

Enzyme Name	Sequence	Count	Cutting Position					
AatI	AGG!CCT	1	2650					
AatII	GACGT!C	4	126	179	262	448		
Acc65I	G!GTACC	0						
AccI	GT!MKAC	0						
AccII	CG!CG	22	61	663	665	794	1025	
			1064	1349	1379	1478	1753	
			2129	2149	2173	2204	2895	
			3196	3634	3718	3781	3850	
			4123	4704				
AccIII	T!CCGGA	0						
AciI	G!CGG	29	371	404	529	624	648	
			713	794	1271	1274	2089	
			2116	2204	2314	2895	2959	
			3060	3063	3343	3398	3496	
			3634	3781	3921	4164	4409	
			4530	4549	4676	4704		
AciI	C!CGC	38	59	87	99	113	280	
			508	590	613	661	792	
			1023	1347	1392	1476	2144	
			2147	2161	2251	2499	2511	
			2520	2532	2542	2553	2599	
			2738	2801	3303	3348	3414	
			3440	3565	3568	3716	3848	
			4155	4299	4795			
AcyI	GR!CGYC	10	123	176	259	445	610	
			716	905	2829	3531	3610	
AfaI	GT!AC	11	220	245	300	333	384	
			541	925	1057	1150	1297	
			3135					
AflI	G!GWCC	2	3345	3790				
AflII	C!TTAAG	1	1711					
AflIII	A!CRYGT	1	4747					
AgeI	A!CCGGT	1	693					
AhaI	CC!SGG	5	645	2833	2993	3606	4370	
AhaII	GR!CGYC	10	123	176	259	445	610	
			716	905	2829	3531	3610	
AhaIII	TTT!AAA	3	1531	3973	3992			
AluI	AG!CT	20	567	936	956	1071	1310	
			1323	1412	1421	1426	1608	
			1775	2032	2327	2615	2935	
			3393	4190	4447	4493	4583	
Alw44I	G!TGCAC	2	1282	4433				
AlwI	GGATCNNNN!	8	695	2697	3008	3254	3646	
			4005	4103	4189			
AlwI	!NNNNNGATCC	9	581	666	682	732	1439	
			3073	3617	4004	4101		
AlwNI	CAGNNN!CTG	2	3930	4338				
AocI	CC!TNAGG	2	2309	3941				
AocII	GDGCH!C	13	866	872	880	1286	1292	
			2020	2775	2868	2942	3132	
			3194	3904	4437			
Aor51HI	AGC!GCT	0						
AosI	TGC!GCA	1	2931					
AosII	GR!CGYC	10	123	176	259	445	610	
			716	905	2829	3531	3610	
ApaI	GGGCC!C	0						

ApaI	G!TGCAC	2	1282	4433				
ApoI	R!AATTY	3	1643	1747	1758			
ApyI	CC!WGG	14	91	284	873	883	1081	
			2365	2420	2437	3216	3795	
			3898	4588	4601	4722		
AquI	C!YCGRG	2	931	1291				
AscI	GG!CGCGCC	0						
AseI	AT!TAAT	1	8					
AsnI	AT!TAAT	1	8					
Asp700I	GAANN!NNTTC	1	943					
Asp718I	G!GTACC	0						
AspHI	GWGCW!C	5	880	1286	2942	3132	4437	
AspI	GACN!NNGTC	1	2947					
AsuI	G!GNCC	11	84	277	642	928	1276	
			1936	3345	3790	3836	3894	
			3927					
AsuII	TT!CGAA	1	3511					
AvaI	C!YCGRG	2	931	1291				
AvaII	G!GWCC	2	3345	3790				
AvaIII	ATGCA!T	3	2401	2473	4804			
AviII	TGC!GCA	1	2931					
AvrI	!CYCGRG	2	930	1290				
AvrII	C!CTAGG	2	2651	3666				
AxyI	CC!TNAGG	2	2309	3941				
BalI	TGG!CCA	4	776	1079	1100	2911		
BamHI	G!GATCC	1	687					
BanI	G!GYRCC	6	466	715	904	1982	2828	
