

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	2	2003	3460			
AatII	GACGT!C	5	126	179	262	448	1206
Acc65I	G!GTACC	0					
AccI	GT!MKAC	2	1206	2154			
AccII	CG!CG	16	61	1547	2288	2563	2939
			2959	2983	3014	3705	4006
			4444	4528	4591	4660	4933
			5514				
AccIII	T!CCGGA	0					
AciI	G!CGG	30	371	404	529	919	1186
			1547	1722	2034	2076	2282
			2899	2926	3014	3124	3705
			3769	3870	3873	4153	4208
			4306	4444	4591	4731	4974
			5219	5340	5359	5486	5514
AciI	C!CGC	42	59	87	99	113	280
			508	590	1243	1351	1366
			1384	1408	1489	1531	1887
			2025	2127	2286	2954	2957
			2971	3061	3309	3321	3330
			3342	3352	3363	3409	3548
			3611	4113	4158	4224	4250
			4375	4378	4526	4658	4965
			5109	5605			
AcyI	GR!CGYC	9	123	176	259	445	1203
			2151	3639	4341	4420	
AfaI	GT!AC	12	220	245	300	333	384
			541	716	1328	1619	1769
			2021	3945			
AflI	G!GWCC	6	927	1033	1069	1095	4155
			4600				
AflII	C!TTAAG	1	2521				
AflIII	A!CRYGT	1	5557				
AgeI	A!CCGGT	1	1555				
AhaI	CC!SGG	7	1340	1504	2148	3643	3803
			4416	5180			
AhaII	GR!CGYC	9	123	176	259	445	1203
			2151	3639	4341	4420	
AhaIII	TTT!AAA	3	2341	4783	4802		
AluI	AG!CT	22	567	619	625	937	959
			1283	1447	1588	1615	1761
			2068	2418	2585	2842	3137
			3425	3745	4203	5000	5257
			5303	5393			
Alw44I	G!TGCAC	2	1654	5243			
AlwI	GGATCNNNN!	8	1557	3507	3818	4064	4456
			4815	4913	4999		
AlwI	!NNNNNGATCC	7	581	1544	2109	3883	4427
			4814	4911			
AlwNI	CAGNNN!CTG	3	2241	4740	5148		
AocI	CC!TNAGG	3	1833	3119	4751		
AocII	GDGCH!C	16	621	1351	1360	1394	1631
			1658	1691	2268	2830	3585
			3678	3752	3942	4004	4714
			5247				

Aor51HI	AGC!GCT	1	597					
AosI	TGC!GCA	1	3741					
AosII	GR!CGYC	9	123	176	259	445	1203	
			2151	3639	4341	4420		
ApaI	GGGCC!C	0						
ApaLI	G!TGCAC	2	1654	5243				
ApoI	R!AATTY	4	630	2453	2557	2568		
ApyI	CC!WGG	21	91	284	660	870	1046	
			1170	1497	1751	1799	1895	
			1910	2039	3175	3230	3247	
			4026	4605	4708	5398	5411	
			5532					
AquI	C!YCGRG	2	614	1393				
AscI	GG!CGCGCC	0						
AseI	AT!TAAT	1	8					
AsnI	AT!TAAT	1	8					
Asp700I	GAANN!NNTTC	0						
Asp718I	G!GTACC	0						
AspHI	GWGCW!C	5	621	1658	3752	3942	5247	
AspI	GACN!NNGTC	1	3757					
AsuI	G!GNCC	18	84	277	927	1024	1033	
			1069	1095	1341	1724	1967	
			2059	2080	2746	4155	4600	
			4646	4704	4737			
AsuII	TT!CGAA	2	628	4321				
AvaI	C!YCGRG	2	614	1393				
AvaII	G!GWCC	6	927	1033	1069	1095	4155	
			4600					
AvaIII	ATGCA!T	3	3211	3283	5614			
AviII	TGC!GCA	1	3741					
AvrI	!CYCGRG	2	613	1392				
AvrII	C!CTAGG	2	3461	4476				
AxyI	CC!TNAGG	3	1833	3119	4751			
BalI	TGG!CCA	3	971	2234	3721			
BamHI	G!GATCC	1	1549					
BanI	G!GYRCC	7	466	1499	1628	1688	2792	
