

Enzyme Name	Sequence	Count	Cutting Position					
AatI	AGG!CCT	1	2577					
AatII	GACGT!C	4	126	179	262	448		
Acc65I	G!GTACC	1	1376					
AccI	GT!MKAC	1	1371					
AccII	CG!CG	20	61	1026	1179	1251	1269	
			1382	1644	1680	2056	2076	
			2100	2131	2822	3123	3561	
			3645	3708	3777	4050	4631	
AccIII	T!CCGGA	1	1331					
AciI	G!CGG	26	371	404	529	627	1128	
			1382	2016	2043	2131	2241	
			2822	2886	2987	2990	3270	
			3325	3423	3561	3708	3848	
			4091	4336	4457	4476	4603	
			4631					
AciI	C!CGC	39	59	87	99	113	280	
			508	590	633	835	898	
			976	1119	1177	1297	1380	
			2071	2074	2088	2178	2426	
			2438	2447	2459	2469	2480	
			2526	2665	2728	3230	3275	
			3341	3367	3492	3495	3643	
			3775	4082	4226	4722		
AcyI	GR!CGYC	7	123	176	259	445	2756	
			3458	3537				
AfaI	GT!AC	9	220	245	300	333	384	
			541	815	1378	3062		
AflI	G!GWCC	2	3272	3717				
AflII	C!TTAAG	0						
AflIII	A!CRYGT	4	1062	1257	1642	4674		
AgeI	A!CCGGT	1	601					
AhaI	CC!SGG	7	1194	1388	1389	2760	2920	
			3533	4297				
AhaII	GR!CGYC	7	123	176	259	445	2756	
			3458	3537				
AhaIII	TTT!AAA	3	1458	3900	3919			
AluI	AG!CT	19	567	775	786	952	1033	
			1096	1349	1355	1535	1702	
			1959	2254	2542	2862	3320	
			4117	4374	4420	4510		
Alw44I	G!TGCAC	2	749	4360				
AlwI	GGATCNNNN!	9	1399	1408	2624	2935	3181	
			3573	3932	4030	4116		
AlwI	!NNNNNGATCC	8	581	802	1183	1386	3000	
			3544	3931	4028			
AlwNI	CAGNNN!CTG	2	3857	4265				
AocI	CC!TNAGG	2	2236	3868				
AocII	GDGCH!C	13	753	825	1296	1351	1388	
			1947	2702	2795	2869	3059	
			3121	3831	4364			
Aor51HI	AGC!GCT	1	597					
AosI	TGC!GCA	1	2858					
AosII	GR!CGYC	7	123	176	259	445	2756	
			3458	3537				
ApaI	GGGCC!C	1	1388					
ApaLI	G!TGCAC	2	749	4360				

ApoI	R!AATTY	4	1360	1570	1674	1685		
ApyI	CC!WGG	12	91	284	866	2292	2347	
			2364	3143	3722	3825	4515	
			4528	4649				
AquI	C!YCGRG	3	878	1344	1387			
AscI	GG!CGCGCC	0						
AseI	AT!TAAT	1	8					
AsnI	AT!TAAT	1	8					
Asp700I	GAANN!NNTTC	0						
Asp718I	G!GTACC	1	1376					
AspHI	GWGCW!C	5	753	1351	2869	3059	4364	
AspI	GACN!NNGTC	1	2874					
AsuI	G!GNCC	16	84	277	629	883	1013	
			1076	1141	1196	1384	1385	
			1863	3272	3717	3763	3821	
			3854					
AsuII	TT!CGAA	2	1358	3438				
AvaI	C!YCGRG	3	878	1344	1387			
AvaII	G!GWCC	2	3272	3717				
AvaIII	ATGCA!T	3	2328	2400	4731			
AviII	TGC!GCA	1	2858					
AvrI	!CYCGRG	3	877	1343	1386			
AvrII	C!CTAGG	2	2578	3593				
AxyI	CC!TNAGG	2	2236	3868				