			2863					
BanII	GRGCY!C	3	2020	3194	3904			
BanIII	AT!CGAT	1	2669					
BbeI	GGCGC!C	3	719	908	2832			
BbiII/AcyI	GR!CGYC	10	123	176	259	445	610	
			716	905	2829	3531	3610	
BbrPI	CAC!GTG	3	1104	1287	1317			
BbsI	!GTCTTC	0						
BbsI	!GAAGAC	0						
BbvI	GCAGCNNNNNNNN!	9	722	1061	1229	1617	2902	
			2944	3465	3918	4418		
BbvI	!NNNNNNNNNNNGCTGC	11	602	605	608	943	2111	
			2776	2960	3053	4119	4325	
			4328					
BbvII	GAAGACNN!	0						
BbvII	!NNNNNNGTCTTC	0						
BcefI	ACGGCNNNNNNNNNN!	7	747	822	918	942	1191	
			3255	3742				
BcefI	!NNNNNNNNNNNGCCGT	4	1367	1971	2789	4248		
BcgI	!GCANNNNNTCG	0						
BcgI	!CGANNNNNTGC	0						
BclI	T!GATCA	0						
BcnI	CC!SGG	5	645	2833	2993	3606	4370	
BfaI	C!TAG	8	593	1484	1678	2096	2652	
			3667	4001	4254			
BfrI	C!TTAAG	1	1711					
BglI	GCCNNNN!NGGC	4	91	213	284	2604		
BglII	A!GATCT	1	1196					
BinI	!NNNNNGATCC	9	581	666	682	732	1439	
			3073	3617	4004	4101		
BinI	GGATCNNNN!	8	695	2697	3008	3254	3646	
			4005	4103	4189			

BlnI	C!CTAGG	2	2651	3666				
BmyI	GDGCH!C	13	866	872	880	1286	1292	
			2020	2775	2868	2942	3132	
			3194	3904	4437			
BpmI	!CTCCAG	1	3626					
BpmI	!CTGGAG	3	872	1119	3647			
Bpu1102I	GC!TNAGC	2	1324	1422				
BsaAI	YAC!GTR	8	341	747	975	1104	1287	
			1317	1945	3133			
BsaBI	GATNN!NNATC	2	1491	2688				
BsaHI	GR!CGYC	10	123	176	259	445	610	
			716	905	2829	3531	3610	
BsaI	GGTCTCN!	0						
BsaI	!NNNNNGAGACC	2	1363	3818				
BsaJI	C!CNNGG	22	361	729	792	871	881	
			1028	1080	1382	2363	2435	
			2558	2593	2602	2651	2992	
			3261	3666	3793	3794	3896	
			3897	4587				
BseAI	T!CCGGA	0						
BsePI	!GCGCGC	2	662	1376				
BsgI	!GTGCAG	1	975					
BsgI	!CTGCAC	1	845					
BsiEI	CGRY!CG	3	698	2738	4413			
BsiWI	C!GTACG	0						
BsiYI	CCNNNNN!NNGG	25	50	681	747	798	872	
			1029	1107	1270	1841	2167	
			2559	2810	3354	3635	3667	
			3668	3799	3834	3884	3885	
			3916	4269	4548	4714	4732	
BslI	CCNNNNN!NNGG	25	50	681	747	798	872	
			1029	1107	1270	1841	2167	
			2559	2810	3354	3635	3667	
			3668	3799	3834	3884	3885	
			3916	4269	4548	4714	4732	
BsmAI	!GTCTC	1	429					
BsmAI	!GAGAC	5	1368	2258	2676	3695	3823	
BsmI	GAATGCN!	1	1580					
BsmI	G!CATTC	1	1673					
Bsp1286I	GDGCH!C	13	866	872	880	1286	1292	
			2020	2775	2868	2942	3132	
			3194	3904	4437			
BspDI	AT!CGAT	1	2669					
BspEI	T!CCGGA	0						
BspHI	T!CATGA	2	2255	4027				
BspMI	ACCTGCNNNN!	2	3097	3547				
BspMI	!NNNNNNNNGCAGGT	1	2716					
BspMII	T!CCGGA	0						
BspWI	GCNNNNN!NNGC	40	91	213	245	284	377	
