

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	1	2547				
AatII	GACGT!C	4	126	179	262	448	
Acc65I	G!GTACC	1	1346				
AccI	GT!MKAC	1	1341				
AccII	CG!CG	19	61	960	1116	1221	1352
			1614	1650	2026	2046	2070
			2101	2792	3093	3531	3615
			3678	3747	4020	4601	
AccIII	T!CCGGA	0					
AciI	G!CGG	31	371	404	529	630	702
			705	837	876	915	1122
			1352	1986	2013	2101	2211
			2792	2856	2957	2960	3240
			3295	3393	3531	3678	3818
			4061	4306	4427	4446	4573
			4601				
AciI	C!CGC	42	59	87	99	113	280
			508	590	639	658	730
			892	943	958	1033	1099
			1216	1219	1350	2041	2044
			2058	2148	2396	2408	2417
			2429	2439	2450	2496	2635
			2698	3200	3245	3311	3337
			3462	3465	3613	3745	4052
			4196	4692			
AcyI	GR!CGYC	8	123	176	259	445	762
			2726	3428	3507		
AfaI	GT!AC	10	220	245	300	333	384
			541	905	1265	1348	3032
AflI	G!GWCC	2	3242	3687			
AflIII	C!TTAAG	0					
AflIII	A!CRYGT	2	1612	4644			
AgeI	A!CCGGT	1	601				
AhaI	CC!SGG	7	1286	1358	1359	2730	2890
			3503	4267			
AhaII	GR!CGYC	8	123	176	259	445	762
			2726	3428	3507		
AhaIII	TTT!AAA	3	1428	3870	3889		
AluI	AG!CT	20	567	694	933	939	1092
			1134	1249	1319	1325	1505
			1672	1929	2224	2512	2832
			3290	4087	4344	4390	4480
Alw44I	G!TGCAC	1	4330				
AlwI	GGATCNNNN!	10	1239	1369	1378	2594	2905
			3151	3543	3902	4000	4086
AlwI	!NNNNNGATCC	6	581	1356	2970	3514	3901
			3998				
AlwNI	CAGNNN!CTG	2	3827	4235			
AocI	CC!TNAGG	2	2206	3838			
AocII	GDGCH!C	15	716	755	989	1056	1202
			1321	1358	1917	2672	2765
			2839	3029	3091	3801	4334
Aor51HI	AGC!GCT	1	597				
AosI	TGC!GCA	1	2828				
AosII	GR!CGYC	8	123	176	259	445	762
			2726	3428	3507		

ApaI	GGGCC!C	2	1202	1358				
ApaLI	G!TGCAC	1	4330					
ApoI	R!AATTY	4	1330	1540	1644	1655		
ApyI	CC!WGG	11	91	284	2262	2317	2334	
			3113	3692	3795	4485	4498	
			4619					
AquI	C!YCGRG	4	719	998	1314	1357		
AscI	GG!CGCGCC	0						
AseI	AT!TAAT	1	8					
AsnI	AT!TAAT	1	8					
Asp700I	GAANN!NNTTC	1	851					
Asp718I	G!GTACC	1	1346					
AspHI	GWGCW!C	6	755	1056	1321	2839	3029	
			4334					
AspI	GACN!NNGTC	1	2844					
AsuI	G!GNCC	13	84	277	727	1198	1199	
			1354	1355	1833	3242	3687	
			3733	3791	3824			
AsuII	TT!CGAA	2	1328	3408				
AvaI	C!YCGRG	4	719	998	1314	1357		
AvaII	G!GWCC	2	3242	3687				
AvaIII	ATGCA!T	3	2298	2370	4701			
AviII	TGC!GCA	1	2828					
AvrI	!CYCGRG	4	718	997	1313	1356		
AvrII	C!CTAGG	2	2548	3563				
