
Restriction enzymes and their isoschizomers

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INTRODUCTION

Since the last compilation of restriction enzymes (1), 156 new entries have been added including 12 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 is presented in the second Table. A copy of the restriction enzyme data base 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 online through the BIONET computer resource. A version corresponding to the printed text is located in the file
<ROBERTS>RESTRICT.NAR Several alternative versions are available and are documented in
<ROBERTS>RESTRICT.DOC

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).

Several enzymes appear in this list with revised names. These revisions were made to avoid confusion with existing enzymes or to increase the uniformity of the names. In each case the name changes were made with the approval of the appropriate authors. The specific changes are as follows. *Eag*KI replaces *Eag*I (5) and *Ani*MI replaces *Ani*I (130), since the latter names had been used for other enzymes. *Nsp*I-V replaces *Nsp*(7524)I-V (6). This had already been used in many publications and catalogs. *Cvi*QI replaces *Cvi*II (7) to show that this enzyme comes from a strain different from that encoding *Cvi*I.

Type II enzymes

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|---------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| <i>Aat</i> II | | GACGT↓C | | MNRU | 8 |
| <i>Acc</i> I | | GT↓MKAC | | ABCIMNPRU | 9 |
| <i>Acy</i> I | | GR↓CGYC | | | 10 |
| <i>Aha</i> II | | GR↓CGYC | | N | 11 |
| <i>Aos</i> II | | GR↓CGYC | | | 12 |
| <i>Ast</i> WI | | GR↓CGYC | | | 13 |
| <i>Asu</i> III | | GR↓CGYC | | | 13 |
| <i>Bbi</i> II | | GR↓CGYC | | A | 14 |
| <i>Hgi</i> DI | | GR↓CGYC | | | 15 |
| <i>Hgi</i> GI | | GR↓CGYC | | | 15 |
| <i>Hgi</i> HII | | GR↓CGYC | | | 16 |
| <i>Nla</i> SII | | GRCGYC | | | 17 |
| <i>Afl</i> II | | C↓TTAAG | | AGN | 18 |
| <i>Afl</i> III | | A↓CRYGT | | G | 18 |
| <i>Aha</i> III | | TTT↓AAA | | | 19 |
| <i>Dra</i> I | | TTT↓AAA | | ABGIMNPRU | 20 |
| <i>Alu</i> I | | AG↓CT | 3(5) | ABGIMNPRU | 17,21-24 |
| <i>Mlt</i> I | | AG↓CT | | | 25,26 |
| <i>Otu</i> I | | AGCT | | | 27 |
| <i>Otu</i> NI | | AGCT | | | 28 |
| <i>Oxa</i> I | | AGCT | | | 29 |
| <i>Ahv</i> NI | | CAGNNN↓CTG | | N | 30 |
| <i>Apa</i> I | | GGGCC↓C | 4(5) | BGIMNPRU | 31,32 |
| <i>Apa</i> LI | | G↓TGCAC | | AGN | 33 |
| <i>Amel</i> | | GTGCAC | | | 27 |
| <i>Sno</i> I | | G↓TGCAC | | | 34,35 |
| <i>Vne</i> I | | G↓TGCAC | | | 36 |
| <i>Asu</i> I | | G↓GNCC | | P | 37 |
| <i>Apu</i> I | | GGNCC | | | 38 |
| <i>Bac</i> 36 | | G↓GNCC | | | 38 |
| <i>Bsp</i> BII | | G↓GNCC | | | 39 |
| <i>Cfr</i> 4I | | GGNCC | | | 41,44 |
| <i>Cfr</i> 8I | | GGNCC | | | 41,44 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| | <i>Cfr</i> 13I | G↓GNCC | 4(5) | AU | 40,41 |
| | <i>Cfr</i> 23I | GGNCC | | | 42 |
| | <i>Cfr</i> 33I | GGNCC | | | 43 |
| | <i>Cfr</i> 45I | GGNCC | | | 43 |
| | <i>Cfr</i> 46I | GGNCC | | | 43 |
| | <i>Cfr</i> 47I | GGNCC | | | 43 |
| | <i>Cfr</i> NI | GGNCC | | | 45 |
| | <i>Eco</i> 39I | GGNCC | | | 46 |
| | <i>Eco</i> 47II | GGNCC | | | 47 |
| | <i>Mja</i> II | GGNCC | | | 48 |
| | <i>Nla</i> DII | GGNCC | | | 49 |
| | <i>Nmu</i> EII | GGNCC | | | 50 |
| | <i>Nmu</i> SI | GGNCC | | | 51 |
| | <i>Nsp</i> IV | G↓GNCC | | R | 6 |
| | <i>Psp</i> I | GGNCC | | | 52 |
| | <i>Sau</i> 96I | G↓GNCC | | BGMNP | 53 |
| | <i>Sdy</i> I | GGNCC | | | 54 |
| <i>Asu</i> II | | TT↓CGAA | | GP | 13,55 |
| | <i>Bst</i> BI | TTCGAA | | N | 56 |
| | <i>Fsp</i> II | TT↓CGAA | | | 57 |
| | <i>Lsp</i> I | TT↓CGAA | | | 34,35 |
| | <i>Mla</i> I | TT↓CGAA | | | 58 |
| | <i>Nsp</i> V | TTCGAA | | AR | 6 |
| | <i>Nsp</i> BI | TTCGAA | | | 59 |
| <i>Ava</i> I | | C↓YCGRG | | ABCIMNPRU | 60,61 |
| | <i>Aqu</i> I | C↓YCGRG | 1(5) | | 62,63 |
| | <i>Avr</i> I | CYCGRG | | | 64 |
| | <i>Bst</i> SI | C↓YCGRG | | | 56 |
| | <i>Eco</i> 88I | CYCGRG | | | 65 |
| | <i>Nsp</i> III | C↓YCGRG | | R | 6 |
| | <i>Nsp</i> SAI | C↓YCGRG | | | 66 |
| <i>Ava</i> II | | G↓GWCC | | ABGIMNPR | 60,61,67,68 |
| | <i>Afl</i> I | G↓GWCC | | | 18 |
| | <i>Asp</i> 697I | GGWCC | | | 69 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|-----------------------------|---------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| <i>Bam</i> N _x I | | G↓GWCC | | | 70-72 |
| <i>Bme</i> 216I | | G↓GWCC | | | 73,74 |
| <i>Btl</i> | | GGWCC | | | 75 |
| <i>Cau</i> I | | G↓GWCC | | | 76,77 |
| <i>Cla</i> II | | GGWCC | | | 78 |
| <i>Eag</i> MI | | G↓GWCC | | | 79 |
| <i>Eco</i> 47I | | G↓GWCC | U | | 47,80 |
| <i>Erp</i> I | | G↓GWCC | | | 38 |
| <i>Fdl</i> I | | G↓GWCC | | | 81,82 |
| <i>Fsp</i> MSI | | G↓GWCC | | | 38 |
| <i>Gsp</i> AI | | GGWCC | | | 83 |
| <i>Hgi</i> BI | | G↓GWCC | | | 15 |
| <i>Hgi</i> CII | | G↓GWCC | | | 15 |
| <i>Hgi</i> EI | | G↓GWCC | | | 15 |
| <i>Hgi</i> HIII | | G↓GWCC | | | 16 |
| <i>Hgi</i> JII | | G↓GWCC | | | 16 |
| <i>Nsp</i> HII | | GGWCC | | | 59 |
| <i>Sfn</i> I | | GGWCC | | | 84 |
| <i>Sin</i> AI | | GGWCC | | | 85 |
| <i>Sin</i> BI | | GGWCC | | | 85 |
| <i>Sin</i> CI | | GGWCC | | | 85 |
| <i>Sin</i> DI | | GGWCC | | | 85 |
| <i>Sin</i> EI | | GGWCC | | | 85 |
| <i>Sin</i> FI | | GGWCC | | | 85 |
| <i>Sin</i> GI | | GGWCC | | | 85 |
| <i>Sin</i> HI | | GGWCC | | | 85 |
| <i>Sin</i> I | | G↓GWCC | P | | 86,87 |
| <i>Sin</i> JII | | GGWCC | | | 85 |
| <i>Tru</i> I | | GGWCC | | | 88 |
| <i>Avr</i> III | | ATGCAT | G | | 89,90 |
| <i>Eco</i> T22I | | ATGCA↓T | U | | 91 |
| <i>Nsi</i> I | | ATGCA↓T | BMNP | | 92 |
| <i>Avr</i> II | | C↓CTAGG | N | | 64,93 |
| <i>Bal</i> I | | TCG↓CCA | 4(5) | ABGIN | 94,32 |
| <i>Bam</i> HI | | G↓GATCC | 5(5) | ABGIMNPRU | 95-97 |