			401	618	621	635	657	
			794	800	815	854	911	
			1184	1223	1418	2056	2086	
			2113	2155	2157	2396	2468	
			2519	2598	2604	2820	2904	
			2927	3066	3072	3189	3272	
			3539	3726	3930	4129	4701	
BsrFI	R!CCGGY	7	693	853	1213	2046	3148	
			3329	3612				
BsrI	ACTGGN!	3	2773	2974	4220			

BsrI	C!CAGT	9	296	655	837	909	1448
			1858	2536	4332	4345	
BssHII	G!CGCGC	2	663	1377			
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	1	3511				
BstEII	G!GTNACC	2	885	1032			
BstI	G!GATCC	1	687				
BstNI	CC!WGG	14	91	284	873	883	1081
			2365	2420	2437	3216	3795
			3898	4588	4601	4722	
BstPI	G!GTNACC	2	885	1032			
BstUI	CG!CG	22	61	663	665	794	1025
			1064	1349	1379	1478	1753
			2129	2149	2173	2204	2895
			3196	3634	3718	3781	3850
			4123	4704			
BstXI	CCANNNNN!NTGG	0					
BstYI	R!GATCY	13	586	671	687	737	1196
			1444	3000	3246	3638	3997
			4009	4095	4106		
Bsu36I	CC!TNAGG	2	2309	3941			
CbiI	TT!CGAA	1	3511				
CcrI	C!TCGAG	0					
CelII	GC!TNAGC	2	1324	1422			
CfoI	GCG!C	29	665	667	718	907	986
			1027	1189	1379	1381	2095
			2103	2129	2151	2160	2173
			2204	2823	2831	2895	2932
			3198	3458	3634	3720	4123
			4232	4406	4506	4573	
Cfr10I	R!CCGGY	7	693	853	1213	2046	3148
			3329	3612			
Cfr13I	G!GNCC	11	84	277	642	928	1276
			1936	3345	3790	3836	3894
			3927				
Cfr9I	C!CCGGG	0					
CfrI	Y!GGCCR	9	774	1077	1098	1238	1473
			2735	2909	3300	3327	
ClaI	AT!CGAT	1	2669				
CpoI	CG!GWCCG	1	3345				
Csp45I	TT!CGAA	1	3511				
CspI	CG!GSCCG	1	642				
CviJI	RG!CY	96	43	85	94	278	567
			629	638	643	651	776
			803	929	936	956	971
			1071	1079	1085	1100	1168
			1178	1240	1277	1310	1323
			1361	1412	1421	1426	1475
			1498	1608	1775	1795	1937
			2018	2032	2046	2327	2368
			2440	2563	2592	2598	2607
			2615	2638	2650	2656	2737
			2754	2762	2789	2814	2898
			2907	2911	2935	2973	3049
			3066	3085	3148	3192	3202
			3219	3302	3329	3333	3370
			3393	3409	3589	3616	3677
			3799	3838	3895	3902	3909
			3928	3939	4190	4219	4262

			4273	4338	4417	4442	4447
			4493	4583	4681	4707	4725
			4736				
CvnI	CC!TNAGG	2	2309	3941			
DdeI	C!TNAG	9	1324	1422	1459	2309	2611
			3492	3941	4064	4473	
DpnI	GA!TC	27	588	673	689	739	1163
			1198	1210	1244	1343	1446
			1488	2668	2672	2691	3002
			3080	3161	3170	3248	3624
			3640	3999	4011	4089	4097
			4108	4183			
DpnII	!GATC	27	586	671	687	737	1161
			1196	1208	1242	1341	1444
			1486	2666	2670	2689	3000
			3078	3159	3168	3246	3622
			3638	3997	4009	4087	4095
			4106	4181			
DraI	TTT!AAA	3	1531	3973	3992		