AxyI	CC!TNAGG	2	2206	3838				
BalI	TGG!CCA	1	2808					
BamHI	G!GATCC	1	1361					
BanI	G!GYRCC	10	466	668	713	761	824	
			986	1346	1879	2725	2760	
BanII	GRGCY!C	6	1202	1321	1358	1917	3091	
			3801					
BanIII	AT!CGAT	1	2566					
BbeI	GGCGC!C	2	765	2729				
BbiIII/AcyI	GR!CGYC	8	123	176	259	445	762	
			2726	3428	3507			
BbrPI	CAC!GTG	2	796	928				
BbsI	!GTCTTC	0						
BbsI	!GAAGAC	0						
BbvI	GCAGCNNNNNNNN!	8	1047	1101	1514	2799	2841	
			3362	3815	4315			
BbvI	!NNNNNNNNNNNGCTGC	8	909	2008	2673	2857	2950	
			4016	4222	4225			
BbvII	GAAGACNN!	0						
BbvII	!NNNNNNGTCTTC	0						
BcefI	ACGGCNNNNNNNNNN!	7	694	823	889	928	1135	
			3152	3639				
BcefI	!NNNNNNNNNNNGCCGT	3	1868	2686	4145			
BcgI	!GCANNNNNNTCG	0						
BcgI	!CGANNNNNNTGC	0						
BclI	T!GATCA	1	1383					
BcnI	CC!SGG	7	1286	1358	1359	2730	2890	
			3503	4267				
BfaI	C!TAG	8	593	1374	1575	1993	2549	
			3564	3898	4151			
BfrI	C!TTAAG	0						
BglI	GCCNNNN!NGGC	4	91	213	284	2501		
BglII	A!GATCT	1	1310					
BinI	!NNNNNGATCC	6	581	1356	2970	3514	3901	

			3998					
BinI	GGATCNNNN!	10	1239	1369	1378	2594	2905	
			3151	3543	3902	4000	4086	
BlnI	C!CTAGG	2	2548	3563				
BmyI	GDGCH!C	15	716	755	989	1056	1202	
			1321	1358	1917	2672	2765	
			2839	3029	3091	3801	4334	
BpmI	!CTCCAG	1	3523					
BpmI	!CTGGAG	1	3544					
Bpu1102I	GC!TNAGC	1	788					
BsaAI	YAC!GTR	5	341	796	928	1842	3030	
BsaBI	GATNN!NNATC	3	1077	1388	2585			
BsaHI	GR!CGYC	8	123	176	259	445	762	
			2726	3428	3507			
BsaI	GGTCTCN!	0						
BsaI	!NNNNNGAGACC	1	3715					
BsaJI	C!CNNGG	24	361	612	642	999	1047	
			1065	1137	1284	1350	1357	
			2260	2332	2455	2490	2499	
			2548	2889	3158	3563	3690	
			3691	3793	3794	4484		
BseAI	T!CCGGA	0						
BsePI	!GCGCGC	0						
BsgI	!GTGCAG	0						
BsgI	!CTGCAC	3	856	922	1057			
BsiEI	CGRY!CG	4	606	958	2635	4310		
BsiWI	C!GTACG	0						
BsiYI	CCNNNNN!NNGG	24	50	724	725	799	949	
			1222	1225	1738	2064	2456	
			2707	3251	3532	3564	3565	
			3696	3731	3781	3782	3813	
			4166	4445	4611	4629		
BsII	CCNNNNN!NNGG	24	50	724	725	799	949	
			1222	1225	1738	2064	2456	
			2707	3251	3532	3564	3565	
			3696	3731	3781	3782	3813	
			4166	4445	4611	4629		
BsmAI	!GTCTC	1	429					
BsmAI	!GAGAC	4	2155	2573	3592	3720		
BsmI	GAATGCN!	1	1477					
BsmI	G!CATTC	1	1570					
Bsp1286I	GDGCH!C	15	716	755	989	1056	1202	
			1321	1358	1917	2672	2765	
			2839	3029	3091	3801	4334	
BspDI	AT!CGAT	1	2566					