			3638	3673				
BanII	GRGCY!C	6	621	1360	1394	2830	4004	
			4714					
BanIII	AT!CGAT	1	3479					
BbeI	GGCGC!C	1	3642					
BbiII/AcyI	GR!CGYC	9	123	176	259	445	1203	
			2151	3639	4341	4420		
BbrPI	CAC!GTG	1	979					
BbsI	!GTCTTC	3	1138	1261	1270			
BbsI	!GAAGAC	3	1867	2100	2168			
BbvI	GCAGCNNNNNNN!	13	661	664	667	946	1503	
			1785	2227	2427	3712	3754	
			4275	4728	5228			
BbvI	!NNNNNNNNNNNGCTGC	13	799	860	1270	1434	1520	
			1904	2921	3586	3770	3863	
			4929	5135	5138			
BbvII	GAAGACNN!	3	1875	2108	2176			
BbvII	!NNNNNNGTCTTC	3	1132	1255	1264			
BcefI	ACGGCNNNNNNNNNN!	7	1031	1786	1930	1981	2047	
			4065	4552				
BcefI	!NNNNNNNNNNNGCCGT	4	1122	2781	3599	5058		
BcgI	!GCANNNNNNTCG	0						
BcgI	!CGANNNNNNTGC	0						

BclI	T!GATCA	0						
BcnI	CC!SGG	7	1340 4416	1504 5180	2148	3643	3803	
BfaI	C!TAG	11	593 2488 5064	944 2906	1280 3462	2253 4477	2294 4811	
BfrI	C!TTAAG	1	2521					
BglI	GCCNNNN!NGGC	5	91	213	284	2043	3414	
BglII	A!GATCT	1	610					
BinI	!NNNNNGATCC	7	581 4814	1544 4911	2109	3883	4427	
BinI	GGATCNNNN!	8	1557 4815	3507 4913	3818 4999	4064	4456	
BlnI	C!CTAGG	2	3461	4476				
BmyI	GDGCH!C	16	621 1658 3678 5247	1351 1691 3752	1360 2268 3942	1394 2830 4004	1631 3585 4714	
BpmI	!CTCCAG	3	862	1906	4436			
BpmI	!CTGGAG	2	1045	4457				
Bpu1102I	GC!TNAGC	0						
BsaAI	YAC!GTR	4	341	979	2755	3943		
BsaBI	GATNN!NNATC	2	2301	3498				
BsaHI	GR!CGYC	9	123 2151	176 3639	259 4341	445 4420	1203	
BsaI	GGTCTCN!	0						
BsaI	!NNNNNGAGACC	2	2195	4628				
BsaJI	C!CNNGG	26	361 1495 1798 3403 4476 5397	1027 1503 1893 3412 4603	1168 1566 3173 3461 4604	1179 1662 3245 3802 4706	1317 1797 3368 4071 4707	
BseAI	T!CCGGA	0						
BsePI	!GCGCGC	0						
BsgI	!GTGCAG	0						
BsgI	!CTGCAC	0						
BsiEI	CGRY!CG	5	1531	1560	2286	3548	5223	
BsiWI	C!GTACG	0						
BsiYI	CCNNNNN!NNGG	30	50 1312 1798 3369 4478 4726	918 1502 1964 3620 4609 5079	1045 1509 2030 4164 4644 5358	1051 1537 2651 4445 4694 5524	1185 1685 2977 4477 4695 5542	
BslI	CCNNNNN!NNGG	30	50 1312 1798 3369 4478 4726	918 1502 1964 3620 4609 5079	1045 1509 2030 4164 4644 5358	1051 1537 2651 4445 4694 5524	1185 1685 2977 4477 4695 5542	
BsmAI	!GTCTC	1	429					
BsmAI	!GAGAC	5	2200	3068	3486	4505	4633	
BsmI	GAATGCN!	1	2390					
BsmI	G!CATTC	2	742	2483				
Bsp1286I	GDGCH!C	16	621 1658 3678 5247	1351 1691 3752	1360 2268 3942	1394 2830 4004	1631 3585 4714	

BspDI	AT!CGAT	1	3479					
BspEI	T!CCGGA	0						
BspHI	T!CATGA	2	3065	4837				
BspMI	ACCTGCNNNN!	2	3907	4357				
BspMI	!NNNNNNNNGCAGGT	1	3526					
BspMII	T!CCGGA	0						
BspWI	GCNNNNN!NNGC	38	91	213	245	284	377	
			401	655	879	1355	1456	
			1497	1514	1533	1542	1612	
			2043	2866	2896	2923	2965	