BalI	TGG!CCA	1	2838					
BamHI	G!GATCC	1	1391					
BanI	G!GYRCC	6	466	923	1376	1909	2755	
			2790					
BanII	GRGCY!C	6	825	1351	1388	1947	3121	
			3831					
BanIII	AT!CGAT	1	2596					
BbeI	GGCGC!C	1	2759					
BbiII/AcyI	GR!CGYC	7	123	176	259	445	2756	
			3458	3537				
BbrPI	CAC!GTG	1	1258					
BbsI	!GTCTTC	0						
BbsI	!GAAGAC	0						
BbvI	GCAGCNNNNNNNN!	9	729	732	1185	1544	2829	
			2871	3392	3845	4345		
BbvI	!NNNNNNNNNNNGCTGC	8	762	2038	2703	2887	2980	
			4046	4252	4255			
BbvII	GAAGACNN!	0						
BbvII	!NNNNNNGTCTTC	0						
BcefI	ACGGCNNNNNNNNNN!	14	691	703	730	751	832	
			862	937	1012	1027	1090	
			1141	1186	3182	3669		
BcefI	!NNNNNNNNNNNGCCGT	5	741	1071	1898	2716	4175	
BcgI	!GCANNNNNTCG	0						
BcgI	!CGANNNNNTGC	0						
BclI	T!GATCA	1	1413					
BcnI	CC!SGG	7	1194	1388	1389	2760	2920	
			3533	4297				
BfaI	C!TAG	8	593	1404	1605	2023	2579	
			3594	3928	4181			
BfrI	C!TTAAG	0						
BglI	GCCNNNN!NGGC	4	91	213	284	2531		
BglII	A!GATCT	1	1340					
BinI	!NNNNNGATCC	8	581	802	1183	1386	3000	
			3544	3931	4028			

BinI	GGATCNNNN!	9	1399 3573	1408 3932	2624 4030	2935 4116	3181
BlnI	C!CTAGG	2	2578	3593			
BmyI	GDGCH!C	13	753 1947 3121	825 2702 3831	1296 2795 4364	1351 2869	1388 3059
BpmI	!CTCCAG	1	3553				
BpmI	!CTGGAG	1	3574				
Bpu1102I	GC!TNAGC	0					
BsaAI	YAC!GTR	4	341	1258	1872	3060	
BsaBI	GATNN!NNATC	2	1418	2615			
BsaHI	GR!CGYC	7	123 3458	176 3537	259	445	2756
BsaI	GGTCTCN!	0					
BsaI	!NNNNNGAGACC	3	758	1049	3745		
BsaJI	C!CNNGG	23	361 1072 2362 2919 3823	612 1192 2485 3188 3824	733 1380 2520 3593 4514	864 1387 2529 3720	879 2290 2578 3721
BseAI	T!CCGGA	1	1331				
BsePI	!GCGCGC	0					
BsgI	!GTGCAG	0					
BsgI	!CTGCAC	0					
BsiEI	CGRY!CG	4	606	898	2665	4340	
BsiWI	C!GTACG	0					
BsiYI	CCNNNNN!NNGG	22	50 1768 3562 3811 4641	786 2094 3594 3812 4659	1036 2486 3595 3843	1138 2737 3726 4196	1270 3281 3761 4475
BslI	CCNNNNN!NNGG	22	50 1768 3562 3811 4641	786 2094 3594 3812 4659	1036 2486 3595 3843	1138 2737 3726 4196	1270 3281 3761 4475
BsmAI	!GTCTC	1	429				
BsmAI	!GAGAC	6	763 3750	1054	2185	2603	3622
BsmI	GAATGCN!	1	1507				
BsmI	G!CATTC	1	1600				
Bsp1286I	GDGCH!C	13	753 1947 3121	825 2702 3831	1296 2795 4364	1351 2869	1388 3059
BspDI	AT!CGAT	1	2596				
BspEI	T!CCGGA	1	1331				
BspHI	T!CATGA	2	2182	3954			
BspMI	ACCTGCNNNN!	3	971	3024	3474		
BspMI	!NNNNNNNNGCAGGT	1	2643				
BspMII	T!CCGGA	1	1331				
