Name	Sequence	Site Length	Overhang	Frequency	Cut Positions
AgsI	TTSAA	5	three_prime	2	243, 341
ApoI	RAATTY	6	five_prime	1	63
BbvCI	CCTCAGC	7	five_prime	1	269
BbvI	GCAGC	5	five_prime	1	179
BcgI	CGANNNNNNTGC	6	three_prime	2	127, 161
BglI	GCCNNNNNGGC	6	three_prime	1	164
Bpu10I	CCTNAGC	6	five_prime	1	269
BseMII	CTCAG	5	three_prime	2	260, 347
BseSI	GKGCMC	6	three_prime	1	121
BsmAI	GTCTC	5	five_prime	1	365
BsrI	ACTGG	5	three_prime	1	234
BstXI	CCANNNNNNTGG	6	three_prime	1	178
CspCI	CAANNNNNGTGG	7	three_prime	2	174, 209
DraII	RGGNCCY	6	five_prime	1	371
EcoP15I	CAGCAG	6	five_prime	1	148
EcoRI	GAATTC	6	five_prime	1	63
EcoRII	CCWGG	5	five_prime	3	67, 84, 285
FokI	GGATG	5	five_prime	1	40
NspI	RCATGY	6	three_prime	1	153
PfoI	TCCNGGA	6	five_prime	1	285
PleI	GAGTC	5	five_prime	2	103, 366
SduI	GDGCHC	6	three_prime	2	121, 147
SfaNI	GCATC	5	five_prime	1	255
SmlI	CTYRAG	6	five_prime	1	90
SpeI	ACTAGT	6	five_prime	1	208
SphI	GCATGC	6	three_prime	1	153
StyI	CCWWGG	6	five_prime	1	112

Name	Sequence	Site Length	Overhang	Frequency	Cut Positions
TatI	WGTACW	6	five_prime	1	205
TseI	GCWGC	5	five_prime	1	167
TspRI	CASTG	5	three_prime	1	241