			2967	3206	3278	3329	3408	
			3414	3630	3714	3737	3876	
			3882	3999	4082	4349	4536	
			4740	4939	5511			
BsrFI	R!CCGGY	6	738	1555	2856	3958	4139	
			4422					
BsrI	ACTGGN!	5	707	2178	3583	3784	5030	
BsrI	C!CAGT	5	296	2668	3346	5142	5155	
BssHII	G!CGCGC	0						
Bst1107I	GTA!TAC	0						
BstBI	TT!CGAA	2	628	4321				
BstEII	G!GTNACC	0						
BstI	G!GATCC	1	1549					
BstNI	CC!WGG	21	91	284	660	870	1046	
			1170	1497	1751	1799	1895	
			1910	2039	3175	3230	3247	
			4026	4605	4708	5398	5411	
			5532					
BstPI	G!GTNACC	0						
BstUI	CG!CG	16	61	1547	2288	2563	2939	
			2959	2983	3014	3705	4006	
			4444	4528	4591	4660	4933	
			5514					
BstXI	CCANNNNN!NTGG	2	979	1225				
BstYI	R!GATCY	11	586	610	1549	2114	3810	
			4056	4448	4807	4819	4905	
			4916					
Bsu36I	CC!TNAGG	3	1833	3119	4751			
CbiI	TT!CGAA	2	628	4321				
CcrI	C!TCGAG	1	614					
CelII	GC!TNAGC	0						
CfoI	GCG!C	24	598	1365	1488	1547	2905	
			2913	2939	2961	2970	2983	
			3014	3633	3641	3705	3742	
			4008	4268	4444	4530	4933	
			5042	5216	5316	5383		
Cfr10I	R!CCGGY	6	738	1555	2856	3958	4139	
			4422					
Cfr13I	G!GNCC	18	84	277	927	1024	1033	
			1069	1095	1341	1724	1967	
			2059	2080	2746	4155	4600	
			4646	4704	4737			
Cfr9I	C!CCGGG	0						
CfrI	Y!GGCCR	10	757	969	1314	1402	2232	
			2283	3545	3719	4110	4137	
ClaI	AT!CGAT	1	3479					
CpoI	CG!GWCCG	1	4155					
Csp45I	TT!CGAA	2	628	4321				
CspI	CG!GSCCG	0						

CviJI	RG!CY	124	43	85	94	278	567
			619	625	658	759	769
			812	868	873	922	937
			959	971	983	1018	1026
			1123	1136	1173	1189	1283
			1316	1343	1358	1392	1398
			1404	1447	1467	1494	1508
			1513	1517	1541	1588	1615
			1678	1725	1754	1761	1839
			1905	1917	1968	1995	2003
			2037	2060	2068	2082	2192
			2228	2234	2285	2308	2418
			2585	2605	2747	2828	2842
			2856	3137	3178	3250	3373
			3402	3408	3417	3425	3448
			3460	3466	3547	3564	3572
			3599	3624	3708	3717	3721
			3745	3783	3859	3876	3895
			3958	4002	4012	4029	4112
			4139	4143	4180	4203	4219
4399	4426	4487	4609	4648			
4705	4712	4719	4738	4749			
5000	5029	5072	5083	5148			
5227	5252	5257	5303	5393			
5491	5517	5535	5546				
CvnI	CC!TNAGG	3	1833	3119	4751		
DdeI	C!TNAG	13	607	725	1078	1451	1575
			1833	2130	3119	3421	4302
			4751	4874	5283		
DpnI	GA!TC	24	588	612	1439	1551	1940
			2090	2116	2298	3478	3482
			3501	3812	3890	3971	3980
			4058	4434	4450	4809	4821
			4899	4907	4918	4993	
DpnII	!GATC	24	586	610	1437	1549	1938
			2088	2114	2296	3476	3480
			3499	3810	3888	3969	3978
			4056	4432	4448	4807	4819
			4897	4905	4916	4991	
DraI	TTT!AAA	3	2341	4783	4802		
DraII	RG!GNCCY	2	1033	4737			
DraIII	CACNNN!GTG	2	2224	2755			
DrdI	GACNNNN!NNGTC	3	2710	3666	5455		