BspWI	GCNNNNN!NNGC	36	91 401 1030 2013 2395 2831 3199 4628	213 768 1248 2040 2446 2854 3466	245 897 1296 2082 2525 2993 3653	284 906 1305 2084 2531 2999 3857	377 972 1983 2323 2747 3116 4056
BsrFI	R!CCGGY	6	601	767	1973	3075	3256

			3539					
BsrI	ACTGGN!	3	2700	2901	4147			
BsrI	C!CAGT	6	296	811	1785	2463	4259	
			4272					
BssHII	G!CGCGC	0						
Bst1107I	GTA!TAC	0						
BstBI	TT!CGAA	2	1358	3438				
BstEII	G!GTNACC	0						
BstI	G!GATCC	1	1391					
BstNI	CC!WGG	12	91	284	866	2292	2347	
			2364	3143	3722	3825	4515	
			4528	4649				
BstPI	G!GTNACC	0						
BstUI	CG!CG	20	61	1026	1179	1251	1269	
			1382	1644	1680	2056	2076	
			2100	2131	2822	3123	3561	
			3645	3708	3777	4050	4631	
BstXI	CCANNNNN!NTGG	0						
BstYI	R!GATCY	12	586	1188	1340	1391	1400	
			2927	3173	3565	3924	3936	
			4022	4033				
Bsu36I	CC!TNAGG	2	2236	3868				
CbiI	TT!CGAA	2	1358	3438				
CcrI	C!TCGAG	1	1344					
CelII	GC!TNAGC	0						
CfoI	GCG!C	28	598	909	1026	1092	1181	
			1251	1269	1310	2022	2030	
			2056	2078	2087	2100	2131	
			2750	2758	2822	2859	3125	
			3385	3561	3647	4050	4159	
			4333	4433	4500			
Cfr10I	R!CCGGY	6	601	767	1973	3075	3256	
			3539					
Cfr13I	G!GNCC	16	84	277	629	883	1013	
			1076	1141	1196	1384	1385	
			1863	3272	3717	3763	3821	
			3854					
Cfr9I	C!CCGGG	1	1387					
CfrI	Y!GGCCR	6	688	708	2662	2836	3227	
			3254					
ClaI	AT!CGAT	1	2596					
CpoI	CG!GWCCG	1	3272					
Csp45I	TT!CGAA	2	1358	3438				
CspI	CG!GSCCG	0						
CviJI	RG!CY	99	43	85	94	278	567	
			630	690	710	775	786	
			793	823	855	885	936	
			952	975	999	1006	1014	
			1033	1077	1096	1100	1110	
			1131	1143	1176	1197	1277	
			1349	1355	1386	1425	1535	
			1702	1722	1864	1945	1959	
			1973	2254	2295	2367	2490	
			2519	2525	2534	2542	2565	
			2577	2583	2664	2681	2689	
			2716	2741	2825	2834	2838	
			2862	2900	2976	2993	3012	
			3075	3119	3129	3146	3229	
			3256	3260	3297	3320	3336	

			3516	3543	3604	3726	3765
			3822	3829	3836	3855	3866
			4117	4146	4189	4200	4265
			4344	4369	4374	4420	4510
			4608	4634	4652	4663	
CvnI	CC!TNAGG	2	2236	3868			
DdeI	C!TNAG	8	887	1337	2236	2538	3419
			3868	3991	4400		
DpnI	GA!TC	23	588	809	1190	1342	1393
			1402	1415	2595	2599	2618
			2929	3007	3088	3097	3175
			3551	3567	3926	3938	4016
			4024	4035	4110		
DpnII	!GATC	23	586	807	1188	1340	1391
			1400	1413	2593	2597	2616
			2927	3005	3086	3095	3173
			3549	3565	3924	3936	4014
			4022	4033	4108		
DraI	TTT!AAA	3	1458	3900	3919		