BspEI	T!CCGGA	0						
BspHI	T!CATGA	2	2152	3924				
BspMI	ACCTGCNNNN!	4	792	1065	2994	3444		
BspMI	!NNNNNNNNGCAGGT	1	2613					
BspMII	T!CCGGA	0						
BspWI	GCNNNNN!NNGC	37	91	213	245	284	377	
			401	627	666	711	759	
			966	1098	1107	1122	1131	
			1953	1983	2010	2052	2054	
			2293	2365	2416	2495	2501	
			2717	2801	2824	2963	2969	
			3086	3169	3436	3623	3827	
			4026	4598				
BsrFI	R!CCGGY	9	601	665	953	989	1298	

			1943	3045	3226	3509	
BsrI	ACTGGN!	3	2670	2871	4117		
BsrI	C!CAGT	5	296	1755	2433	4229	4242
BssHII	G!CGCGC	0					
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	2	1328	3408			
BstEII	G!GTNACC	0					
BstI	G!GATCC	1	1361				
BstNI	CC!WGG	11	91	284	2262	2317	2334
			3113	3692	3795	4485	4498
			4619				
BstPI	G!GTNACC	0					
BstUI	CG!CG	19	61	960	1116	1221	1352
			1614	1650	2026	2046	2070
			2101	2792	3093	3531	3615
			3678	3747	4020	4601	
BstXI	CCANNNNN!NTGG	0					
BstYI	R!GATCY	11	586	1310	1361	1370	2897
			3143	3535	3894	3906	3992
			4003				
Bsu36I	CC!TNAGG	2	2206	3838			
CbiI	TT!CGAA	2	1328	3408			
CcrI	C!TCGAG	1	1314				
CelII	GC!TNAGC	1	788				
CfoI	GCG!C	24	598	764	1116	1169	1992
			2000	2026	2048	2057	2070
			2101	2720	2728	2792	2829
			3095	3355	3531	3617	4020
			4129	4303	4403	4470	
Cfr10I	R!CCGGY	9	601	665	953	989	1298
			1943	3045	3226	3509	
Cfr13I	G!GNCC	13	84	277	727	1198	1199
			1354	1355	1833	3242	3687
			3733	3791	3824		
Cfr9I	C!CCGGG	1	1357				
CfrI	Y!GGCCR	5	955	2632	2806	3197	3224
ClaI	AT!CGAT	1	2566				
CpoI	CG!GWCCG	1	3242				
Csp45I	TT!CGAA	2	1328	3408			
CspI	CG!GSCCG	0					
CviJI	RG!CY	95	43	85	94	278	567
			633	694	729	777	792
			804	810	840	879	933
			939	953	957	993	1092
			1110	1125	1134	1149	1200
			1249	1319	1325	1356	1395
			1505	1672	1692	1834	1915
			1929	1943	2224	2265	2337
			2460	2489	2495	2504	2512
			2535	2547	2553	2634	2651
			2659	2686	2711	2795	2804
			2808	2832	2870	2946	2963
			2982	3045	3089	3099	3116
			3199	3226	3230	3267	3290
			3306	3486	3513	3574	3696
			3735	3792	3799	3806	3825
			3836	4087	4116	4159	4170
			4235	4314	4339	4344	4390
			4480	4578	4604	4622	4633

CvnI	CC!TNAGG	2	2206	3838				
DdeI	C!TNAG	7	788	2206	2508	3389	3838	
			3961	4370				
DpnI	GA!TC	27	588	650	965	1013	1028	
			1080	1233	1312	1363	1372	
			1385	2565	2569	2588	2899	
			2977	3058	3067	3145	3521	
			3537	3896	3908	3986	3994	
			4005	4080				
DpnII	!GATC	27	586	648	963	1011	1026	
