</i> I)	<i>Sth</i> KI (<i>Kpn</i> I)	<i>Sth</i> LI (<i>Kpn</i> I)
<i>Sth</i> MI (<i>Kpn</i> I)	<i>Sth</i> NI (<i>Kpn</i> I)	<i>Stu</i> I	<i>Sry</i> I
<i>Sua</i> I (<i>Hae</i> III)	<i>Sul</i> I (<i>Hae</i> III)	<i>Sur</i> 2I (<i>Bam</i> HI)	<i>Sve</i> 194I (<i>Xho</i> I)
<i>Taq</i> I	<i>Taq</i> II	<i>Taq</i> XI (<i>Eco</i> RII)	<i>Tce</i> I (<i>Mbo</i> II)
<i>Tfi</i> I	<i>Tfi</i> I (<i>Taq</i> I)	<i>Tgi</i> I (<i>Sac</i> II)	<i>Tha</i> I (<i>Fnu</i> DII)
<i>Tma</i> I (<i>Fnu</i> DII)	<i>Tmu</i> II (<i>Cau</i> II)	<i>Tru</i> I (<i>Ava</i> II)	<i>Tru</i> II (<i>Mbo</i> I)
<i>Tru</i> 9I (<i>Mse</i> I)	<i>Tru</i> 20II (<i>Xho</i> II)	<i>Tsp</i> I (<i>Tth</i> 111I)	<i>Tsp</i> 45I
<i>Tsp</i> EI	<i>Tsp</i> ZNI (<i>Hae</i> III)	<i>Tie</i> I (<i>Tth</i> 111I)	<i>Tte</i> AI (<i>Hae</i> III)
<i>Tth</i> 111I	<i>Tth</i> 111II	<i>Tth</i> HB8I (<i>Taq</i> I)	<i>Tmi</i> I (<i>Hae</i> III)
<i>Tir</i> I (<i>Tth</i> 111I)	<i>Uba</i> II (<i>Ppu</i> MI)	<i>Uba</i> 6I (<i>Mlu</i> I)	<i>Uba</i> 1102I (<i>Esp</i> I)
<i>Uba</i> 1103I (<i>Ava</i> III)	<i>Uba</i> 1103II (<i>Dpn</i> I)	<i>Uba</i> 1103II (<i>Dpn</i> I)	<i>Uba</i> 1104I (<i>Ksp</i> 632I)
<i>Uba</i> 1105I	<i>Uba</i> 1106I (<i>Ppu</i> MI)	<i>Uba</i> 1107I (<i>Sna</i> I)	<i>Uba</i> 1108I
<i>Uba</i> 1109I (<i>Bbv</i> I)	<i>Uba</i> 1110I (<i>Bam</i> HI)	<i>Uur</i> 960I (<i>Fnu</i> 4HI)	<i>Van</i> I (<i>Bgl</i> I)
<i>Var</i> 9II (<i>Pfi</i> MI)	<i>Vha</i> I (<i>Hae</i> III)	<i>Vha</i> 464I (<i>Afi</i> II)	<i>Vne</i> I (<i>Apa</i> LI)
<i>Vne</i> AI (<i>Dra</i> II)	<i>Vni</i> I (<i>Hae</i> III)	<i>Vsp</i> I	<i>Xam</i> I (<i>Sal</i> I)
<i>Xba</i> I	<i>Xca</i> I (<i>Sna</i> I)	<i>Xci</i> I (<i>Sal</i> I)	<i>Xcm</i> I
<i>Xcy</i> I (<i>Sma</i> I)	<i>Xgl</i> 3216I (<i>Pvu</i> I)	<i>Xgl</i> 3217I (<i>Pvu</i> I)	<i>Xgl</i> 3218I (<i>Pvu</i> I)
<i>Xgl</i> 3219I (<i>Pvu</i> I)	<i>Xgl</i> 3220I (<i>Pvu</i> I)	<i>Xho</i> I	<i>Xho</i> II
<i>Xma</i> I (<i>Sma</i> I)	<i>Xma</i> II (<i>Pst</i> I)	<i>Xma</i> III	<i>Xmi</i> I (<i>Pvu</i> I)
<i>Xmi</i> AI (<i>Pvu</i> I)	<i>Xmn</i> I	<i>Xni</i> I (<i>Pvu</i> I)	<i>Xor</i> I (<i>Pst</i> I)