DraII	RG!GNCCY	2	883	3854			
DraIII	CACNNN!GTG	4	757	1207	1291	1872	
DrdI	GACNNNN!NNGTC	3	1827	2783	4572		
DsaI	C!CRYGG	7	361	612	733	1072	1380
			2485	3188			
DsaV	!CCNGG	19	89	282	864	1192	1386
			1387	2290	2345	2362	2758
			2918	3141	3531	3720	3823
			4295	4513	4526	4647	
EaeI	Y!GGCCR	6	688	708	2662	2836	3227
			3254				
EagI	C!GGCCG	1	2662				
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	3	2227	3104	3314		
Ecl136II	GAG!CTC	1	1349				
EclXI	C!GGCCG	1	2662				
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	3	758	1049	3745		
Eco47I	G!GWCC	2	3272	3717			
Eco47III	AGC!GCT	1	597				
Eco52I	C!GGCCG	1	2662				
Eco57I	!CTGAAG	2	2880	3312			
Eco57I	!CTTCAG	1	4146				
Eco81I	CC!TNAGG	2	2236	3868			
EcoNI	CCTNN!NNNAGG	0					
EcoO109I	RG!GNCCY	2	883	3854			
EcoO65I	G!GTNACC	0					
EcoRI	G!AATTC	1	1360				
EcoRII	!CCWGG	12	89	282	864	2290	2345
			2362	3141	3720	3823	4513
			4526	4647			
EcoRV	GAT!ATC	0					
EcoT14I	C!CWGGG	6	361	612	2485	2578	3188
			3593				
EcoT22I	ATGCA!T	3	2328	2400	4731		
EcoT38I	GRGCY!C	6	825	1351	1388	1947	3121
			3831				
EheI	GGC!GCC	1	2757				

Esp3I	CGTCTCN!	0						
Esp3I	!NNNNNGAGACG	0						
EspI	GC!TNAGC	0						
FbaI	T!GATCA	1	1413					
FinI	!GTCCC	4	2286	2358	2422	3717		
FinI	!GGGAC	7	161	312	480	1232	2892	
			3424	3533				
Fnu4HI	GC!NGC	36	628	718	721	776	976	
			1129	1174	1177	1533	2052	
			2074	2088	2526	2665	2717	
			2728	2818	2823	2860	2901	
			2988	2991	2994	3230	3326	
			3367	3381	3495	3834	3849	
			4060	4266	4269	4334	4477	
			4632					
FokI	GGATGNNNNNNNNN!	4	2623	3080	3105	3559		
FokI	!NNNNNNNNNNNNNCATCC	2	1028	2429				
FspI	TGC!GCA	1	2858					
GdiII	!NNNNNYGGCCG	4	702	2656	3221	3248		
GdiII	CGGCCRN!	2	694	2668				
GsuI	!CTCCAG	1	3553					
GsuI	!CTGGAG	1	3574					
HaeI	WGG!CCW	7	1100	1277	2577	2838	4200	
			4652	4663				
HaeII	RGCGC!Y	6	599	1311	2023	2031	2759	
			4434					
HaeIII	GG!CC	31	85	278	630	690	710	
			885	1014	1077	1100	1131	
			1143	1197	1277	1386	1722	
			1864	2519	2525	2534	2577	
			2664	2838	3229	3256	3765	
			3822	3855	4200	4634	4652	
			4663					
HapII	C!CGG	22	602	768	1194	1332	1388	
			1398	1974	2661	2738	2760	
			2788	2919	3009	3076	3257	
			3532	3540	3632	4106	4296	
			4322	4469				
HgaI	GACGCNNNNN!	3	527	3466	3545			
HgaI	!NNNNNNNNNNNGCGTC	3	2089	3984	4562			
HgiAI	GWGCW!C	5	753	1351	2869	3059	4364	
HgiEII	!ACCNNNNNNGGT	1	4086					
HhaI	GCG!C	28	598	909	1026	1092	1181	