			1078	1231	1310	1361	1370	
			1383	2563	2567	2586	2897	
			2975	3056	3065	3143	3519	
			3535	3894	3906	3984	3992	
			4003	4078				
DraI	TTT!AAA	3	1428	3870	3889			
DraII	RG!GNCCY	3	1198	1199	3824			
DraIII	CACNNN!GTG	1	1842					
DrdI	GACNNNN!NNGTC	3	1797	2753	4542			
DsaI	C!CRYGG	9	361	612	642	1047	1065	
			1137	1350	2455	3158		
DsaV	!CCNGG	18	89	282	1284	1356	1357	
			2260	2315	2332	2728	2888	
			3111	3501	3690	3793	4265	
			4483	4496	4617			
EaeI	Y!GGCCR	5	955	2632	2806	3197	3224	
EagI	C!GGCCG	2	955	2632				
Eam1105I	GACNNN!NNGTC	0						
EarI	!CTCTTC	0						
EarI	!GAAGAG	4	747	2197	3074	3284		
Ecl136II	GAG!CTC	1	1319					
EclXI	C!GGCCG	2	955	2632				
Eco105I	TAC!GTA	1	341					
Eco31I	GGTCTCN!	0						
Eco31I	!NNNNNGAGACC	1	3715					
Eco47I	G!GWCC	2	3242	3687				
Eco47III	AGC!GCT	1	597					
Eco52I	C!GGCCG	2	955	2632				
Eco57I	!CTGAAG	2	2850	3282				
Eco57I	!CTTCAG	4	771	933	1104	4116		
Eco81I	CC!TNAGG	2	2206	3838				
EcoNI	CCTNN!NNNAGG	0						
EcoO109I	RG!GNCCY	3	1198	1199	3824			
EcoO65I	G!GTNACC	0						
EcoRI	G!AATTC	1	1330					
EcoRII	!CCWGG	11	89	282	2260	2315	2332	
			3111	3690	3793	4483	4496	
			4617					
EcoRV	GAT!ATC	0						
EcoT14I	C!CWGG	8	361	612	642	1065	2455	
			2548	3158	3563			
EcoT22I	ATGCA!T	3	2298	2370	4701			
EcoT38I	GRGCY!C	6	1202	1321	1358	1917	3091	
			3801					
EheI	GGC!GCC	2	763	2727				
Esp3I	CGTCTCN!	0						
Esp3I	!NNNNNGAGACG	0						
EspI	GC!TNAGC	1	788					
FbaI	T!GATCA	1	1383					

FinI	!GTCCC	4	2256	2328	2392	3687	
FinI	!GGGAC	6	161 3503	312	480	2862	3394
Fnu4HI	GC!NGC	41	631 877 1090 2044 2698 2958 3337 4030 4602	658 916 1123 2058 2788 2961 3351 4236	703 923 1219 2496 2793 2964 3465 4239	730 958 1503 2635 2830 3200 3804 4304	838 1036 2022 2687 2871 3296 3819 4447
FokI	GGATGNNNNNNNNN!	6	1097 3529	1301	2593	3050	3075
FokI	!NNNNNNNNNNNNNCATCC	4	1016	1157	1169	2399	
FspI	TGC!GCA	1	2828				
GdiII	!NNNNNYGGCCG	4	949	2626	3191	3218	
GdiII	CGGCCRN!	2	961	2638			
GsuI	!CTCCAG	1	3523				
GsuI	!CTGGAG	1	3544				
HaeI	WGG!CCW	5	2547	2808	4170	4622	4633
HaeII	RGCGC!Y	7	599 2729	765 4404	1170	1993	2001
HaeIII	GG!CC	25	85 957 2489 2808 3825	278 1200 2495 3199 4170	633 1356 2504 3226 4604	729 1692 2547 3735 4622	953 1834 2634 3792 4633
HapII	C!CGG	24	602 1299 2708 3046 4076	666 1358 2730 3227 4266	954 1368 2758 3502 4292	990 1944 2889 3510 4439	1286 2631 2979 3602