<i>Xor</i> I (<i>Pvu</i> I)	<i>Xpa</i> I (<i>Xho</i> I)	<i>Xph</i> I (<i>Pst</i> I)	<i>Yen</i> I (<i>Pst</i> I)
<i>Yen</i> AI (<i>Pst</i> I)	<i>Yen</i> BI (<i>Pst</i> I)	<i>Yen</i> CI (<i>Pst</i> I)	<i>Yen</i> DI (<i>Pst</i> I)
<i>Yen</i> EI (<i>Pst</i> I)	<i>Zan</i> I (<i>Eco</i> RII)	<i>Zsp</i> 2I (<i>Ava</i> III)	

unknown recognition sequences

<i>Aam</i> I	<i>Acy</i> II	<i>Aim</i> I	<i>Ani</i> I
<i>Atu</i> AI	<i>Atu</i> BVI	<i>Atu</i> LAMI	<i>Bbe</i> II
<i>Bbe</i> AI	<i>Bbe</i> SI	<i>Bbi</i> IV	<i>Bce</i> 1229I
<i>Bce</i> 14579I	<i>Blo</i> I	<i>Bme</i> I	<i>Bme</i> 205I
<i>Bme</i> 899I	<i>Bpr</i> I	<i>Bsi</i> BI	<i>Bsi</i> EI
<i>Bsi</i> FI	<i>Bsi</i> GI	<i>Bsi</i> II	<i>Bsi</i> NI
<i>Bsi</i> PI	<i>Bsi</i> RI	<i>Bsi</i> TI	<i>Bsi</i> UI
<i>Bsi</i> VI	<i>Bsi</i> YI	<i>Bsp</i> 12II	<i>Bsr</i> PI
<i>Bss</i> PI	<i>Bst</i> AI	<i>Bst</i> EI	<i>Bsu</i> 1145I
<i>Bsu</i> 1259I	<i>Cal</i> I	<i>Chi</i> I	<i>Clf</i> III
<i>Clu</i> I	<i>Csu</i> I	<i>Cve</i> I	<i>Cvi</i> I
<i>Cvi</i> PI	<i>Cvi</i> QII	<i>Dmo</i> I	<i>Eco</i> CKI
<i>Eco</i> O34I	<i>Eco</i> O44I	<i>Esp</i> II	<i>Fnu</i> 48I
<i>Fsa</i> I	<i>Ggi</i> I	<i>Gsp</i> AIII	<i>Hag</i> I
<i>Hap</i> I	<i>Hcu</i> I	<i>Hgi</i> FI	<i>Hgi</i> KI
<i>Hhl</i> II	<i>Hin</i> 1056II	<i>Hsa</i> I	<i>Lpn</i> I
<i>Lpn</i> II	<i>Mbv</i> I	<i>Mgi</i> II	<i>Mgi</i> II
<i>Mnn</i> III	<i>Mno</i> II (<i>Mnn</i> III)	<i>Msi</i> II	<i>Mvi</i> I
<i>Mvi</i> II	<i>Nfi</i> II	<i>Nfi</i> III	<i>Ngo</i> DII
<i>Nme</i> I	<i>Nme</i> II	<i>Nme</i> III	<i>Nme</i> IV
<i>Nop</i> II	<i>Nov</i> I	<i>Nsp</i> EII	<i>Nsp</i> LIII
<i>Nsp</i> LIV	<i>Nun</i> I	<i>Oxa</i> II	<i>Pfi</i> I
<i>Pgi</i> II	<i>Pmi</i> I	<i>Pss</i> II	<i>Rhp</i> II
<i>Rle</i> I	<i>Rme</i> I	<i>Rrb</i> I	<i>Rrh</i> II
<i>ft2Rru</i> AI	<i>Sac</i> III	<i>Sal</i> II	<i>San</i> I
<i>Sbr</i> I	<i>Sex</i> II	<i>Sgr</i> I	<i>Shy</i> TI
<i>Sin</i> MII	<i>Sis</i> I	<i>Sod</i> I	<i>Sod</i> II
<i>Ssa</i> I	<i>Ssc</i> I	<i>Ssp</i> XI	<i>Sst</i> III (<i>Sac</i> III)
<i>Stm</i> I	<i>Sry</i> D4I	<i>Tmi</i> I	<i>Tru</i> III