			1251	1269	1310	2022	2030	
			2056	2078	2087	2100	2131	
			2750	2758	2822	2859	3125	
			3385	3561	3647	4050	4159	
			4333	4433	4500			
Hin1I	GR!CGYC	7	123	176	259	445	2756	
			3458	3537				
HinP1I	G!CGC	28	596	907	1024	1090	1179	
			1249	1267	1308	2020	2028	
			2054	2076	2085	2098	2129	
			2748	2756	2820	2857	3123	
			3383	3559	3645	4048	4157	
			4331	4431	4498			
HincII	GTY!RAC	2	1372	1519				
HindIII	A!AGCTT	1	1353					
HinfI	G!ANTC	13	411	1328	1335	1799	1821	

			2231	3241	3375	3427	3485
			3522	4304	4700		
HpaI	GTT!AAC	1	1519				
HpaII	C!CGG	22	602	768	1194	1332	1388
			1398	1974	2661	2738	2760
			2788	2919	3009	3076	3257
			3532	3540	3632	4106	4296
			4322	4469			
HphI	GGTGANNNNNNN!	5	377	694	982	3827	3943
HphI	!NNNNNNNTCACC	7	691	973	1105	1210	1294
			1873	2934			
KasI	G!GCGCC	1	2755				
KpnI	GGTAC!C	1	1380				
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	3	2223	3100	3310		
KspI	CCGC!GG	1	1383				
MaeI	C!TAG	8	593	1404	1605	2023	2579
			3594	3928	4181		
MaeII	A!CGT	16	123	135	176	259	340
			445	747	1257	1816	1828
			1871	1981	2872	3059	3840
			3973				
MaeIII	!GTNAC	13	62	149	498	1207	1544
			2045	2057	2876	3182	3872
			4138	4254	4317		
MamI	GATNN!NNATC	2	1418	2615			
MboI	!GATC	23	586	807	1188	1340	1391
			1400	1413	2593	2597	2616
			2927	3005	3086	3095	3173
			3549	3565	3924	3936	4014
			4022	4033	4108		
MboII	GAAGANNNNNNN!	5	2012	2240	3117	3327	3946
MboII	!NNNNNNNTCTTC	6	658	820	3407	3574	3774
			4017				
McrI	!CGRYCG	4	602	894	2661	4336	
MfeI	!CAATTG	1	1505				
MflI	R!GATCY	12	586	1188	1340	1391	1400
			2927	3173	3565	3924	3936
			4022	4033			
MluI	A!CGCGT	1	1642				
MmeI	!TCCRAC	1	1823				
MmeI	!GTYGGA	2	4299	4483			
MnlI	CCTCNNNNNNN!	9	1475	1484	2539	2545	3357
			3563	3877	4247	4571	
MnlI	!NNNNNNNGAGG	21	541	615	672	873	1223
			1263	1317	1434	1899	2230
			2508	2514	2551	2554	2566
			2606	2670	2806	3163	3593
			4496				
MroI	T!CCGGA	1	1331				
MscI	TGG!CCA	1	2838				
MseI	T!TAA	15	8	1457	1518	1639	1660
			1671	1683	1694	1711	1809
			2080	3899	3913	3918	3970
MspI	C!CGG	22	602	768	1194	1332	1388
			1398	1974	2661	2738	2760
			2788	2919	3009	3076	3257
			3532	3540	3632	4106	4296
			4322	4469			

MstI	TGC!GCA	1	2858					
MstII	CC!TNAGG	2	2236	3868				
MunI	C!AATTG	1	1506					
MvaI	CC!WGG	12	91	284	866	2292	2347	
			2364	3143	3722	3825	4515	
			4528	4649				
MvnI	CG!CG	20	61	1026	1179	1251	1269	
			1382	1644	1680	2056	2076	