| Enzyme ¹ | Iisoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|------------------|--------------------------------------|-------------------------|-----------------------------------|------------|
| | <i>AacI</i> | GGATCC | | | 98 |
| | <i>AaeI</i> | GGATCC | | | 98 |
| | <i>AccEBI</i> | G↓GATCC | | | 99 |
| | <i>Ali12257I</i> | GGATCC | | | 100 |
| | <i>Ali12258I</i> | GGATCC | | | 100 |
| | <i>AliI</i> | G↓GATCC | | | 101 |
| | <i>BamFI</i> | GGATCC | | | 102 |
| | <i>BamKI</i> | GGATCC | | | 102 |
| | <i>BamNI</i> | GGATCC | | | 71 |
| | <i>BstI</i> | G↓GATCC | | GR | 103,104 |
| | <i>BstQI</i> | GGATCC | | | 56 |
| | <i>CelI</i> | GGATCC | | | 105 |
| | <i>DdsI</i> | GGATCC | | | 106 |
| | <i>GdiI</i> | GGATCC | | | 98 |
| | <i>GinI</i> | GGATCC | | | 107 |
| | <i>GoxI</i> | GGATCC | | | 98 |
| | <i>MleI</i> | GGATCC | | | 25 |
| | <i>NasBI</i> | GGATCC | | | 25 |
| | <i>NspSAIV</i> | G↓GATCC | | | 66 |
| | <i>RhsI</i> | GGATCC | | | 108 |
| <i>BbvI</i> | | GCAGC(8/12) | | GIN | 97,109-111 |
| | <i>AtwXI</i> | GCAGC(8/12) | | | 112 |
| <i>BbvII</i> | | GAAGAC(2/6) | | | 113 |
| <i>BclII</i> | | T↓GATCA | | ABGIMNRU | 114 |
| | <i>AtuCI</i> | TGATCA | | | 115 |
| | <i>BspXII</i> | T↓GATCA | | | 116 |
| | <i>BstGI</i> | TGATCA | | | 117 |
| | <i>BstKI</i> | TGATCA | | | 56 |
| | <i>CpeI</i> | TGATCA | | | 118 |
| | <i>CthI</i> | TGATCA | | | 119 |
| | <i>FbaI</i> | TGATCA | | | 84 |
| | <i>PovI</i> | T↓GATCA | | | 120 |
| | <i>SstIV</i> | TGATCA | | | 121 |
| <i>BglII</i> | | GCCNNNN↓NGGC | | BGIMNPRU | 122-125 |
| | <i>VanI</i> | GCCNNNNNGGC | | | 126 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| <i>Bgl</i> III | | A↓GATCT | | ABGIMNPRU | 122,123,127 |
| | <i>Nsp</i> MACI | A↓GATCT | | | 128 |
| <i>Bin</i> I | | GGATC(4/5) | | | 129 |
| | <i>Alw</i> I | GGATC(4/5) | N | | 130 |
| <i>Bse</i> PI | <i>Bth</i> II | GGATC | | | 131 |
| | | GCGCC | | | 117 |
| <i>Bso</i> PI | | GCGCC | | | 117 |
| | <i>Bsr</i> HI | GCGCC | | | 117 |
| <i>Bss</i> HII | | G↓CGCC | GN | | 117,476 |
| | | GTCTC | | | 132 |
| <i>Bsm</i> AI | | GAATGC(1/-1) | GN | | 55 |
| <i>Bsm</i> I | | T↓CATGA | N | | 133 |
| <i>Bsp</i> HI | <i>Rsp</i> XI | T↓CATGA | G | | 134 |
| <i>Bsp</i> MI | | ACCTGC(4/8) | N | | 130,135 |
| <i>Bsp</i> MII | | T↓CCGA | N | | 130,135 |
| | <i>Acc</i> III | T↓CCGA | AG | | 55,136 |
| | <i>Kpn</i> 2I | TCCGA | | | 137 |
| | <i>Mro</i> I | T↓CCGA | U | | 138 |
| <i>Bsr</i> I | | ACTGG(1/-1) | | | 27 |
| <i>Bst</i> EII | | G↓GTNACC | BGMNRU | | 139,140 |
| | <i>Asp</i> AI | G↓GTNACC | | | 35 |
| | <i>Bst</i> 3I | GGTNACC | | | 141 |
| | <i>Bst</i> DI | GGTNACC | | | 56 |
| | <i>Bst</i> PI | G↓GTNACC | | | 142 |
| | <i>Cfr</i> 7I | GGTNACC | | | 41 |
| | <i>Cfr</i> 19I | GGTNACC | | | 43 |
| | <i>Eca</i> I | G↓GTNACC | | | 143 |
| | <i>Eco</i> 91I | GGTNACC | | | 144 |
| | <i>Eco</i> O65I | G↓GTNACC | | | 145,146 |
| | <i>Kox</i> I | G↓GTNACC | | | 147 |
| | <i>Nsp</i> SAII | G↓GTNACC | | | 66 |
| <i>Bst</i> XI | | CCANNNNN↓NTGG | GNP | | 117,148 |
| | <i>Bss</i> GI | CCANNNNNNTGG | | | 117 |
| | <i>Bst</i> TI | CCANNNNNNTGG | | | 117 |
| <i>Cau</i> II | | CC↓SGG | | | 76,77,149 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|------------------|--------------------------------------|-------------------------|-----------------------------------|----------------|
| | <i>Aha</i> I | CC↓SGG | | | 11 |
| | <i>Ase</i> II | CC↓SGG | | | 150 |
| | <i>Bcn</i> I | CC↓SGG | 2(4) | A | 151-155 |
| | <i>Hgs</i> S21I | CCSGG | | | 156 |
| | <i>Nci</i> I | CC↓SGG | | BGMNU | 157,158 |
| | <i>Rsh</i> II | CCSGG | | | 73 |
| <i>Cfr</i> 10I | | R↓CCGGY | 2(5) | AU | 41,44,155,159 |
| <i>Cfr</i> I | | Y↓GGCCR | 4(5) | | 44,155,160,161 |
| | <i>Cfr</i> 14I | YGGCCR | | | 41 |
| | <i>Cfr</i> 38I | YGGCCR | | | 42 |
| | <i>Cfr</i> 39I | YGGCCR | | | 43 |
| | <i>Cfr</i> 40I | YGGCCR | | | 43 |
| | <i>Eae</i> I | Y↓GGCCR | 4(5) | GN | 162,163 |
| | <i>Eco</i> 90I | YGGCCR | | | 144 |
| | <i>Eco</i> 164I | YGGCCR | | | 164 |
| | <i>Eco</i> HI | YGGCCR | | | 28 |
| <i>Clal</i> | | AT↓CGAT | | ABGMNR | 165 |
| | <i>Asp</i> 707I | ATCGAT | | | 69 |
| | <i>Ban</i> III | ATCGAT | | U | 8 |
| | <i>Bcm</i> I | AT↓CGAT | | | 166 |
| | <i>Bsc</i> I | AT↓CGAT | | | 35,167 |
| | <i>Bsp</i> 106I | AT↓CGAT | | | 168 |
| | <i>Bsp</i> XI | AT↓CGAT | | | 116 |
| <i>Cvi</i> JI | | RG↓CY | | | 169 |
| | <i>Cvi</i> KI | RGCY | | | 170 |
| | <i>Cvi</i> LI | RGCY | | | 170 |
| | <i>Cvi</i> MI | RGCY | | | 170 |
| | <i>Cvi</i> NI | RGCY | | | 170 |
| | <i>Cvi</i> OI | RGCY | | | 170 |
| <i>Dde</i> I | | C↓TNAG | 1(5) | BGIMNPRU | 171-173 |
| <i>Dpn</i> I* | | GA↓TC | | ABGIMNP | 174-176 |
| | <i>Cfu</i> I* | GA↓TC | | | 177,178 |
| | <i>Nan</i> II* | GATC | | | 179 |
| | <i>Ngo</i> DIII* | GATC | | | 180 |
| | <i>Nmu</i> DI* | GATC | | | 50 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|-----------------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| | <i>NmuEI*</i> | GATC | | | 181 |
| | <i>NsuDI*</i> | GATC | | | 50 |
| <i>Dra</i> II | | RG↓GNCCY | | GM | 182-184 |
| | <i>EcoO109I</i> | RG↓GNCCY | | AGN | 185 |
| | <i>Pss</i> I | RGGNC↓CY | | I | 106,186 |
| <i>Dra</i> III | | CACNNN↓GTC | | MN | 182-184 |
| <i>Dsa</i> I | | C↓CRYGG | | | 187 |
| <i>Ear</i> I | | CTCTTC | | | 27 |
| <i>Eco31</i> I | | GGTCTC(1/5) | | | 188 |
| | <i>Eco42</i> I | GGTCTC | | | 189 |
| | <i>Eco51</i> I | GGTCTC | | | 164 |
| | <i>Eco95</i> I | GGTCTC | | | 191 |
| | <i>Eco97</i> I | GGTCTC | | | 192 |
| | <i>Eco101</i> I | GGTCTC | | | 189 |
| | <i>Eco120</i> I | GGTCTC | | | 189 |
| | <i>Eco127</i> I | GGTCTC | | | 190 |
| | <i>Eco129</i> I | GGTCTC | | | 190 |
| | <i>Eco155</i> I | GGTCTC | | | 189 |
| | <i>Eco156</i> I | GGTCTC | | | 189 |
| | <i>Eco157</i> I | GGTCTC | | | 189 |
| | <i>Eco162</i> I | GGTCTC | | | 190 |
| | <i>Ppa</i> I | GGTCTC | | | 130 |
| <i>Eco47</i> III | | AGC↓GCT | | AU | 47 |
| | <i>Aii</i> I | AGC↓GCT | | | 193 |
| <i>Eco57</i> I | | CTGAAG(16/14) | | | 194 |
| | <i>Fs</i> fl | CTGAAG | | | 195 |
| <i>Eco</i> NI | | CCTNNN↓NNNAGG | | | 196 |
| | <i>Bst</i> WI | CCTNNNNNNAGG | | | 132 |
| <i>Eco</i> RI | | G↓AATT | 3(6) | ABGIMNPRU | 197-199 |
| | <i>Eco82</i> I | GAATT | | | 191 |
| | <i>Eco159</i> I | GAATT | | | 190 |
| | <i>Rsr</i> I | G↓AATT | | | 200,201 |
| | <i>Sso</i> I | G↓AATT | | | 202 |
| <i>Eco</i> RII ⁵ | | ↓CCWGG | 2(5) | BG | 203-205 |
| | <i>Aeu</i> I | CC↓WGG | | | 150 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| + | <i>Aor</i> I | CC↓WGG | | | 98 |
| + | <i>Apy</i> I | CC↓WGG | | M | 206 |
| | <i>Atu</i> BI | CCWGG | | | 207 |
| | <i>Atu</i> II | CCWGG | | | 208 |
| | <i>Bir</i> SI | CCWGG | | | 131 |
| | <i>Bst</i> GII | CCWGG | | | 117 |
| + | <i>Bst</i> NI | CC↓WGG | | N | 209 |
| | <i>Bst</i> OI | CCWGG | | | 56 |
| | <i>Cdi</i> 27I | CCWGG | | | 210 |
| | <i>Cfr</i> 5I | CCWGG | | | 41,44 |
| | <i>Cfr</i> 11I | CCWGG | | | 41,44 |
| | <i>Cfr</i> 20I | CCWGG | | | 43 |
| | <i>Cfr</i> 22I | CCWGG | | | 43 |
| | <i>Cfr</i> 24I | CCWGG | | | 43 |
| | <i>Cfr</i> 25I | CCWGG | | | 43 |
| | <i>Cfr</i> 27I | CCWGG | | | 43 |
| | <i>Cfr</i> 28I | CCWGG | | | 43 |
| | <i>Cfr</i> 29I | CCWGG | | | 43 |
| | <i>Cfr</i> 30I | CCWGG | | | 43 |
| | <i>Cfr</i> 31I | CCWGG | | | 43 |
| | <i>Cfr</i> 35I | CCWGG | | | 43 |
| | <i>Cfr</i> S37I | CCWGG | | | 156 |
| | <i>Cth</i> II | CC↓WGG | | | 119 |
| | <i>Eag</i> KI | CCWGG | | | 5 |
| | <i>Ea</i> II | CCWGG | | | 55 |
| | <i>Ecl</i> 66I | CCWGG | | | 192 |
| | <i>Ecl</i> II | CCWGG | | | 211 |
| | <i>Ecl</i> 539I | CCWGG | | | 156 |
| | <i>Eco</i> 38I | CCWGG | | | 46 |
| | <i>Eco</i> 40I | CCWGG | | | 46 |
| | <i>Eco</i> 41I | CCWGG | | | 46 |
| | <i>Eco</i> 60I | CCWGG | | | 44 |
| | <i>Eco</i> 61I | CCWGG | | | 44 |
| | <i>Eco</i> 67I | CCWGG | | | 191 |
| | <i>Eco</i> 70I | CCWGG | | | 191 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-------------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| + | <i>Eco</i> 71I | CCWGG | | | 212 |
| | <i>Eco</i> 128I | CCWGG | | | 190 |
| | <i>Eco</i> 136I | CCWGG | | | 190 |
| | <i>Eco</i> 165I | CCWGG | | | 164 |
| | <i>Mph</i> I | CCWGG | | | 213 |
| | <i>Msp</i> I | CC↓WGG | 2(4) | AU | 214,215 |
| | <i>Sgr</i> II | CCWGG | | | 216 |
| | <i>Taq</i> XI | CC↓WGG | | | 217 |
| | <i>Zan</i> I | CC↓WGG | | | 218 |
| | <i>EcoRV</i> | GAT↓ATC | | ABGIMNPRU | 219,220 |
| <i>Bst</i> RI | | GATATC | | | 56 |
| | <i>Ceq</i> I | GAT↓ATC | | | 221 |
| | <i>Eco</i> 32I | GAT↓ATC | | | 44,222 |
| | <i>Hpa</i> I | GATATC | | | 477 |
| | <i>Nan</i> I | GATATC | | | 179 |
| | <i>Nla</i> II | GATATC | | | 223 |
| | <i>Nsi</i> CI | GAT↓ATC | | | 224 |
| | <i>Esp</i> I | GC↓TNAGC | | G | 225 |
| <i>Fin</i> I | <i>Cel</i> II | GCTNAG | | | 105 |
| | | GTCCC | | | 130 |
| <i>Fnu</i> 4HI | | GC↓NGC | | N | 226 |
| <i>Fnu</i> DII | <i>Fbr</i> I | GC↓NGC | | | 84 |
| | | CG↓CG | | | 227 |
| <i>Acc</i> II | | CG↓CG | | AG | 9,228 |
| | <i>Bce</i> FI | CGCG | | | 229 |
| | <i>Bce</i> RI | CGCG | | | 102 |
| | <i>Bep</i> I | CG↓CG | | | 230 |
| | <i>Bst</i> UI | CG↓CG | | N | 56 |
| | <i>Bsu</i> 1192II | CGCG | | | 54 |
| | <i>Bsu</i> 1193I | CGCG | | | 54,102 |
| | <i>Bsu</i> 6633I | CGCG | | | 102,231 |
| | <i>Bsu</i> EII | CGCG | | | 54,232 |
| | <i>Fsp</i> MI | CGCG | | | 130 |
| <i>Hin</i> 1056I | | CGCG | | | 233 |
| | <i>Mvn</i> I | CG↓CG | | M | 234 |