DsaI	C!CRYGG	6	361	1027	1179	1566	3368
			4071				
DsaV	!CCNGG	28	89	282	658	868	1044
			1168	1338	1495	1502	1749
			1797	1893	1908	2037	2146
			3173	3228	3245	3641	3801
			4024	4414	4603	4706	5178
5396	5409	5530					
EaeI	Y!GGCCR	10	757	969	1314	1402	2232
			2283	3545	3719	4110	4137
EagI	C!GGCCG	2	2283	3545			
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	5	1056	1582	3110	3987	4197
Ecl136II	GAG!CTC	1	619				
EclXI	C!GGCCG	2	2283	3545			

Eco105I	TAC!GTA	1	341					
Eco31I	GGTCTCN!	0						
Eco31I	!NNNNNGAGACC	2	2195	4628				
Eco47I	G!GWCC	6	927	1033	1069	1095	4155	
			4600					
Eco47III	AGC!GCT	1	597					
Eco52I	C!GGCCG	2	2283	3545				
Eco57I	!CTGAAG	3	2062	3763	4195			
Eco57I	!CTTCAG	2	786	5029				
Eco81I	CC!TNAGG	3	1833	3119	4751			
EcoNI	CCTNN!NNNAGG	1	1049					
EcoO109I	RG!GNCCY	2	1033	4737				
EcoO65I	G!GTNACC	0						
EcoRI	G!AATTC	1	630					
EcoRII	!CCWGG	21	89	282	658	868	1044	
			1168	1495	1749	1797	1893	
			1908	2037	3173	3228	3245	
			4024	4603	4706	5396	5409	
			5530					
EcoRV	GAT!ATC	0						
EcoT14I	C!CWGGG	8	361	1027	1179	1566	3368	
			3461	4071	4476			
EcoT22I	ATGCA!T	3	3211	3283	5614			
EcoT38I	GRGCY!C	6	621	1360	1394	2830	4004	
			4714					
EheI	GGC!GCC	1	3640					
Esp3I	CGTCTCN!	0						
Esp3I	!NNNNNGAGACG	0						
EspI	GC!TNAGC	0						
FbaI	T!GATCA	0						
FinI	!GTCCC	6	927	1239	3169	3241	3305	
			4600					
FinI	!GGGAC	8	161	312	480	1031	1081	
			3775	4307	4416			
Fnu4HI	GC!NGC	49	650	653	656	813	874	
			920	935	1187	1284	1366	
			1448	1489	1492	1534	1723	
			1774	1918	2035	2216	2283	
			2286	2416	2935	2957	2971	
			3409	3548	3600	3611	3701	
			3706	3743	3784	3871	3874	
			3877	4113	4209	4250	4264	
			4378	4717	4732	4943	5149	
			5152	5217	5360	5515		
FokI	GGATGNNNNNNNNN!	6	709	1475	3506	3963	3988	
			4442					
FokI	!NNNNNNNNNNNNNCATCC	8	926	1289	1322	1645	1733	
			1790	1945	3312			
FspI	TGC!GCA	1	3741					
GdiII	!NNNNNYGGCCG	6	751	1308	2277	3539	4104	
			4131					
GdiII	CGGCCRN!	3	1408	2289	3551			
GsuI	!CTCCAG	3	862	1906	4436			
GsuI	!CTGGAG	2	1045	4457				
HaeI	WGG!CCW	8	971	2003	2234	3460	3721	
			5083	5535	5546			
HaeII	RGCGC!Y	6	599	1366	2906	2914	3642	
			5317					
HaeIII	GG!CC	36	85	278	759	971	1026	

			1136	1316	1343	1404	1541
			1725	1968	2003	2037	2060
			2082	2192	2234	2285	2605
			2747	3402	3408	3417	3460
			3547	3721	4112	4139	4648
			4705	4738	5083	5517	5535
			5546				
HapII	C!CGG	23	602	739	1120	1339	1503
			1556	2148	2857	3544	3621
			3643	3671	3802	3892	3959
			4140	4415	4423	4515	4989
			5179	5205	5352		
HgaI	GACGCNNNNN!	3	527	4349	4428		
HgaI	!NNNNNNNNNGCGTC	4	2140	2972	4867	5445	
HgiAI	GWGCW!C	5	621	1658	3752	3942	5247