			2100	2131	2822	3123	3561	
			3645	3708	3777	4050	4631	
NaeI	GCC!GGC	3	1975	3258	3541			
NarI	GG!CGCC	1	2756					
NciI	CC!SGG	7	1194	1388	1389	2760	2920	
			3533	4297				
NcoI	C!CATGG	4	361	612	2485	3188		
NdeI	CA!TATG	1	235					
NdeII	!GATC	23	586	807	1188	1340	1391	
			1400	1413	2593	2597	2616	
			2927	3005	3086	3095	3173	
			3549	3565	3924	3936	4014	
			4022	4033	4108			
NgoMI	G!CCGGC	3	1973	3256	3539			
NheI	G!CTAGC	1	592					
NlaIII	CATG!	21	305	365	616	655	784	
			931	1024	1066	2186	2326	
			2398	2489	2630	2975	3161	
			3192	3218	3574	3958	4678	
			4729					
NlaIV	GGN!NCC	21	468	631	925	1015	1078	
			1378	1386	1393	1911	1932	
			1944	2135	2296	2368	2757	
			2792	3640	3719	3764	4607	
			4646					
NotI	GC!GGCCGC	0						
NruI	TCG!CGA	0						
NsiI	ATGCA!T	3	2328	2400	4731			
Nsp(7524)I	RCATG!Y	5	1066	2326	2398	3161	4678	
Nsp(7524)V	TT!CGAA	2	1358	3438				
NspBII	CMG!CKG	7	1033	1096	1382	2254	2862	
			4091	4336				
NspII	GDGCH!C	13	753	825	1296	1351	1388	
			1947	2702	2795	2869	3059	
			3121	3831	4364			
NspIII	C!YCGRG	3	878	1344	1387			
NspIV	G!GNCC	16	84	277	629	883	1013	
			1076	1141	1196	1384	1385	
			1863	3272	3717	3763	3821	
			3854					
NunII	GG!CGCC	1	2756					
PacI	TTAAT!TAA	0						
Paer71	C!TCGAG	1	1344					
PalI	GG!CC	31	85	278	630	690	710	
			885	1014	1077	1100	1131	
			1143	1197	1277	1386	1722	
			1864	2519	2525	2534	2577	
			2664	2838	3229	3256	3765	
			3822	3855	4200	4634	4652	
			4663					
PflMI	CCANNNN!NTGG	2	786	1036				

PleI	GAGTCNNNN!	2	1807	2239				
PleI	!NNNNNGACTC	5	405	1329	1815	3421	4298	
PmaCI	CAC!GTG	1	1258					
PmeI	GTTT!AAAC	0						
PmlI	CAC!GTG	1	1258					
PpuMI	RG!GWCCY	0						
PshAI	GACNN!NNGTC	0						
PstI	CTGCA!G	1	1369					
PvuI	CGAT!CG	0						
PvuII	CAG!CTG	4	1033	1096	2254	2862		
RmaI	C!TAG	8	593	1404	1605	2023	2579	
			3594	3928	4181			
RsaI	GT!AC	9	220	245	300	333	384	
			541	815	1378	3062		
RsrI	G!AATTC	1	1360					
RsrII	CG!GWCCG	1	3272					
SacI	GAGCT!C	1	1351					
SacII	CCGC!GG	1	1383					
SalI	G!TCGAC	1	1370					
Sau3AI	!GATC	23	586	807	1188	1340	1391	
			1400	1413	2593	2597	2616	
			2927	3005	3086	3095	3173	
			3549	3565	3924	3936	4014	
			4022	4033	4108			
Sau96I	G!GNCC	16	84	277	629	883	1013	
			1076	1141	1196	1384	1385	
			1863	3272	3717	3763	3821	
			3854					
SauI	CC!TNAGG	2	2236	3868				
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	19	91	284	866	1194	1388	