| Enzyme ¹ | Iisoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| | <i>PflAI</i> | CGCG | | | 27 |
| | <i>ThaI</i> | CG↓CG | | BI | 235 |
| <i>FokI</i> | | GGATG(9/13) | | AMN | 236 |
| | <i>HinGII</i> | GGATG | | | 237-239 |
| <i>GdiII</i> | | YGGCCG(-5/-1) | | | 240 |
| <i>GsuI</i> | | CTGGAG(16/14) | | | 241,242 |
| <i>HaeI</i> | | WGG↓CCW | | | 238 |
| <i>HaeII</i> | | RGC ₁ GC ₂ ↓Y | | ABGIMNR | 243,244 |
| | <i>HinHI</i> | RGC ₁ GC ₂ Y | | | 245 |
| | <i>NgoI</i> | RGC ₁ GC ₂ Y | | | 246 |
| <i>HaeIII</i> | | GG↓CC | 3(5) | ABGIMNPRU | 247,248,249 |
| | <i>Asp742I</i> | GGCC | | | 156 |
| | <i>Bce71I</i> | GGCC | | | 250 |
| | <i>BliI</i> | GGCC | | | 251 |
| | <i>BluII</i> | GGCC | | | 240 |
| | <i>BseI</i> | GGCC | | | 252 |
| | <i>BshAI</i> | GGCC | | | 83 |
| | <i>BshBI</i> | GGCC | | | 83 |
| | <i>BshCI</i> | GGCC | | | 83 |
| | <i>BshDI</i> | GGCC | | | 83 |
| | <i>BshEI</i> | GGCC | | | 83 |
| | <i>BshFI</i> | GGCC | | | 83 |
| | <i>BshI</i> | GGCC | | | 83 |
| | <i>Bsp71I</i> | GGCC | | | 250 |
| | <i>Bsp211I</i> | GG↓CC | | | 250 |
| | <i>Bsp226I</i> | GGCC | | | 250 |
| | <i>BspRI</i> | GG↓CC | | | 253-255 |
| | <i>BssCI</i> | GGCC | | | 117 |
| | <i>BstCI</i> | GGCC | | | 117 |
| | <i>BstJI</i> | GGCC | | | 56 |
| | <i>Bsu1076I</i> | GGCC | | | 102 |
| | <i>Bsu1114I</i> | GGCC | | | 102 |
| | <i>BsuRI</i> | GG↓CC | 3(5) | G | 248,256,257 |
| | <i>ClaI</i> | GGCC | | | 78 |
| | <i>CltI</i> | GG↓CC | | | 258 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|----------------------------|----------------------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| <i>Dsal</i> II | | GG↓CC | | | 187 |
| <i>FinS</i> I | | GGCC | | | 84 |
| <i>Fnu</i> DI | | GG↓CC | | | 227 |
| <i>Hhg</i> I | | GGCC | | | 55 |
| <i>Mnl</i> I | | GGCC | | | 213 |
| <i>Mnn</i> II | | GGCC | | | 259 |
| <i>Ngo</i> II | | GGCC | | | 260 |
| <i>Ngo</i> P ^{II} | | GG↓CC | | | 224 |
| <i>Ngo</i> S ^I | | GGCC | | | 49 |
| <i>Nla</i> I | | GGCC | | | 261 |
| <i>Pai</i> I | | GGCC | | | 107 |
| <i>Pal</i> I | | GG↓CC | R | | 262,263 |
| <i>Ppu</i> I | | GGCC | | | 107 |
| <i>Sfa</i> I | | GG↓CC | | | 264 |
| <i>Sph</i> III | | GGCC | | | 265 |
| <i>Sua</i> I | | GG↓CC | | | 266 |
| <i>Sul</i> I | | GGCC | | | 267 |
| <i>Tsp</i> ZNI | | GGCC | | | 268 |
| <i>Ttn</i> I | | GGCC | | | 82 |
| <i>Vha</i> I | | GGCC | | | 108 |
| <i>Hga</i> I | | GACGC(5/10) | N | | 181,245,269 |
| <i>Hgi</i> A ^I | | GWGCW↓C | N | | 270 |
| | <i>Asp</i> H ^{II} | GWGCW↓C | | | 479 |
| <i>Hgi</i> C ^I | | G↓GYRCC | | | 15,271 |
| | <i>Ban</i> I | G↓GYRCC | GIMNRU | | 8,271 |
| | <i>Eco</i> 50I | GGYRCC | | | 164 |
| | <i>Eco</i> 64I | GGYRCC | | | 191 |
| | <i>Hgi</i> H ^{II} | G↓GYRCC | | | 16 |
| <i>Hgi</i> E ^{II} | | ACCNNNNNNNGGT | | | 15 |
| <i>Hgi</i> J ^{II} | | GRGCY↓C | | | 16 |
| | <i>Ban</i> II | GRGCY↓C | GIMNRU | | 8 |
| | <i>Bvu</i> I | GRGCY↓C | | | 272 |
| | <i>Cfr</i> 48I | GRGCYC | | | 43 |
| | <i>Eco</i> 24I | GRGCYC | | | 210 |
| | <i>Eco</i> 25I | GRGCYC | | | 210 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|--------------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| | <i>Eco</i> 26I | GRGCYC | | | 44 |
| | <i>Eco</i> 35I | GRGCYC | | | 210 |
| | <i>Eco</i> 68I | GRGCYC | | | 192 |
| | <i>Eco</i> 113I | GRGCYC | | | 189 |
| | <i>Eco</i> T38I | GRGCYC | | | 91 |
| | <i>Kox</i> II | GRGCY↓C | | | 147 |
| <i>Hha</i> I | | GCG↓C | 2(5) | ABGNPRU | 273,274 |
| | <i>Bca</i> I | GCGC | | | 28 |
| | <i>Cfr</i> I | GCGC | | BIMP | 106 |
| | <i>Fnu</i> DIII | GCG↓C | | | 227 |
| | <i>Hin</i> GUI | GCGC | | | 237,275 |
| | <i>Hin</i> P1I | G↓CGC | | N | 276 |
| | <i>Hin</i> S1I | GCGC | | | 276 |
| | <i>Hin</i> S2I | GCGC | | | 276 |
| | <i>Mnn</i> IV | GCGC | | | 259 |
| | <i>Sci</i> NI | G↓CGC | | | 277 |
| <i>Hind</i> II | | GTY↓RAC | 5(6) | M | 278-281 |
| | <i>Chu</i> II | GTYRAC | | | 282 |
| | <i>Hin</i> 1160II | GTYRAC | | | 233 |
| | <i>Hin</i> 1161II | GTYRAC | | | 233 |
| | <i>Hin</i> JCI | GTY↓RAC | | | 283 |
| | <i>Hinc</i> II | GTY↓RAC | | ABGINPRU | 284 |
| | <i>Mnn</i> I | GTYRAC | | | 259 |
| <i>Hind</i> III | | A↓AGCTT | 1(6) | ABGIMNPRU | 280,281,285 |
| | <i>Asp</i> 52I | AAGCTT | | | 156 |
| | <i>Bbr</i> I | AAGCTT | | | 55 |
| | <i>Bpu</i> I | AAGCTT | | | 286,287 |
| | <i>Bst</i> FI | A↓AGCTT | | | 288 |
| | <i>Cfr</i> 32I | AAGCTT | | | 43 |
| | <i>Chu</i> I | AAGCTT | | | 282 |
| | <i>Eco</i> 65I | AAGCTT | | | 192 |
| | <i>Eco</i> 98I | AAGCTT | | | 192 |
| | <i>Eco</i> VIII | A↓AGCTT | | | 289 |
| | <i>Hin</i> 173I | AAGCTT | | | 237 |
| | <i>Hin</i> 1076III | AAGCTT | | | 233 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|------------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| <i>HinfI</i> | <i>HinJ</i> CII | AAGCTT | | | 283 |
| | <i>Hinb</i> III | AAGCTT | | | 55,290 |
| | <i>Hinf</i> II | AAGCTT | | | 291 |
| | <i>Hsu</i> I | A↓AGCTT | | | 55 |
| | <i>Mki</i> I | AAGCTT | | | 213 |
| | | G↓ANTC | | ABGIMNPRU | 290,292,293 |
| | <i>Cvi</i> BI | G↓ANTC | | | 294 |
| | <i>Cvi</i> CI | GANTC | | | 294 |
| | <i>Cvi</i> DI | GANTC | | | 294 |
| | <i>Cvi</i> EI | GANTC | | | 294 |
| | <i>Cvi</i> FI | GANTC | | | 294 |
| | <i>Cvi</i> GI | GANTC | | | 294 |
| | <i>Fnu</i> AI | G↓ANTC | | | 227 |
| <i>Hha</i> II | | G↓ANTC | 2(6) | | 295-297 |
| | <i>Nci</i> I | GANTC | | | 298 |
| | <i>Nov</i> II | GANTC | | | 298 |
| | <i>Nsi</i> HI | GANTC | | | 299 |
| | | GTT↓AAC | 5(6) | ABGIMNPRU | 300-302 |
| <i>Hpa</i> I | <i>Bse</i> II | GTAAAC | | | 252 |
| | | C↓CGG | 2(5) | BGMNPRU | 249,300,301 |
| | <i>Asp</i> 748I | CCGG | | | 156 |
| | <i>Bsu</i> 1192I | CCGG | | | 54,102 |
| | <i>Bsu</i> FI | CCGG | 1(5) | | 54,102,232 |
| | <i>Fin</i> II | CCGG | | | 130 |
| | <i>Hap</i> II | C↓CGG | | AGI | 245,303 |
| | <i>Mnl</i> II | CCGG | | | 213 |
| | <i>Mnol</i> | C↓CGG | | | 55,304 |
| | <i>Msp</i> I | C↓CGG | 1(5) | ABGIMNPRU | 111,305,306 |
| <i>Hph</i> I | <i>Sec</i> II | CCGG | | | 307 |
| | <i>Sfa</i> GUI | CCGG | | | 308 |
| | | GGTGA(8/7) | | N | 290,309 |
| | <i>Ngo</i> BI | GGTGA | -2(5) | | 49,480 |
| | | GGTAC↓C | | ABGIMNPRU | 310,311 |
| <i>Kpn</i> I | <i>Asp</i> 718I | G↓GTACC | | M | 312 |
| | <i>Eco</i> 149I | GGTACC | | | 190 |