HgiEII	!ACCNNNNNNGGT	1	4969				
HhaI	GCG!C	24	598	1365	1488	1547	2905
			2913	2939	2961	2970	2983
			3014	3633	3641	3705	3742
			4008	4268	4444	4530	4933
			5042	5216	5316	5383	
Hin1I	GR!CGYC	9	123	176	259	445	1203
			2151	3639	4341	4420	
HinPII	G!CGC	24	596	1363	1486	1545	2903
			2911	2937	2959	2968	2981
			3012	3631	3639	3703	3740
			4006	4266	4442	4528	4931
			5040	5214	5314	5381	
HincII	GTY!RAC	1	2402				
HindIII	A!AGCTT	1	623				
HinfI	G!ANTC	17	411	605	792	1704	1854
			2181	2290	2682	2704	3114
			4124	4258	4310	4368	4405
			5187	5583			
HpaI	GTT!AAC	1	2402				
HpaII	C!CGG	23	602	739	1120	1339	1503
			1556	2148	2857	3544	3621
			3643	3671	3802	3892	3959
			4140	4415	4423	4515	4989
			5179	5205	5352		
HphI	GGTGANNNNNNN!	6	377	842	1066	1960	4710
			4826				
HphI	!NNNNNNNTCACC	6	977	1258	1407	1849	2756
			3817				
KasI	G!GCGCC	1	3638				
KpnI	GGTAC!C	0					
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	5	1052	1578	3106	3983	4193
KspI	CCGC!GG	0					
MaeI	C!TAG	11	593	944	1280	2253	2294
			2488	2906	3462	4477	4811
			5064				
MaeII	A!CGT	18	123	135	176	259	340
			445	978	1203	1932	2208
			2699	2711	2754	2864	3755
			3942	4723	4856		
MaeIII	!GTNAC	14	62	149	498	670	1855
			2427	2928	2940	3759	4065
			4755	5021	5137	5200	

MamI	GATNN!NNATC	2	2301	3498				
MboI	!GATC	24	586	610	1437	1549	1938	
			2088	2114	2296	3476	3480	
			3499	3810	3888	3969	3978	
			4056	4432	4448	4807	4819	
			4897	4905	4916	4991		
MboII	GAAGANNNNNNNN!	12	1015	1069	1595	1880	1993	
			2113	2181	2895	3123	4000	
			4210	4829				
MboII	!NNNNNNNTCTTC	11	772	778	834	892	1132	
			1255	1264	4290	4457	4657	
			4900					
McrI	!CGRYCG	5	1527	1556	2282	3544	5219	
MfeI	!CAATTG	1	2388					
MflI	R!GATCY	11	586	610	1549	2114	3810	
			4056	4448	4807	4819	4905	
			4916					
MluI	A!CGCGT	0						
MmeI	!TCCRAC	2	1960	2706				
MmeI	!GTYGGA	2	5182	5366				
MnlI	CCTCNNNNNNN!	21	646	974	1297	1316	1340	
			1421	1737	1916	1931	2014	
			2147	2258	2358	2367	3422	
			3428	4240	4446	4760	5130	
			5454					
MnlI	!NNNNNNNGAGG	32	541	671	682	1014	1041	
			1044	1052	1125	1311	1617	
			1656	1677	1710	1827	1936	
			1992	2181	2214	2317	2782	
			3113	3391	3397	3434	3437	
			3449	3489	3553	3689	4046	
			4476	5379				
MroI	T!CCGGA	0						
MscI	TGG!CCA	3	971	2234	3721			
MseI	T!TAA	18	8	1593	1818	2274	2340	
			2401	2522	2543	2554	2566	
			2577	2594	2692	2963	4782	
			4796	4801	4853			
MspI	C!CGG	23	602	739	1120	1339	1503	
			1556	2148	2857	3544	3621	
			3643	3671	3802	3892	3959	
			4140	4415	4423	4515	4989	
			5179	5205	5352			
MstI	TGC!GCA	1	3741					
MstII	CC!TNAGG	3	1833	3119	4751			
MunI	C!AATTG	1	2389					