			1389	2292	2347	2364	2760	
			2920	3143	3533	3722	3825	
			4297	4515	4528	4649		
SduI	GDGCH!C	13	753	825	1296	1351	1388	
			1947	2702	2795	2869	3059	
			3121	3831	4364			
SecI	C!CNNGG	23	361	612	733	864	879	
			1072	1192	1380	1387	2290	
			2362	2485	2520	2529	2578	
			2919	3188	3593	3720	3721	
			3823	3824	4514			
SexI	!CTCGAG	1	1343					
SfaNI	GCATCNNNNN!	10	858	900	987	1308	1572	
			2335	2407	3055	3119	3394	
SfaNI	!NNNNNNNNNGATGC	5	358	2714	2969	3185	4577	
SfcI	C!TRYAG	4	1365	2091	4218	4409		
SfiI	GGCCNNNN!NGCC	1	2531					
SfuI	TT!CGAA	2	1358	3438				
SgrAI	CR!CCGGYG	0						
SinI	G!GWCC	2	3272	3717				
SmaI	CCC!GGG	1	1389					
SnaBI	TAC!GTA	1	341					
SnaI	!GTATAC	0						
SnoI	G!TGCAC	2	749	4360				
SpeI	A!CTAGT	0						
SphI	GCATG!C	3	2326	2398	3161			
SplI	C!GTACG	0						
SrfI	GCCC!GGGC	0						

Sse8387I	CCTGCA!GG	0						
SspI	AAT!ATT	2	1664	2217				
SstI	GAGCT!C	1	1351					
SstII	CCGC!GG	1	1383					
SstIII	!ACGT	16	122	134	175	258	339	
			444	746	1256	1815	1827	
			1870	1980	2871	3058	3839	
			3972					
StuI	AGG!CCT	1	2577					
StyI	C!CWGG	6	361	612	2485	2578	3188	
			3593					
StySJI	!GAGNNNNNGTRC	0						
StySJI	!GYACNNNNNNCTC	0						
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	19	672	894	912	1149	1185	
			1345	1358	1371	1905	2596	
			2869	3025	3049	3085	3247	
			3438	3483	3738	4576		
TaqII	GACCGANNNNNNNNN!	3	1776	2793	3461			
TaqII	!NNNNNNNNNTCGGTC	1	3705					
TaqII	CACCCANNNNNNNNN!	0						
TaqII	!NNNNNNNNNTGGGTG	3	173	2660	3252			
TfiI	G!AWTC	6	1328	3241	3375	3485	3522	
			4700					
ThaI	CG!CG	20	61	1026	1179	1251	1269	
			1382	1644	1680	2056	2076	
			2100	2131	2822	3123	3561	
			3645	3708	3777	4050	4631	
Tsp45I	!GTSAC	4	1207	2045	2876	3182		
TspEI	!AATT	11	19	1360	1506	1570	1648	
			1674	1685	2332	2404	2496	
			3914					
Tth111I	GACN!NNGTC	1	2874					
Tth111II	CAARCANNNNNNNNN!	2	1238	4083				
Tth111III	!NNNNNNNNNTGYTTG	3	3191	4051	4090			
TthHB8I	T!CGA	19	672	894	912	1149	1185	
			1345	1358	1371	1905	2596	
			2869	3025	3049	3085	3247	
			3438	3483	3738	4576		
Van91I	CCANNNN!NTGG	2	786	1036				
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	1403					
XcmI	!CCANNNNNNNNTCG	1	3031					
XcyI	C!CCGGG	1	1387					
XhoI	C!TCGAG	1	1344					
XhoII	R!GATCY	12	586	1188	1340	1391	1400	
			2927	3173	3565	3924	3936	
			4022	4033				
XmaI	C!CCGGG	1	1387					
XmaIII	C!GGCCG	1	2662					
XmnI	GAANN!NNTTC	0						
XorII	CGAT!CG	0						