| Enzyme ¹ | Isochizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|------------|
| | <i>Kpn</i> K14I | GGTACC | | | 156 |
| | <i>Nml</i> | GGTACC | | | 313 |
| | <i>Sth</i> A1 | GGTACC | | | 85 |
| | <i>Sth</i> B1 | GGTACC | | | 85 |
| | <i>Sth</i> C1 | GGTACC | | | 85 |
| | <i>Sth</i> D1 | GGTACC | | | 85 |
| | <i>Sth</i> E1 | GGTACC | | | 85 |
| | <i>Sth</i> F1 | GGTACC | | | 85 |
| | <i>Sth</i> G1 | GGTACC | | | 85 |
| | <i>Sth</i> H1 | GGTACC | | | 85 |
| | <i>Sth</i> I | G↓GTACC | | | 85 |
| | <i>Sth</i> J1 | GGTACC | | | 85 |
| | <i>Sth</i> K1 | GGTACC | | | 85 |
| | <i>Sth</i> L1 | GGTACC | | | 85 |
| | <i>Sth</i> M1 | GGTACC | | | 85 |
| | <i>Sth</i> N1 | GGTACC | | | 314 |
| <i>Ksp</i> 632I | | CTCTTC(1/4) | | | 478 |
| <i>Mae</i> I | | C↓TAG | M | | 315 |
| | <i>Mja</i> I | CTAG | | | 48 |
| <i>Mae</i> II | | A↓CGT | M | | 315 |
| <i>Mae</i> III | | ↓GTNAC | M | | 315 |
| <i>Mbo</i> I ⁶ | | ↓GATC | BGINR | | 316 |
| + | <i>Bce</i> 243I | ↓GATC | | | 317 |
| | <i>Bsa</i> PI | GATC | | | 117 |
| | <i>Bsp</i> 64I | GATC | | | 250 |
| + | <i>Bsp</i> 67I | ↓GATC | | | 250 |
| | <i>Bsp</i> 74I | GATC | | | 250 |
| | <i>Bsp</i> 76I | GATC | | | 250 |
| | <i>Bsp</i> 105I | ↓GATC | | | 250 |
| + | <i>Bsp</i> AI | ↓GATC | | | 39 |
| + | <i>Bsr</i> PII | GATC | | | 117 |
| | <i>Bss</i> GII | GATC | | | 117 |
| | <i>Bst</i> EIII | GATC | | | 55,139,318 |
| | <i>Bst</i> XII | GATC | | | 117 |
| | <i>Cpa</i> I | GATC | | | 254 |
| + | <i>Cpf</i> I | ↓GATC | | | 182 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| | <i>CviAI</i> | ↓GATC | 2(6) | | 319,320 |
| | <i>CviBII</i> | GATC | | | 321 |
| | <i>CviHI</i> | GATC | | | 170 |
| | <i>DpnII</i> | GATC | | | 174,176 |
| | <i>FnuAII</i> | GATC | | | 55,227 |
| | <i>FnuCI</i> | ↓GATC | | | 227 |
| + | <i>FnuEI</i> | ↓GATC | | | 227 |
| | <i>HaeI</i> | ↓GATC | | | 322 |
| | <i>MeuI</i> | GATC | | | 25 |
| | <i>MmeII</i> | GATC | | | 323 |
| | <i>MboIII</i> | GATC | | | 55 |
| | <i>MosI</i> | GATC | | | 316 |
| | <i>Msp67II</i> | GATC | | | 38 |
| | <i>MthI</i> | GATC | | | 117 |
| | <i>NdeII</i> | ↓GATC | B | | 298 |
| | <i>NflAII</i> | GATC | | | 223 |
| | <i>NflBI</i> | GATC | | | 324 |
| | <i>NflII</i> | GATC | | | 298 |
| | <i>NlaDI</i> | GATC | | | 49 |
| | <i>NlaII</i> | ↓GATC | | | 261 |
| | <i>NmeCI</i> | ↓GATC | | | 224 |
| | <i>NphI</i> | ↓GATC | | | 224 |
| | <i>NsiAI</i> | GATC | | | 325 |
| | <i>NspAI</i> | GATC | | | 25 |
| | <i>NsuI</i> | GATC | | | 50 |
| | <i>PflI</i> | GATC | | | 305 |
| | <i>SalAI</i> | GATC | | | 196 |
| | <i>SalHI</i> | GATC | | | 196 |
| + | <i>Sau3AI</i> | ↓GATC | 4(5) | ABGIMNPRU | 326,327 |
| | <i>Sau6782I</i> | GATC | | | 328 |
| | <i>SinMI</i> | GATC | | | 50 |
| | <i>TruII</i> | GATC | | | 88 |
| <i>MboII</i> | | GAAGA(8/7) | 5(6) | BGINR | 316,329-331 |
| | <i>NcuI</i> | GAAGA | | | 332 |
| | <i>TceI</i> | GAAGA | | | 267 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| <i>Mlu</i> I | | A↓CGCGT | | ABGIMNPRU | 236 |
| | <i>Ape</i> I | ACGCGT | | | 196 |
| <i>Mme</i> I | | TCCRAC(20/18) | | | 323 |
| <i>Mn</i> II | | CCTC(7/7) | | GN | 111,333 |
| <i>Mse</i> I | | T↓TAA | | N | 130 |
| <i>Mst</i> I | | TCC↓GCA | | | 110,334 |
| | <i>Aoe</i> I | TGC↓GCA | G | | 12 |
| | <i>Avi</i> II | TGC↓GCA | | | 87 |
| | <i>Fdi</i> II | TGC↓GCA | | | 81,82 |
| | <i>Fsp</i> I | TGC↓GCA | | GN | 57,335 |
| | <i>Gsp</i> AII | TGCGCA | | | 83 |
| <i>Nae</i> I | | GCC↓GGC | | GMNU | 336 |
| | <i>Ame</i> II | GCCGGC | | | 27 |
| | <i>Ani</i> MI | GCCGGC | | | 130 |
| | <i>Ape</i> AI | GCCGGC | | | 27 |
| | <i>Apr</i> I | GCCGGC | | | 25 |
| | <i>Eco</i> 56I | GCCGGC | | | 44 |
| | <i>Mis</i> I | GCCGGC | | | 139 |
| | <i>Nas</i> WI | GCCGGC | | | 25 |
| | <i>Nba</i> I | GCCGGC | | | 54 |
| | <i>Nbr</i> I | GCCGGC | | | 54 |
| | <i>Ngo</i> MI | GCCGGC | | | 337 |
| | <i>Nmu</i> FI | GCCGGC | | | 50 |
| | <i>Nmu</i> I | GCCGGC | | | 298 |
| | <i>Nsp</i> WI | GCCGGC | | | 25 |
| | <i>Ntb</i> SII | GCCGGC | | | 84 |
| | <i>Pgl</i> II | GCCGGC | | | 338 |
| | <i>Psp</i> 61I | GCCGGC | | | 38 |
| | <i>Rlu</i> I | GCCGGC | | | 339-341 |
| | <i>Sac</i> AI | GCCGGC | | | 45 |
| | <i>Sal</i> CI | GCCGGC | | | 27 |
| | <i>Sao</i> I | GCCGGC | | | 342 |
| | <i>Sau</i> AI | GCCGGC | | | 324 |
| | <i>Sau</i> BMKI | GCC↓GGC | | | 343 |
| | <i>Sal</i> I | GCCGGC | | | 50 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| <i>NarI</i> | | GG↓CGCC | | BGN | 344 |
| | <i>BbeAI</i> | GGCGCC | | | 131 |
| | <i>BbeI</i> | GGCGC↓C | | A | 345 |
| | <i>BinSII</i> | GGCGCC | | | 131 |
| | <i>Eco78I</i> | GGC↓GCC | | | 346 |
| | <i>NamI</i> | GGCGCC | | | 347 |
| | <i>NdaI</i> | GG↓CGCC | | | 348 |
| | <i>NunII</i> | GG↓CGCC | | G | 117 |
| | <i>SfoI</i> | GGCGCC | | | 130 |
| <i>NcoI</i> | | C↓CATGG | | ABGMNPR | 117 |
| | <i>NspSAIII</i> | CCATGG | | | 66 |
| <i>NdeI</i> | | CA↓TATG | | BNR | 349 |
| <i>NheI</i> | | G↓CTAGC | | BGMNPR | 313 |
| <i>NlaIII</i> | | CATG↓ | | N | 261 |
| <i>NlaIV</i> | | GGN↓NCC | | N | 261 |
| <i>BcrI</i> | | GGNNCC | | | 193 |
| | | GC↓GGCCGC | | AGMNPRU | 350,351 |
| | | TCG↓CGA | | ABGMNRU | 313 |
| <i>NruI</i> | <i>AmI</i> | TCGCGA | | | 108 |
| | <i>SalDI</i> | TCGCGA | | | 27 |
| | <i>Sbo13I</i> | TCG↓CGA | | | 91 |
| <i>NspI</i> | | RCATG↓Y | | A | 6 |
| | <i>NspHI</i> | RCATG↓Y | | G | 59 |
| <i>NspBII</i> | | CMG↓CKG | | G | 59 |
| <i>PflMI</i> | | CCANNNNN↓NTGG | | N | 130 |
| <i>PleI</i> | | GAGTC(4/5) | | N | 352 |
| <i>PmaCI</i> | | CAC↓GTG | | | 353 |
| <i>Eco72I</i> | | CAC↓GTG | | | 354 |
| | | RG↓GWCCY | | N | 130,355 |
| <i>PpuMI</i> | | CTGCA↓G | 5(6) | ABGMNPRU | 332,356,357 |
| <i>PstI</i> | | CTGCA↓G | | | |
| <i>Alu2882I</i> | | CTGCAG | | | 100 |
| <i>AluAJI</i> | | CTGCA↓G | | | 358 |
| <i>Asp36I</i> | | CTGCAG | | | 210 |
| <i>Asp708I</i> | | CTGCAG | | | 69 |
| <i>BbvI</i> | | CTGCAG | | | 14 |