HgaI	GACGCNNNNN!	3	527	3436	3515		
HgaI	!NNNNNNNNNNNGCGTC	3	2059	3954	4532		
HgiAI	GWGCW!C	6	755 4334	1056	1321	2839	3029
HgiEII	!ACCNNNNNNNGGT	1	4056				
HhaI	GCG!C	24	598 2000 2101 3095 4129	764 2026 2720 3355 4303	1116 2048 2728 3531 4403	1169 2057 2792 3617 4470	1992 2070 2829 4020
Hin1I	GR!CGYC	8	123 2726	176 3428	259 3507	445	762
HinP1I	G!CGC	24	596 1998 2099 3093 4127	762 2024 2718 3353 4301	1114 2046 2726 3529 4401	1167 2055 2790 3615 4468	1990 2068 2827 4018
HincII	GTY!RAC	2	1342	1489			
HindIII	A!AGCTT	1	1323				
HinfI	G!ANTC	11	411 3345 4670	1769 3397	1791 3455	2201 3492	3211 4274
HpaI	GTT!AAC	1	1489				
HpaII	C!CGG	24	602 1299 2708	666 1358 2730	954 1368 2758	990 1944 2889	1286 2631 2979

			3046	3227	3502	3510	3602
			4076	4266	4292	4439	
HphI	GGTGANNNNNNNN!	5	377	991	1313	3797	3913
HphI	!NNNNNNNTCACC	5	655	1009	1087	1843	2904
KasI	G!GCGCC	2	761	2725			
KpnI	GGTAC!C	1	1350				
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	4	743	2193	3070	3280	
KspI	CCGC!GG	1	1353				
MaeI	C!TAG	8	593	1374	1575	1993	2549
			3564	3898	4151		
MaeII	A!CGT	16	123	135	176	259	340
			445	795	927	1786	1798
			1841	1951	2842	3029	3810
			3943				
MaeIII	!GTNAC	12	62	149	498	1514	2015
			2027	2846	3152	3842	4108
			4224	4287			
MamI	GATNN!NNATC	3	1077	1388	2585		
MboI	!GATC	27	586	648	963	1011	1026
			1078	1231	1310	1361	1370
			1383	2563	2567	2586	2897
			2975	3056	3065	3143	3519
			3535	3894	3906	3984	3992
			4003	4078			
MboII	GAAGANNNNNNNN!	7	760	1316	1982	2210	3087
			3297	3916			
MboII	!NNNNNNNTCTTC	5	1006	3377	3544	3744	3987
McrI	!CGRYCG	4	602	954	2631	4306	
MfeI	!CAATTG	1	1475				
MflI	R!GATCY	11	586	1310	1361	1370	2897
			3143	3535	3894	3906	3992
			4003				
MluI	A!GCGGT	1	1612				
MmeI	!TCCRAC	1	1793				
MmeI	!GTYGGA	2	4269	4453			
MnlI	CCTCNNNNNNN!	9	1445	1454	2509	2515	3327
			3533	3847	4217	4541	
MnlI	!NNNNNNNGAGG	21	541	702	900	942	993
			1218	1221	1404	1869	2200
			2478	2484	2521	2524	2536
			2576	2640	2776	3133	3563
			4466				
MroI	T!CCGGA	0					
MscI	TGG!CCA	1	2808				
MseI	T!TAA	15	8	1427	1488	1609	1630
			1641	1653	1664	1681	1779
			2050	3869	3883	3888	3940
MspI	C!CGG	24	602	666	954	990	1286
			1299	1358	1368	1944	2631
			2708	2730	2758	2889	2979
			3046	3227	3502	3510	3602
			4076	4266	4292	4439	
MstI	TGC!GCA	1	2828				
MstII	CC!TNAGG	2	2206	3838			
MunI	C!AATTG	1	1476				