MvaI	CC!WGG	21	91	284	660	870	1046	
			1170	1497	1751	1799	1895	
			1910	2039	3175	3230	3247	
			4026	4605	4708	5398	5411	
			5532					
MvnI	CG!CG	16	61	1547	2288	2563	2939	
			2959	2983	3014	3705	4006	
			4444	4528	4591	4660	4933	
			5514					
NaeI	GCC!GGC	3	2858	4141	4424			
NarI	GG!CGCC	1	3639					
NciI	CC!SGG	7	1340	1504	2148	3643	3803	
			4416	5180				

NcoI	C!CATGG	6	361	1027	1179	1566	3368
			4071				
NdeI	CA!TATG	1	235				
NdeII	!GATC	24	586	610	1437	1549	1938
			2088	2114	2296	3476	3480
			3499	3810	3888	3969	3978
			4056	4432	4448	4807	4819
			4897	4905	4916	4991	
NgoMI	G!CCGGC	3	2856	4139	4422		
NheI	G!CTAGC	2	592	1279			
NlaIII	CATG!	30	305	365	799	917	976
			1031	1147	1183	1279	1570
			1606	1612	1624	1702	1768
			1848	2059	3069	3209	3281
			3372	3513	3858	4044	4075
			4101	4457	4841	5561	5612
NlaIV	GGN!NCC	23	468	929	1034	1035	1468
			1501	1551	1630	1690	2081
			2794	2815	2827	3018	3179
			3251	3640	3675	4523	4602
			4647	5490	5529		
NotI	GC!GGCCGC	1	2283				
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	3211	3283	5614		
Nsp(7524)I	RCATG!Y	5	1606	3209	3281	4044	5561
Nsp(7524)V	TT!CGAA	2	628	4321			
NspBII	CMG!CKG	6	1533	2027	3137	3745	4974
			5219				
NspII	GDGCH!C	16	621	1351	1360	1394	1631
			1658	1691	2268	2830	3585
			3678	3752	3942	4004	4714
			5247				
NspIII	C!YCGRG	2	614	1393			
NspIV	G!GNCC	18	84	277	927	1024	1033
			1069	1095	1341	1724	1967
			2059	2080	2746	4155	4600
			4646	4704	4737		
NunII	GG!CGCC	1	3639				
PacI	TTAAT!TAA	0					
Paer71	C!TCGAG	1	614				
PalI	GG!CC	36	85	278	759	971	1026
			1136	1316	1343	1404	1541
			1725	1968	2003	2037	2060
			2082	2192	2234	2285	2605
			2747	3402	3408	3417	3460
			3547	3721	4112	4139	4648
			4705	4738	5083	5517	5535
			5546				
PflMI	CCANNNN!NTGG	0					
PleI	GAGTCNNNN!	3	1862	2690	3122		
PleI	!NNNNGACTC	6	405	599	2284	2698	4304
			5181				
PmaCI	CAC!GTG	1	979				
PmeI	GTTT!AAAC	0					
PmlI	CAC!GTG	1	979				
PpuMI	RG!GWCCY	1	1033				
PshAI	GACNN!NNGTC	2	1100	2207			
PstI	CTGCA!G	2	817	936			
PvuI	CGAT!CG	0					

PvuII	CAG!CTG	2	3137	3745				
RmaI	C!TAG	11	593	944	1280	2253	2294	
			2488	2906	3462	4477	4811	
			5064					
RsaI	GT!AC	12	220	245	300	333	384	
			541	716	1328	1619	1769	
			2021	3945				
RsrI	G!AATTC	1	630					
RsrII	CG!GWCCG	1	4155					
SacI	GAGCT!C	1	621					
SacII	CCGC!GG	0						
SalI	G!TCGAC	0						
Sau3AI	!GATC	24	586	610	1437	1549	1938	
			2088	2114	2296	3476	3480	
			3499	3810	3888	3969	3978	
			4056	4432	4448	4807	4819	
			4897	4905	4916	4991		
Sau96I	G!GNCC	18	84	277	927	1024	1033	
			1069	1095	1341	1724	1967	
			2059	2080	2746	4155	4600	
			4646	4704	4737			
SauI	CC!TNAGG	3	1833	3119	4751			