| Enzyme ¹ | Isochizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|--------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| <i>Bce</i> 170I | | CTGCAG | | | 102 |
| <i>Bsp</i> 63I | | CTGCA↓G | | | 250 |
| <i>Bsp</i> 78I | | CTGCAG | | | 250 |
| <i>Bsp</i> BI | | CTGCA↓G | | | 39 |
| <i>Bsu</i> BI | | CTGCAG | | | 102,359 |
| <i>Cau</i> III | | CTGCAG | | | 360 |
| <i>Cfl</i> I | | CTGCA↓G | | | 322 |
| <i>Cfr</i> A4I | | CTGCA↓G | | | 361 |
| <i>Eae</i> PI | | CTGCAG | | | 229 |
| <i>Ecl</i> 77I | | CTGCAG | | | 192 |
| <i>Ecl</i> 593I | | CTGCAG | | | 156 |
| <i>Eco</i> 48I | | CTGCAG | | | 164 |
| <i>Eco</i> 49I | | CTGCAG | | | 164 |
| <i>Eco</i> 83I | | CTGCAG | | | 192 |
| <i>Eco</i> 133I | | CTGCAG | | | 190 |
| <i>Eco</i> 141I | | CTGCAG | | | 189 |
| <i>Eco</i> 161I | | CTGCAG | | | 190 |
| <i>Eco</i> 167I | | CTGCAG | | | 190 |
| <i>Mau</i> I | | CTGCAG | | | 107 |
| <i>Mkr</i> I | | CTGCAG | | | 25 |
| <i>Nas</i> I | | CTGCAG | | | 84 |
| <i>Ngbl</i> | | CTGCAG | | | 25 |
| <i>Noc</i> I | | CTGCAG | | | 313 |
| <i>Pma</i> 44I | | CTGCA↓G | | | 361 |
| <i>Pma</i> I | | CTGCAG | | | 111 |
| <i>Pmy</i> I | | CTGCAG | | | 362 |
| <i>Sa</i> PI | | CTGCA↓G | | | 363,364 |
| <i>Sf</i> II | | CTGCA↓G | | | 213 |
| <i>Sk</i> II | | CTGCAG | | | 50 |
| <i>Xma</i> II | | CTGCAG | | | 365 |
| <i>Xor</i> I | | CTGCAG | | | 366 |
| <i>Xph</i> I | | CTGCAG | | | 367 |
| <i>Yen</i> AI | | CTGCAG | | | 368 |
| <i>Yen</i> BI | | CTGCAG | | | 368 |
| <i>Yen</i> CI | | CTGCAG | | | 368 |

| Enzyme ¹ | Iisoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|------------------|--------------------------------------|-------------------------|-----------------------------------|---------------|
| <i>Pvu</i> I | <i>Yen</i> DI | CTGCAG | | | 368 |
| | <i>Yen</i> EI | CTGCAG | | | 368 |
| | <i>Yen</i> I | CTGCA↓G | | | 368 |
| | | CGAT↓CG | | ABGMNPRU | 369 |
| | <i>Bma</i> AI | CGATCG | | | 27 |
| | <i>Bma</i> BI | CGATCG | | | 27 |
| | <i>Bma</i> CI | CGATCG | | | 27 |
| | <i>Bma</i> DI | CGATCG | | | 27 |
| | <i>Bma</i> I | CGATCG | | | 28 |
| | <i>Ecl</i> JI | CGAT↓CG | | | 370 |
| <i>Pvu</i> II | <i>Nb</i> II | CGAT↓CG | | | 111 |
| | <i>Rsh</i> I | CGAT↓CG | | | 371 |
| | <i>Rsp</i> I | CGATCG | | | 372 |
| | <i>Xn</i> I | CGATCG | | | 259 |
| | <i>Xor</i> II | CGAT↓CG | B | | 366,369 |
| | | CAG↓CTG | 4(4) | ABCIMNPRU | 369,373 |
| | <i>Bav</i> I | CAG↓CTG | | | 374 |
| | <i>Cfr</i> 6I | CAG↓CTG | 4(4) | | 41,44,373,375 |
| | <i>Mz</i> I | CAGCTG | | | 376 |
| | <i>Rsa</i> I | GT↓AC | | ABCIMNPRU | 377 |
| <i>Rsr</i> II | <i>Cvi</i> QI | G↓TAC | | | 7 |
| | | CG↓GWCCG | | GN | 378 |
| <i>Sac</i> I | <i>Cpo</i> I | CGGWCCG | | | 27 |
| | <i>Sac</i> I | GAGCT↓C | | AGIMNPRU | 379 |
| | <i>Eco</i> 136II | GAGCTC | | | 190 |
| | <i>Eco</i> ICRI | GAGCTC | | | 107 |
| | <i>Nas</i> SI | GAGCTC | | | 84 |
| | <i>Sco</i> I | GAGCTC | | | 342 |
| | <i>Sst</i> I | GAGCT↓C | B | | 380,381 |
| | | CCGC↓GG | | GINPRU | 379 |
| | <i>Bac</i> I | CCGGGG | | | 55,258 |
| | <i>Cfr</i> 37I | CCGGGG | | | 42 |
| <i>Sac</i> II | <i>Cfr</i> 41I | CCGGGG | | | 42 |
| | <i>Cfr</i> 42I | CCGGGG | | | 43 |
| | <i>Cfr</i> 43I | CCGGGG | | | 43 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|---------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| <i>Cfr</i> 45II | | CCGC ₄ GG | | | 43 |
| <i>Csc</i> I | | CCGC↓GG | | | 382 |
| <i>Eco</i> I | | CCGC ₄ GG | | | 55,383 |
| <i>Ecl</i> 28I | | CCGC ₄ GG | | | 210 |
| <i>Ecl</i> 37I | | CCGC ₄ GG | | | 46 |
| <i>Eco</i> 55I | | CCGC ₄ GG | | | 164 |
| <i>Eco</i> 92I | | CCGC ₄ GG | | | 192 |
| <i>Eco</i> 96I | | CCGC ₄ GG | | | 192 |
| <i>Eco</i> 99I | | CCGC ₄ GG | | | 192 |
| <i>Eco</i> 100I | | CCGC ₄ GG | | | 189 |
| <i>Eco</i> 104I | | CCGC ₄ GG | | | 212 |
| <i>Eco</i> 134I | | CCGC ₄ GG | | | 190 |
| <i>Eco</i> 135I | | CCGC ₄ GG | | | 190 |
| <i>Eco</i> 158I | | CCGC ₄ GG | | | 189 |
| <i>Ga</i> I | | CCGC↓GG | | | 322 |
| <i>Gce</i> GLI | | CCGC↓GG | | | 384 |
| <i>Gce</i> I | | CCGC↓GG | | | 322 |
| <i>Mra</i> I | | CCGC ₄ GG | | | 385 |
| <i>Ngo</i> DI | | CCGC ₄ GG | | | 180 |
| <i>Ngo</i> III | | CCGC ₄ GG | | | 386 |
| <i>Ngo</i> PIII | | CCGC↓GG | | | 224 |
| <i>Nla</i> DIII | | CCGC ₄ GG | | | 49 |
| <i>Nla</i> SI | | CCGC ₄ GG | | | 17 |
| <i>Pac</i> AI | | CCGC↓GG | | | 387 |
| <i>Saa</i> I | | CCGC ₄ GG | | | 130 |
| <i>Sab</i> I | | CCGC ₄ GG | | | 342 |
| <i>Sak</i> I | | CCGC ₄ GG | | | 45 |
| <i>Sbo</i> I | | CCGC ₄ GG | | | 388,389 |
| <i>Sfr</i> I | | CCGC ₄ GG | | | 388,389 |
| <i>Shy</i> I | | CCGC ₄ GG | | | 390 |
| <i>Sst</i> II | | CCGC↓GG | B | | 380 |
| <i>Tg</i> II | | CCGC ₄ GG | | | 109 |
| <i>Sai</i> I | | G↓TCGAC | | ABCIMNPRU | 391 |
| <i>Hgi</i> CIII | | G↓TCGAC | | | 15 |
| <i>Hgi</i> DII | | G↓TCGAC | | | 15 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|------------|
| | <i>NopI</i> | G↓TCGAC | | | 111 |
| | <i>RheI</i> | GTCGAC | | | 108 |
| | <i>RhpI</i> | GTCGAC | | | 108 |
| | <i>RrhI</i> | GTCGAC | | | 54 |
| | <i>RroI</i> | GTCGAC | | | 54 |
| | <i>XamI</i> | GTCGAC | | | 391 |
| | <i>XciI</i> | G↓TCGAC | | | 392 |
| <i>SauI</i> | | CC↓TNAGG | M | | 393 |
| | <i>AacI</i> | CC↓TNAGG | | | 87 |
| | <i>AxyI</i> | CC↓TNAGG | G | | 394 |
| | <i>Bsu36I</i> | CC↓TNAGG | N | | 395 |
| | <i>CvnI</i> | CC↓TNAGG | B | | 396,397 |
| | <i>Eco76I</i> | CCTNAGG | | | 192 |
| | <i>Eco81I</i> | CC↓TNAGG | AU | | 398 |
| | <i>Eco115I</i> | CCTNAGG | | | 189 |
| | <i>Eco118I</i> | CCTNAGG | | | 189 |
| | <i>MstII</i> | CC↓TNAGG | | | 111 |
| | <i>OxaNI</i> | CC↓TNAGG | | | 112 |
| | <i>SecIII</i> | CCTNAGG | | | 307 |
| <i>Scal</i> | | AGT↓ACT | ABGMNPRU | | 399,400 |
| | <i>Asp763I</i> | AGTACT | | | 156 |
| | <i>BstMI</i> | AGTACT | | | 56 |
| <i>ScrFI</i> | | CC↓NGG | N | | 401 |
| | <i>Eco43I</i> | CCNGG | | | 402 |
| | <i>Eco51II</i> | CCNGG | | | 164 |
| | <i>Eco80I</i> | CCNGG | | | 192 |
| | <i>Eco85I</i> | CCNGG | | | 402 |
| | <i>Eco93I</i> | CCNGG | | | 192 |
| | <i>Eco153I</i> | CCNGG | | | 189 |
| | <i>Msp67I</i> | CC↓NGG | | | 38 |
| | <i>SsoII</i> | ↓CCNGG | 2(5) | | 202,403 |
| <i>SduI</i> | | GDGCH↓C | G | | 404,405 |
| | <i>AocII</i> | GDGCH↓C | | | 87 |
| | <i>Bsp1286I</i> | GDGCH↓C | N | | 55,102,406 |
| | <i>NspII</i> | GDGCH↓C | R | | 6 |