MvaI	CC!WGG	11	91	284	2262	2317	2334
			3113	3692	3795	4485	4498
			4619				

MvnI	CG!CG	19	61 1614 2101 3678	960 1650 2792 3747	1116 2026 3093 4020	1221 2046 3531 4601	1352 2070 3615
NaeI	GCC!GGC	4	955	1945	3228	3511	
NarI	GG!CGCC	2	762	2726			
NciI	CC!SGG	7	1286 3503	1358 4267	1359	2730	2890
NcoI	C!CATGG	6	361 3158	612	642	1065	2455
NdeI	CA!TATG	1	235				
NdeII	!GATC	27	586 1078 1383 2975 3535 4003	648 1231 2563 3056 3894 4078	963 1310 2567 3065 3906	1011 1361 2586 3143 3984	1026 1370 2897 3519 3992
NgoMI	G!CCGGC	4	953	1943	3226	3509	
NheI	G!CTAGC	1	592				
NlaIII	CATG!	21	305 1069 2368 3162 4699	365 1156 2459 3188	616 1207 2600 3544	646 2156 2945 3928	736 2296 3131 4648
NlaIV	GGN!NCC	25	468 988 1356 2105 3610	670 1199 1363 2266 3689	715 1200 1881 2338 3734	763 1201 1902 2727 4577	826 1348 1914 2762 4616
NotI	GC!GGCCGC	0					
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	2298	2370	4701		
Nsp(7524)I	RCATG!Y	5	1156	2296	2368	3131	4648
Nsp(7524)V	TT!CGAA	2	1328	3408			
NspBII	CMG!CKG	6	837 4306	1352	2224	2832	4061
NspII	GDGCH!C	15	716 1321 2839	755 1358 3029	989 1917 3091	1056 2672 3801	1202 2765 4334
NspIII	C!YCGRG	4	719	998	1314	1357	
NspIV	G!GNCC	13	84 1354 3733	277 1355 3791	727 1833 3824	1198	1199 3687
NunII	GG!CGCC	2	762	2726			
PacI	TTAAT!TAA	0					
Paer71	C!TCGAG	1	1314				
PalI	GG!CC	25	85 957 2489 2808 3825	278 1200 2495 3199 4170	633 1356 2504 3226 4604	729 1692 2547 3735 4622	953 1834 2634 3792 4633
PflMI	CCANNNN!NTGG	1	799				
PleI	GAGTCNNNN!	2	1777	2209			
PleI	!NNNNNGACTC	4	405	1785	3391	4268	
PmaCI	CAC!GTG	2	796	928			
PmeI	GTTT!AAAC	0					
PmlI	CAC!GTG	2	796	928			
PpuMI	RG!GWCCY	0					
PshAI	GACNN!NNGTC	0					

PstI	CTGCA!G	2	1191	1339				
PvuI	CGAT!CG	0						
PvuII	CAG!CTG	2	2224	2832				
RmaI	C!TAG	8	593	1374	1575	1993	2549	
			3564	3898	4151			
RsaI	GT!AC	10	220	245	300	333	384	
			541	905	1265	1348	3032	
RsrI	G!AATTC	1	1330					
RsrII	CG!GWCCG	1	3242					
SacI	GAGCT!C	1	1321					
SacII	CCGC!GG	1	1353					
SalI	G!TCGAC	1	1340					
Sau3AI	!GATC	27	586	648	963	1011	1026	
			1078	1231	1310	1361	1370	
			1383	2563	2567	2586	2897	
			2975	3056	3065	3143	3519	
			3535	3894	3906	3984	3992	
			4003	4078				
Sau96I	G!GNCC	13	84	277	727	1198	1199	
			1354	1355	1833	3242	3687	
			3733	3791	3824			
SauI	CC!TNAGG	2	2206	3838				
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	18	91	284	1286	1358	1359	