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	28	91	284	660	870	1046	
			1170	1340	1497	1504	1751	
			1799	1895	1910	2039	2148	
			3175	3230	3247	3643	3803	
			4026	4416	4605	4708	5180	
			5398	5411	5532			
SduI	GDGCH!C	16	621	1351	1360	1394	1631	
			1658	1691	2268	2830	3585	
			3678	3752	3942	4004	4714	
			5247					
SecI	C!CNNGG	26	361	1027	1168	1179	1317	
			1495	1503	1566	1662	1797	
			1798	1893	3173	3245	3368	
			3403	3412	3461	3802	4071	
			4476	4603	4604	4706	4707	
			5397					
SexI	!CTCGAG	1	613					
SfaNI	GCATCNNNNN!	9	1302	1308	1812	2455	3218	
			3290	3938	4002	4277		
SfaNI	!NNNNNNNNNGATGC	8	358	1965	2004	2133	3597	
			3852	4068	5460			
SfcI	C!TRYAG	5	813	932	2974	5101	5292	
SfiI	GGCCNNNN!NGGCC	1	3414					
SfuI	TT!CGAA	2	628	4321				
SgrAI	CR!CCGGYG	0						
SinI	G!GWCC	6	927	1033	1069	1095	4155	
			4600					
SmaI	CCC!GGG	0						
SnaBI	TAC!GTA	1	341					
SnaI	!GTATAC	0						
SnoI	G!TGCAC	2	1654	5243				
SpeI	A!CTAGT	0						
SphI	GCATG!C	3	3209	3281	4044			
SplI	C!GTACG	0						
SrfI	GCCC!GGGC	0						
Sse8387I	CCTGCA!GG	0						

SspI	AAT!ATT	2	2547	3100				
SstI	GAGCT!C	1	621					
SstII	CCGC!GG	0						
SstIII	!ACGT	18	122	134	175	258	339	
			444	977	1202	1931	2207	
			2698	2710	2753	2863	3754	
			3941	4722	4855			
StuI	AGG!CCT	2	2003	3460				
StyI	C!CWWGG	8	361	1027	1179	1566	3368	
			3461	4071	4476			
StySJI	!GAGNNNNNGTRC	0						
StySJI	!GYACNNNNNNCTC	1	714					
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	16	615	628	1716	1743	2211	
			2788	3479	3752	3908	3932	
			3968	4130	4321	4366	4621	
			5459					
TaqII	GACCGANNNNNNNNNN!	3	2659	3676	4344			
TaqII	!NNNNNNNNNTCGGTC	2	763	4588				
TaqII	CACCCANNNNNNNNNN!	2	1706	1811				
TaqII	!NNNNNNNNNTGGGTG	3	173	3543	4135			
TfiI	G!AWTC	8	792	1704	2181	4124	4258	
			4368	4405	5583			
ThaI	CG!CG	16	61	1547	2288	2563	2939	
			2959	2983	3014	3705	4006	
			4444	4528	4591	4660	4933	
			5514					
Tsp45I	!GTSAC	5	670	1855	2928	3759	4065	
TspEI	!AATT	13	19	630	896	2275	2389	
			2453	2531	2557	2568	3215	
			3287	3379	4797			
Tth111I	GACN!NNGTC	1	3757					
Tth111II	CAARCANNNNNNNNNN!	2	1442	4966				
Tth111III	!NNNNNNNNNTGYTTG	4	1118	4074	4934	4973		
TthHB8I	T!CGA	16	615	628	1716	1743	2211	
			2788	3479	3752	3908	3932	
			3968	4130	4321	4366	4621	
			5459					
Van91I	CCANNNN!NTGG	0						
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	2293					
XcmI	!CCANNNNNNNNNTCG	1	3914					
XcyI	C!CCGGG	0						
XhoI	C!TCGAG	1	614					
XhoII	R!GATCY	11	586	610	1549	2114	3810	
			4056	4448	4807	4819	4905	
			4916					
XmaI	C!CCGGG	0						
XmaIII	C!GGCCG	2	2283	3545				
XmnI	GAANN!NNTTC	0						
XorII	CGAT!CG	0						