| Enzyme ¹ | Isochizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|-----------------|--------------------------------------|-------------------------|-----------------------------------|---------------|
| <i>Secl</i> | | C↓CNNGG | | | 307 |
| <i>SfaNI</i> | | GCATC(5/9) | | N | 115 |
| <i>SfiI</i> | | GGCCNNNN↓NGGCC | | GNPRU | 407 |
| <i>SmaI</i> | | CCC↓GGG | 3(5) | ABGMNPRU | 365,408,409 |
| | <i>Cfr9I</i> | C↓CCCCG | 2(4) | | 41,44,375,410 |
| | <i>XbaI</i> | C↓CCCCG | | R | 411 |
| | <i>XmaI</i> | C↓CCCCG | | INP | 365 |
| <i>SnaBI</i> | | TAC↓GTA | | GMN | 412 |
| | <i>Eco105I</i> | TAC↓GTA | | | 413 |
| | <i>Eco158II</i> | TACGTA | | | 189 |
| <i>SnaI</i> | | GTATAC | | | 414 |
| | <i>XcaI</i> | GTA↓TAC | | | 415 |
| <i>SpeI</i> | | A↓CTAGT | | BMN | 313 |
| <i>SphI</i> | | GCATG↓C | | ABGMNPRU | 416 |
| | <i>PaeI</i> | GCATG↓C | | | 417 |
| | <i>SpaXI</i> | GCATGC | | | 107 |
| <i>SpII</i> | | C↓GTACG | | A | 265 |
| | <i>PfuI</i> | CGTACG | | | 27 |
| <i>SspI</i> | | AAT↓ATT | | BMN | 400 |
| <i>StuI</i> | | AGG↓CCT | | ABGMNPR | 418 |
| | <i>AatI</i> | AGG↓CCT | | IU | 8 |
| | <i>Asp78I</i> | AGGCCT | | | 156 |
| | <i>ChyI</i> | AGGCCT | | | 28 |
| | <i>Eco147I</i> | AGGCCT | | | 137 |
| | <i>GdiI</i> | AGG↓CCT | | | 240 |
| | <i>NtaSI</i> | AGGCCT | | | 84 |
| <i>StyI</i> | | C↓CWG | | BGNP | 419 |
| | <i>Eco130I</i> | CCWWGG | | | 137 |
| | <i>EcoT14I</i> | C↓CWG | | A | 420 |
| | <i>EcoT104I</i> | CCWWGG | | | 420 |
| | <i>SbaI</i> | CCWWGG | | | 421 |
| | <i>SbIBI</i> | CCWWGG | | | 421 |
| | <i>SbICl</i> | CCWWGG | | | 421 |
| <i>TaqI</i> | | T↓CGA | 4(6) | BGIMNPRU | 422,423 |
| | <i>CviBIII</i> | TCGA | | | 321 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------------|----------------|--------------------------------------|-------------------------|-----------------------------------|-----------|
| | <i>TflI</i> | TCGA | | | 423 |
| | <i>TthHB8I</i> | T↓CGA | 4(6) | A | 423-426 |
| <i>TaqII</i> ⁷ | | GACCGA(11/9) | | | 55,427 |
| <i>TaqII</i> ⁷ | | CACCCA(11/9) | | | |
| <i>Tsp45I</i> | | GTSAC | | | 428 |
| <i>TspEI</i> | | AATT | | | 429 |
| <i>Tth111I</i> | | GACN↓NNGTC | | AGINR | 310 |
| | <i>FsuI</i> | GACNNNGTC | | | 342 |
| | <i>NtaI</i> | GACNNNGTC | | | 84 |
| | <i>SplII</i> | GACNNNGTC | | | 265 |
| | <i>TteI</i> | GACNNNGTC | | | 310 |
| | <i>TtrI</i> | GACNNNGTC | | | 310 |
| <i>Tth111II</i> | | CAARCA(11/9) | | | 430 |
| <i>VspI</i> | | AT↓TAAT | | | 431 |
| | <i>AseI</i> | AT↓TAAT | | N | 150 |
| <i>XbaI</i> | | T↓CTAGA | | ABGIMNPRU | 432 |
| <i>XhoI</i> | | C↓TCGAG | | ABGIMNPRU | 433 |
| | <i>AbrI</i> | C↓TCGAG | | | 434 |
| | <i>Asp47I</i> | CTCGAG | | | 156 |
| | <i>Asp703I</i> | CTCGAG | | | 69 |
| | <i>BbvIII</i> | CTCGAG | | | 14 |
| | <i>BluI</i> | C↓TCGAG | | | 433 |
| | <i>BssHII</i> | CTCGAG | | | 117 |
| | <i>BstHI</i> | CTCGAG | | | 117 |
| | <i>BstLI</i> | CTCGAG | | | 56 |
| | <i>BstVI</i> | CTCGAG | | | 435 |
| | <i>BsuMI</i> | CTCGAG | 3(5) | | 102,232 |
| | <i>BthI</i> | CTCGAG | | | 131 |
| | <i>CcrI</i> | C↓TCGAG | | | 436 |
| | <i>DdeII</i> | CTCGAG | | | 55 |
| | <i>McaI</i> | CTCGAG | | | 69 |
| | <i>MecI</i> | CTCGAG | | | 376 |
| | <i>MpuI</i> | CTCGAG | | | 376 |
| | <i>MsiI</i> | CTCGAG | | | 55,290 |
| | <i>PaeR7I</i> | C↓TCGAG | 5(6) | N | 437,438 |

| Enzyme ¹ | Isoschizomers | Recognition ² Sequence | Me ³ site | Commercial ⁴ source | Reference |
|---------------------|------------------|--------------------------------------|-------------------------|-----------------------------------|-------------|
| | <i>Pan</i> I | C↓TCGAG | | | 107 |
| | <i>Pfl</i> NI | CTCGAG | | | 28 |
| | <i>Pfl</i> WI | CTCGAG | | | 439 |
| | <i>Sau</i> 3239I | C↓TCGAG | | | 440,441 |
| | <i>ScI</i> | CTC↓GAG | | | 38 |
| | <i>Scu</i> I | CTCGAG | | | 388 |
| | <i>Ser</i> I | CTCGAG | | | 388 |
| | <i>Sga</i> I | CTCGAG | | | 388 |
| | <i>Sgo</i> I | CTCGAG | | | 388 |
| | <i>Sla</i> I | C↓TCGAG | | | 442 |
| | <i>Slu</i> I | CTCGAG | | | 389 |
| | <i>Spa</i> I | CTCGAG | | | 388 |
| | <i>Xpa</i> I | C↓TCGAG | | | 433 |
| <i>Xba</i> II | | R↓GATCY | | GM | 109,233,443 |
| | <i>Ait</i> AI | RGATCY | | | 27 |
| | <i>Ait</i> II | RGATCY | | | 193 |
| | <i>Bst</i> YI | R↓GATCY | | N | 56 |
| | <i>Mf</i> I | R↓GATCY | | A | 444 |
| <i>Xma</i> III | | C↓GGCCG | 4(5) | B | 32,445 |
| | <i>Bst</i> ZI | CGGCCG | | | 56 |
| | <i>Eag</i> I | C↓GGCCG | | N | 446 |
| | <i>Eco</i> 52I | C↓GGCCG | | AU | 34,80 |
| <i>Xmn</i> I | | GAANN↓NNNTTC | | GN | 54,447 |
| | <i>Asp</i> 700I | GAANN↓NNNTTC | | M | 69 |

Type I enzymes

| Enzyme | Recognition sequence | Me site ³ | Reference |
|-----------------|-------------------------|----------------------|-----------|
| <i>Eco</i> AI | GAGNNNNNNNGTCA | 2(6) -3(6) | 448,449 |
| <i>Eco</i> BI | TGANNNNNNNNNTGCT | 3(6) | 450-454 |
| <i>Eco</i> DI | TTANNNNNNNGTCY | | 455 |
| <i>Eco</i> DXXI | TCANNNNNNNNATTCA | | 456,457 |
| <i>Eco</i> KI | AACNNNNNNNGTGC | | 458-461 |

| Enzyme | Recognition sequence | Me site ³ | Reference |
|-------------------|----------------------|----------------------|-----------|
| <i>EcoR124I</i> | GAANNNNNNRTCG | | 462 |
| <i>EcoR124/3I</i> | GAANNNNNNNRTCG | -3(6) | 462 |
| <i>StySBI</i> | GAGNNNNNNRTAYG | 2(6) -4(6) | 463 |
| <i>StySPI</i> | AACNNNNNNCTRC | 2(6) -3(6) | 463 |
| <i>StySQI</i> | AACNNNNNNRTAYG | | 464 |

Type III enzymes

| Enzyme | Isoschizomers | Recognition Sequence | Me ³ site | Reference |
|----------------|---------------|----------------------|----------------------|-----------|
| <i>EcoP15I</i> | | CAGCAG | | 465,466 |
| <i>EcoPI</i> | | AGACC | 3(6) | 467-471 |
| <i>HinfII</i> | | CGAAT | | 472,473 |
| <i>HincI</i> | | CGAAT | | 474 |

FOOTNOTES

- * signifies that *DpnI* and its isoschizomers require the presence of 6-methyladenosine within the recognition sequence GATC.
- Recognition sequences are given using the standard abbreviations (475) 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. 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 (4/87)
 - B Bethesda Research Laboratories (9/87)
 - G Anglian Biotechnology Ltd. (1/88)
 - I International Biotechnologies Inc (7/87)
 - M Boehringer-Mannheim (3/88)
 - N New England Biolabs (3/88)
 - P Pharmacia P-L Biochemicals (3/88)
 - R Promega Biotec (9/87)
 - U United States Biochemical Corporation (2/88)
5. *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 +.
6. *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 +.
7. *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

| | | |
|---------------------------------|-----------------------------------|-----------------------------------|
| <i>AacI</i> (<i>BamHI</i>) | <i>AaeI</i> (<i>BamHI</i>) | <i>AatI</i> (<i>StuI</i>) |
| <i>AaiII</i> | <i>AbI</i> (<i>XbaI</i>) | <i>AccEBI</i> (<i>BamHI</i>) |
| <i>AccI</i> | <i>AccII</i> (<i>FnuDII</i>) | <i>AccIII</i> (<i>BspMII</i>) |
| <i>AcyI</i> | <i>AeuI</i> (<i>EcoRII</i>) | <i>AflI</i> (<i>AvaII</i>) |
| <i>AfII</i> | <i>AfIII</i> | <i>AhaI</i> (<i>CauII</i>) |
| <i>AhaII</i> (<i>AcyI</i>) | <i>AhaIII</i> | <i>AitAI</i> (<i>XbaII</i>) |
| <i>AiiI</i> (<i>Eco47III</i>) | <i>AiiII</i> (<i>XbaII</i>) | <i>AiiI</i> (<i>BamHI</i>) |
| <i>Ali2882I</i> (<i>PstI</i>) | <i>Ali12257I</i> (<i>BamHI</i>) | <i>Ali12258I</i> (<i>BamHI</i>) |
| <i>AlaJAI</i> (<i>PstI</i>) | <i>AluI</i> | <i>AlwI</i> (<i>BinI</i>) |
| <i>AlwNI</i> | <i>AlwXI</i> (<i>BbvI</i>) | <i>Amal</i> (<i>NruI</i>) |
| <i>Amel</i> (<i>ApaLI</i>) | <i>Amel</i> (<i>Nael</i>) | <i>AniMI</i> (<i>Nael</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>Acyl</i>) | <i>Apal</i> |
| <i>ApaLI</i> | <i>Apel</i> (<i>MluI</i>) | <i>ApeAI</i> (<i>Nael</i>) |
| <i>AprI</i> (<i>Nael</i>) | <i>Apul</i> (<i>AsuI</i>) | <i>ApyI</i> (<i>EcoRII</i>) |