			2262	2317	2334	2730	2890	
			3113	3503	3692	3795	4267	
			4485	4498	4619			
SduI	GDGCH!C	15	716	755	989	1056	1202	
			1321	1358	1917	2672	2765	
			2839	3029	3091	3801	4334	
SecI	C!CNNGG	24	361	612	642	999	1047	
			1065	1137	1284	1350	1357	
			2260	2332	2455	2490	2499	
			2548	2889	3158	3563	3690	
			3691	3793	3794	4484		
SexI	!CTCGAG	1	1313					
SfaNI	GCATCNNNNN!	10	669	903	1191	1263	1542	
			2305	2377	3025	3089	3364	
SfaNI	!NNNNNNNNNGATGC	7	358	1279	1285	2684	2939	
			3155	4547				
SfcI	C!TRYAG	6	1129	1187	1335	2061	4188	
			4379					
SfiI	GGCCNNNN!NGGCC	1	2501					
SfuI	TT!CGAA	2	1328	3408				
SgrAI	CR!CCGGYG	0						
SinI	G!GWCC	2	3242	3687				
SmaI	CCC!GGG	1	1359					
SnaBI	TAC!GTA	1	341					
SnaI	!GTATAC	0						
SnoI	G!TGCAC	1	4330					
SpeI	A!CTAGT	0						
SphI	GCATG!C	3	2296	2368	3131			
SplI	C!GTACG	0						
SrfI	GCCC!GGGC	0						
Sse8387I	CCTGCA!GG	0						
SspI	AAT!ATT	2	1634	2187				
SstI	GAGCT!C	1	1321					
SstII	CCGC!GG	1	1353					
SstIII	!ACGT	16	122	134	175	258	339	

			444	794	926	1785	1797
			1840	1950	2841	3028	3809
			3942				
StuI	AGG!CCT	1	2547				
StyI	C!CWWGG	8	361	612	642	1065	2455
			2548	3158	3563		
StySJI	!GAGNNNNNNGTRC	0					
StySJI	!GYACNNNNNNCTC	0					
SwaI	ATTT!AAAT	0					
TaqI	T!CGA	17	651	690	897	1315	1328
			1341	1875	2566	2839	2995
			3019	3055	3217	3408	3453
			3708	4546			
TaqII	GACCGANNNNNNNNNN!	3	1746	2763	3431		
TaqII	!NNNNNNNNNTCGGTC	1	3675				
TaqII	CACCCANNNNNNNNNN!	0					
TaqII	!NNNNNNNNNTGGGTG	3	173	2630	3222		
TfiI	G!AWTC	5	3211	3345	3455	3492	4670
ThaI	CG!CG	19	61	960	1116	1221	1352
			1614	1650	2026	2046	2070
			2101	2792	3093	3531	3615
			3678	3747	4020	4601	
Tsp45I	!GTSAC	3	2015	2846	3152		
TspEI	!AATT	11	19	1330	1476	1540	1618
			1644	1655	2302	2374	2466
			3884				
Tth111I	GACN!NNGTC	1	2844				
Tth111II	CAARCANNNNNNNNNN!	1	4053				
Tth111III	!NNNNNNNNNTGYTTG	3	3161	4021	4060		
TthHB8I	T!CGA	17	651	690	897	1315	1328
			1341	1875	2566	2839	2995
			3019	3055	3217	3408	3453
			3708	4546			
Van91I	CCANNNN!NTGG	1	799				
VspI	AT!TAAT	1	8				
XbaI	T!CTAGA	1	1373				
XcmI	!CCANNNNNNNNTCG	2	884	3001			
XcyI	C!CCGGG	1	1357				
XhoI	C!TCGAG	1	1314				
XhoII	R!GATCY	11	586	1310	1361	1370	2897
			3143	3535	3894	3906	3992
			4003				
XmaI	C!CCGGG	1	1357				
XmaIII	C!GGCCG	2	955	2632			
XmnI	GAANN!NNTTC	1	851				
XorII	CGAT!CG	0					