| | | |
|-------------------|-------------------|--------------------|
| AquI (AvaI) | AseI (VspI) | AselII (CauII) |
| Asp36I (PstI) | Asp47I (Xhol) | Asp52I (HindIII) |
| Asp78I (StuI) | Asp697I (AvaiII) | Asp700I (XmnI) |
| Asp703I (Xhol) | Asp707I (ClaI) | Asp708I (PstI) |
| Asp718I (KpnI) | Asp742I (HaeIII) | Asp748I (HpaII) |
| Asp763I (ScaI) | AspAI (BstEII) | AspHI (HgiAI) |
| AsiWI (AcyI) | AsuI | AsuII |
| AsuIII (AcyI) | AtuII (EcoRII) | AtuBI (EcoRII) |
| AtuCI (BclI) | AvAI | AvaiII |
| AvaiIII | AviII (MstI) | AvrI (AvaI) |
| AvrII | AryI (SauI) | Bac36I (AsuI) |
| BacI (SacII) | BalI | BamFI (BamHI) |
| BamHI | BamKI (BamHI) | BamNI (BamHI) |
| BamNxI (AvaiII) | BanI (HgiCI) | BanII (HgiJI) |
| BanIII (ClaI) | BavI (PvuII) | BbeAI (NarI) |
| Bbl (NarI) | BbiI (PstI) | BbiII (AcyI) |
| BbrII (Xhol) | BbrI (HindIII) | BbvI |
| BbvII | BcaI (Hhal) | Bce71I (HaeIII) |
| Bce170I (PstI) | Bce243I (MboI) | BceFI (FnuDII) |
| BceRI (FnuDII) | BclI | BcmI (ClaI) |
| BcnI (CauII) | BcrI (NlaIV) | BepI (FnuDII) |
| BglI | BgI | BinI |
| BinSI (EcoRII) | BinSI (NarI) | BliI (HaeIII) |
| BluI (Xhol) | BluII (HaeIII) | BmaI (PvuI) |
| BmaAI (PvuI) | BmaBI (PvuI) | BmaCI (PvuI) |
| BmaDI (PvuI) | Bme216I (AvaiII) | BpeI (HindIII) |
| BsaPI (MboI) | BscI (ClaI) | BseI (HaeIII) |
| BseII (HpaI) | BsePI | BshI (HaeIII) |
| BshAI (HaeIII) | BshBI (HaeIII) | BshCI (HaeIII) |
| BshDI (HaeIII) | BshEI (HaeIII) | BshFI (HaeIII) |
| BsmI | BsmAI | BsoPI (BsePI) |
| Bsp63I (PstI) | Bsp64I (MboI) | Bsp67I (MboI) |
| Bsp71I (HaeIII) | Bsp74I (MboI) | Bsp76I (MboI) |
| Bsp78I (PstI) | Bsp105I (MboI) | Bsp106I (ClaI) |
| Bsp211I (HaeIII) | Bsp226I (HaeIII) | Bsp1286I (SdU) |
| BspAI (MboI) | BspBI (PstI) | BspBII (AsuI) |
| BspHI | BspMI | BspMII |
| BspRI (HaeIII) | BspXI (ClaI) | BspXII (BclI) |
| BsrI | BsrHI (BsePI) | BsrPII (MboI) |
| BssCI (HaeIII) | BssGI (BstXI) | BssGII (MboI) |
| BssHI (Xhol) | BssHII (BsePI) | BstII (BamHI) |
| Bst31I (BstEII) | BstBI (AsuII) | BstCI (HaeIII) |
| BstDI (BstEII) | BstEII | BstEIII (MboI) |
| BstFI (HindIII) | BstGI (BclI) | BstGII (EcoRII) |
| BstHI (Xhol) | BstII (HaeIII) | BstKI (BclI) |
| BstLI (Xhol) | BstMI (ScaI) | BstNI (EcoRII) |
| BstOI (EcoRII) | BstPI (BstEII) | BstQI (BamHI) |
| BstRI (EcoRV) | BstSI (AvaI) | BstTI (BstXI) |
| BstUI (FnuDII) | BstVI (Xhol) | BstWI (EcoNI) |
| BstXI | BstXII (MboI) | BstYI (Xhol) |
| BstZI (XmaIII) | Bsu36I (SauI) | Bsu1076I (HaeIII) |
| Bsu1114I (HaeIII) | Bsu1192I (HpaII) | Bsu1192II (FnuDII) |
| Bsu1193I (FnuDII) | Bsu6633I (FnuDII) | BsuBI (PstI) |
| BsuEII (FnuDII) | BsuFI (HpaII) | BsuMI (Xhol) |
| BsuRI (HaeIII) | BthI (Xhol) | BthII (BlnI) |
| BtlI (AvaiII) | BvuI (HgiJI) | CauI (AvaiII) |
| CauII | CauIII (PstI) | CcrI (Xhol) |

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| Cdi27I (EcoRII) | CelII (BamHI) | CelII (EspI) |
| Ceq1 (EcoRV) | CflI (PstI) | CfoI (HhaI) |
| Cfr1 | Cfr4I (AsuI) | Cfr5I (EcoRII) |
| Cfr6I (PvuII) | Cfr7I (BstEII) | Cfr8I (AsuI) |
| Cfr9I (SmaI) | Cfr10I | Cfr11I (EcoRII) |
| Cfr13I (AsuI) | Cfr14I (CfrI) | Cfr19I (BstEII) |
| Cfr20I (EcoRII) | Cfr22I (EcoRII) | Cfr23I (AsuI) |
| Cfr24I (EcoRII) | Cfr25I (EcoRII) | Cfr27I (EcoRII) |
| Cfr28I (EcoRII) | Cfr29I (EcoRII) | Cfr30I (EcoRII) |
| Cfr31I (EcoRII) | Cfr32I (HindIII) | Cfr33I (AsuI) |
| Cfr35I (EcoRII) | Cfr37I (SacII) | Cfr38I (CfrI) |
| Cfr39I (CfrI) | Cfr40I (CfrI) | Cfr41I (SacII) |
| Cfr42I (SacII) | Cfr43I (SacII) | Cfr45I (AsuI) |
| Cfr45II (SacII) | Cfr46I (AsuI) | Cfr47I (AsuI) |
| Cfr48I (HgiJII) | CfrA4I (PstI) | CfrNI (AsuI) |
| Cfrs37I (EcoRII) | CfuI (DpnI) | ChuI (HindIII) |
| ChuII (HindII) | ChyI (StuI) | ClaI |
| CImI (HaeIII) | CImII (AvaiI) | CltI (HaeIII) |
| CpaI (MboI) | CpeI (BclI) | CpfI (MboI) |
| CpoI (RsrII) | CscI (SacII) | CthI (BclI) |
| CthII (EcoRII) | CviAI (MboI) | CviBI (HinfI) |
| CviBII (MboI) | CviBIII (TaqI) | CviCI (HinfI) |
| CviDI (HinfI) | CviEI (HinfI) | CviFI (HinfI) |
| CviGI (HinfI) | CviHI (MboI) | CviJI |
| CviKI (CviJI) | CviLI (CviJI) | CviMI (CviJI) |
| CviNI (CviJI) | CviOI (CviJI) | CviQI (RsaI) |
| CvnI (SauI) | DdeI | DdeII (XbaI) |
| DdsI (BamHI) | DpnI | DpnII (MboI) |
| DraI (AhaIII) | DraII | DraIII |
| DsaI | DsaII (HaeIII) | EaeI (CfrI) |
| EaePI (PstI) | EagI (XmaIII) | EagKI (EcoRII) |
| EagMI (AvaiI) | EarI | EcaI (BstEII) |
| EcaII (EcoRII) | EccI (SacII) | EclII (EcoRII) |
| Ecl28I (SacII) | Ecl37I (SacII) | Ecl66I (EcoRII) |
| Ecl77I (PstI) | Ecl593I (PstI) | EclJII (PvuI) |
| Ecs39I (EcoRII) | Eco24I (HgiJII) | Eco25I (HgiJII) |
| Eco26I (HgiJII) | Eco31I | Eco32I (EcoRV) |
| Eco35I (HgiJII) | Eco38I (EcoRII) | Eco39I (AsuI) |
| Eco40I (EcoRII) | Eco41I (EcoRII) | Eco42I (Eco31I) |
| Eco43I (ScrFI) | Eco47I (AvaiI) | Eco47II (AsuI) |
| Eco47III | Eco48I (PstI) | Eco49I (PstI) |
| Eco50I (HgiCI) | Eco51I (Eco31I) | Eco51II (ScrFI) |
| Eco52I (XmaIII) | Eco55I (SacII) | Eco56I (Nael) |
| Eco57I | Eco60I (EcoRII) | Eco61I (EcoRII) |
| Eco64I (HgiCI) | Eco65I (HindIII) | Eco67I (EcoRII) |
| Eco68I (HgiJII) | Eco70I (EcoRII) | Eco71I (EcoRII) |
| Eco72I (PmaCI) | Eco76I (SauI) | Eco78I (NarI) |
| Eco80I (ScrFI) | Eco81I (SauI) | Eco82I (EcoRI) |
| Eco83I (PstI) | Eco85I (ScrFI) | Eco88I (AvaiI) |
| Eco90I (CfrI) | Eco91I (BstEII) | Eco92I (SacII) |
| Eco93I (ScrFI) | Eco95I (Eco31I) | Eco96I (SacII) |
| Eco97I (Eco31I) | Eco98I (HindIII) | Eco99I (SacII) |
| Eco100I (SacII) | Eco101I (Eco31I) | Eco104I (SacII) |
| Eco105I (SnaBI) | Eco113I (HgiJII) | Eco115I (SauI) |
| Eco118I (SauI) | Eco120I (Eco31I) | Eco127I (Eco31I) |
| Eco128I (EcoRII) | Eco129I (Eco31I) | Eco130I (StyI) |
| Eco133I (PstI) | Eco134I (SacII) | Eco135I (SacII) |

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| Eco136I (EcoRI) | Eco136II (SacI) | Eco141I (PstI) |
| Eco147I (StuI) | Eco149I (KpnI) | Eco153I (SceFI) |
| Eco155I (Eco31I) | Eco156I (Eco31I) | Eco157I (Eco31I) |
| Eco158I (SacII) | Eco158II (SnaBI) | Eco159I (EcoRI) |
| Eco161I (PstI) | Eco162I (Eco31I) | Eco164I (CfrI) |
| Eco165I (EcoRII) | Eco167I (PstI) | EcoHI (CfrI) |
| EcoICRI (SacI) | EcoNI | EcoO65I (BstEII) |
| EcoO109I (DraII) | EcoRI | EcoRII |
| EcoRV | EcoT14I (Styl) | EcoT22I (AvaIII) |
| EcoT38I (HgiJII) | EcoT104I (Styl) | EcoVIII (HindIII) |
| ErpI (Avall) | EspI | FbaI (BclI) |
| FbrI (Fnu4HI) | FaiI (Avall) | FdiI (MstI) |
| FinI | FinII (HpaII) | FinSI (HaeIII) |
| Fnu4HI | FnuAI (HinfI) | FnuAII (MboI) |
| FnuCI (MboI) | FnuDI (HaeIII) | FnuDII |
| FnuDIII (HhaI) | FnuEI (MboI) | FokI |
| FspI (Eco57I) | FspI (MstI) | FspI (AsuI) |
| FspMI (FnuDII) | FspMSI (Avall) | FsuI (Tth111I) |
| Gall (SacII) | GceI (SacII) | GceGLI (SacII) |
| GdiI (StuI) | GdiII | GdiI (BamHI) |
| GinI (BamHI) | GoxI (BamHI) | GspAI (AvaII) |
| GspAII (MstI) | GsuI | HacI (MboI) |
| HaeI | HaeII | HaeIII |
| HapII (HpaII) | HgaI | HgiAI |
| HgiBI (Avall) | HgiCI | HgiCII (Avall) |
| HgiCIII (SalI) | HgiDI (AcyI) | HgiDII (SalI) |
| HgiEI (Avall) | HgiEII | HgiGI (AcyI) |
| HgiHI (HgiCI) | HgiHII (AcyI) | HgiHIII (Avall) |
| HgiJI (Avall) | HgiJII | HgsS21I (CauI) |
| Hhal | HhalII (HinfI) | HhgI (HaeIII) |
| Hin173I (HindIII) | Hin1056I (FnuDII) | Hin1076III (HindIII) |
| Hin1160II (HindII) | Hin1161II (HindII) | HinGII (HhaI) |
| HinGIII (FokI) | HinHII (HaeII) | HinJCI (HindII) |
| HinJCI (HindIII) | HinPII (HhaI) | HinSII (HhaI) |
| HinS2I (HhaI) | HinbIII (HindIII) | HincII (HindII) |
| HindII | HindIII | HinfI |
| HinfII (HindIII) | HjaI (EcoRV) | HpaI |
| HpaII | HphI | HsuI (HindIII) |
| KoxI (BstEII) | KoxII (HgiJII) | KpnI |
| KpnK14I (KpnI) | Kpn2I (BspMI) | Ksp632I |
| LspI (AsuII) | MaeI | MaeII |
| MaeIII | Maui (PstI) | MboI |
| MboII | McaI (Xhol) | MecI (Xhol) |
| MeuI (MboI) | MfII (XhoII) | MisI (NaeI) |
| MjaI (MaeI) | MjaII (AsuI) | MkiI (HindIII) |
| MkrI (PstI) | MlaI (AsuII) | MleI (BamHI) |
| MliI (AluI) | MluI | MmeI |
| MmeII (MboI) | MniI (HaeIII) | MniII (HpaII) |
| MnII | MnnI (HindII) | MnnII (HaeIII) |
| MnnIV (HhaI) | MnoI (HpaII) | MnoIII (MboI) |
| MosI (MboI) | MphiI (EcoRII) | MpuI (Xhol) |
| MraI (SacII) | Mrol (BspMI) | MseI |
| MsiI (Xhol) | MspI (HpaII) | Msp67I (SceFI) |
| Msp67II (MboI) | MstI | MstII (SauI) |
| MthI (MboI) | MvaI (EcoRII) | MvnI (FnuDII) |
| MzI (PvuII) | Nael | NamI (NarI) |
| NanI (EcoRV) | NanII (DpnI) | NarI |

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| NasI (PstI) | NasBI (BamHI) | NasSI (SacI) |
| NasWI (NaeI) | NbaI (NaeI) | NblII (PvuI) |
| NbrI (NaeI) | NcaI (HinfI) | NciI (CeuII) |
| NcoI | NcuI (MboI) | NdaI (NarI) |
| NdeI | NdeII (MboI) | NfII (MboI) |
| NflAI (EcoRV) | NflAII (MboI) | NfbI (MboI) |
| NgbI (PstI) | NgoI (HaeII) | NgoII (HaeIII) |
| NgoIII (SacII) | NgoBI (HphI) | NgoDI (SacII) |
| NgoDIII (DpnI) | NgoMI (NaeI) | NgoPII (HaeIII) |
| NgoPIII (SacII) | NgoSI (HaeIII) | NheI |
| NlaI (HaeIII) | NlaII (MboI) | NlaIII |
| NlaIV | NlaDI (MboI) | NlaDII (AsuI) |
| NlaDIII (SacII) | NmeCI (MboI) | NlaSI (SacII) |
| NlaSII (AcyI) | NmuDI (DpnI) | NmlI (KpnI) |
| NmuI (NaeI) | NmuFI (NaeI) | NmuEI (DpnI) |
| NmuEII (AsuI) | NopI (SalI) | NmuSI (AsuI) |
| NoI (PstI) | NphI (MboI) | NotI |
| NovII (HinfI) | NsiAI (MboI) | NruI |
| NsiII (AvaIII) | NspI | NsiCI (EcoRV) |
| NsiIII (HinfI) | NspIV (AsuI) | NspII (SdI) |
| NspIII (AvaI) | NspBI (AsuI) | NspV (AsuII) |
| NspAI (MboI) | NspHII (AvaII) | NspBII |
| NspHI (NspI) | NspSAII (BstEII) | NspMACI (BglII) |
| NspSAI (AvaI) | NspWI (NaeI) | NspSAIII (NcoI) |
| NspSAIV (BamHI) | NtaI (Tth111I) | NsuDI (DpnI) |
| NsuI (MboI) | NunII (NarI) | NtsI (SstI) |
| NtsII (NaeI) | OxaI (AluI) | OtuI (AluI) |
| OtuNI (AluI) | PaaI (SacII) | OxaNI (SacI) |
| Pael (SphI) | PaiI (HaeIII) | PaeR7I (XbaI) |
| Pail (HaeIII) | PfIAI (FnuDII) | Pani (XbaI) |
| Pfal (MboI) | PfTWI (XbaI) | PfIMI |
| PfTNI (XbaI) | PleI | PfUI (SphI) |
| PgII (NaeI) | PmaCI | Pmal (PstI) |
| Pma44I (PstI) | PpaI (Eco31I) | PmyI (PstI) |
| PovI (BclI) | PspI (AsuI) | PpuI (HaeIII) |
| PpuMI | PstI | Psp61I (NaeI) |
| PsiI (DraII) | RheI (SalI) | PvuI |
| PvuII | RliI (NaeI) | RhpI (SalI) |
| RhsI (BamHI) | RsaI | RriI (SalI) |
| RvoI (SalI) | RspI (PvuI) | RshI (PvuI) |
| RshII (CeuII) | RsrII | RspXI (BspHI) |
| RsrI (EcoRI) | SacI | SaaI (SacII) |
| SabI (SacII) | Saki (SacII) | SacII |
| SacAI (NaeI) | SaICl (NaeI) | SaII |
| SaIAl (MboI) | SaIPI (PstI) | SaIDI (NruI) |
| SaIHI (MboI) | Sau3AI (MboI) | SaoI (NaeI) |
| SauI | Sau6782I (MboI) | Sau96I (AsuI) |
| Sau3239I (XbaI) | SbIAl (StyI) | SauAI (NaeI) |
| SauBMKI (NaeI) | Sbo13I (NruI) | SbIBI (StyI) |
| SbICl (StyI) | ScI (XbaI) | SboI (SacII) |
| Scal | ScRFI | ScrNI (HhaI) |
| Scal (SacI) | SdI (AsuI) | ScuI (XbaI) |
| SdI | SecII (SacI) | SecI |
| SecII (HpaII) | SecIII (SacI) | SexI (XbaI) |
| SfxGUI (HpaII) | SfaI (HaeIII) | SfaNI |
| SfiI | SfiI (PstI) | SfmI (AvaII) |
| SfoI (NarI) | SfrI (SacII) | SgaI (XbaI) |

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|--------------------------|--------------------------|-------------------------|
| SgoI (<i>Xba</i> I) | SgrII (<i>Eco</i> RII) | ShyI (<i>Sac</i> II) |
| SinI (<i>Ava</i> II) | SinAI (<i>Ava</i> II) | SinBI (<i>Ava</i> II) |
| SinCI (<i>Ava</i> II) | SinDI (<i>Ava</i> II) | SinEI (<i>Ava</i> II) |
| SinFI (<i>Ava</i> II) | SinGI (<i>Ava</i> II) | SinHI (<i>Ava</i> II) |
| SinJI (<i>Ava</i> II) | SinMI (<i>Mbo</i> I) | SkaI (<i>Nae</i> I) |
| SkaII (<i>Pst</i> I) | SlaI (<i>Xba</i> I) | SluI (<i>Xba</i> I) |
| SmaI | SnaI | SnaBI |
| SnoI (<i>Apal</i> II) | SpaI (<i>Xba</i> I) | SpaXI (<i>Sph</i> I) |
| SpeI | SphI | SpI |
| SpiII (<i>Tth</i> 111I) | SpiIII (<i>Hae</i> III) | SsoI (<i>Eco</i> RI) |
| SsoI (<i>Scr</i> FI) | SspI | SstI (<i>Sac</i> I) |
| SstII (<i>Sac</i> II) | SstIV (<i>Bcl</i> II) | SthI (<i>Kpn</i> I) |
| SthAI (<i>Kpn</i> I) | SthBI (<i>Kpn</i> I) | SthCI (<i>Kpn</i> I) |
| SthDI (<i>Kpn</i> I) | SthEI (<i>Kpn</i> I) | SthFI (<i>Kpn</i> I) |
| SthGI (<i>Kpn</i> I) | SthHII (<i>Kpn</i> I) | SthJI (<i>Kpn</i> I) |
| SthKI (<i>Kpn</i> I) | SthLI (<i>Kpn</i> I) | SthMI (<i>Kpn</i> I) |
| SthNI (<i>Kpn</i> I) | StuI | StyI |
| SuaI (<i>Hae</i> III) | SuII (<i>Hae</i> III) | TaqI |
| TaqII | TaqXI (<i>Eco</i> RII) | TceI (<i>Mbo</i> II) |
| TfiI (<i>Taq</i> I) | TgI (<i>Sac</i> II) | ThaI (<i>Fnu</i> DII) |
| TruI (<i>Ava</i> II) | TruII (<i>Mbo</i> I) | Tsp45I |
| TspEI | TspZNI (<i>Hae</i> III) | TteI (<i>Tth</i> 111I) |
| Tth111I | Tth111II | TthHB8I (<i>Taq</i> I) |
| TtnI (<i>Hae</i> III) | TtrI (<i>Tth</i> 111D) | VanI (<i>Bgl</i> I) |
| VhaI (<i>Hae</i> III) | VneI (<i>Apal</i> II) | VspI |
| XamI (<i>Sal</i> I) | XbaI | XcaI (<i>Sna</i> I) |
| XciI (<i>Sal</i> I) | XcyI (<i>Sma</i> I) | XhoI |
| XhoII | XmaI (<i>Sma</i> I) | XmaII (<i>Pst</i> I) |
| XmaIII | XmnI | XniI (<i>Pvu</i> I) |
| XorI (<i>Pst</i> I) | XorII (<i>Pvu</i> I) | XpaI (<i>Xba</i> I) |
| XphiI (<i>Pst</i> I) | YenI (<i>Pst</i> I) | YenAI (<i>Pst</i> I) |
| YenBI (<i>Pst</i> I) | YenCI (<i>Pst</i> I) | YenDI (<i>Pst</i> I) |
| YenEI (<i>Pst</i> I) | ZanI (<i>Eco</i> RII) | |

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