DraII	RG!GNCCY	1	3927				
DraIII	CACNNN!GTG	1	1945				
DrdI	GACNNNN!NNGTC	3	1900	2856	4645		
DsaI	C!CRYGG	6	361	729	792	1382	2558
			3261				
DsaV	!CCNGG	19	89	282	643	871	881
			1079	2363	2418	2435	2831
			2991	3214	3604	3793	3896
			4368	4586	4599	4720	
EaeI	Y!GGCCR	9	774	1077	1098	1238	1473
			2735	2909	3300	3327	
EagI	C!GGCCG	1	2735				
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	4	1181	2300	3177	3387	
Ecl136II	GAG!CTC	0					
EclXI	C!GGCCG	1	2735				
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	2	1363	3818			
Eco47I	G!GWCC	2	3345	3790			
Eco47III	AGC!GCT	0					
Eco52I	C!GGCCG	1	2735				
Eco57I	!CTGAAG	5	936	1179	1362	2953	3385
Eco57I	!CTTCAG	2	782	4219			
Eco81I	CC!TNAGG	2	2309	3941			
EcoNI	CCTNN!NNNAGG	0					
EcoO109I	RG!GNCCY	1	3927				
EcoO65I	G!GTNACC	2	885	1032			
EcoRI	G!AATTC	0					
EcoRII	!CCWGG	14	89	282	871	881	1079
			2363	2418	2435	3214	3793
			3896	4586	4599	4720	
EcoRV	GAT!ATC	0					
EcoT14I	C!CWGGG	5	361	2558	2651	3261	3666
EcoT22I	ATGCA!T	3	2401	2473	4804		
EcoT38I	GRGCY!C	3	2020	3194	3904		
EheI	GGC!GCC	3	717	906	2830		
Esp3I	CGTCTCN!	0					
Esp3I	!NNNNNGAGACG	0					

EspI	GC!TNAGC	2	1324	1422				
FbaI	T!GATCA	0						
FinI	!GTCCC	4	2359	2431	2495	3790		
FinI	!GGGAC	6	161	312	480	2965	3497	
			3606					
Fnu4HI	GC!NGC	44	613	616	619	622	649	
			661	711	714	795	957	
			1050	1218	1272	1275	1392	
			1476	1606	2125	2147	2161	
			2599	2738	2790	2801	2891	
			2896	2933	2974	3061	3064	
			3067	3303	3399	3440	3454	
			3568	3907	3922	4133	4339	
			4342	4407	4550	4705		
FokI	GGATGNNNNNNNNN!	8	784	844	1108	1345	2696	
			3153	3178	3632			
FokI	!NNNNNNNNNNNNNNCATCC	2	775	2502				
FspI	TGC!GCA	1	2931					
GdiII	!NNNNNYGGCCG	5	1232	1467	2729	3294	3321	
GdiII	CGGCCRN!	1	2741					
GsuI	!CTCCAG	1	3626					
GsuI	!CTGGAG	3	872	1119	3647			
HaeI	WGG!CCW	9	776	1079	1100	1168	2650	
			2911	4273	4725	4736		
HaeII	RGC GC!Y	7	719	908	1190	2096	2104	
			2832	4507				
HaeIII	GG!CC	29	85	278	643	776	929	
			1079	1100	1168	1178	1240	
			1277	1475	1795	1937	2592	
			2598	2607	2650	2737	2911	
			3302	3329	3838	3895	3928	
			4273	4707	4725	4736		
HapII	C!CGG	20	645	694	854	1214	2047	
			2734	2811	2833	2861	2992	
			3082	3149	3330	3605	3613	
			3705	4179	4369	4395	4542	
HgaI	GACGCNNNNN!	4	527	618	3539	3618		
HgaI	!NNNNNNNNNNNGCGTC	3	2162	4057	4635			
HgiAI	GWGCW!C	5	880	1286	2942	3132	4437	
HgiEII	!ACCNNNNNNGGT	1	4159					
HhaI	GCG!C	29	665	667	718	907	986	
			1027	1189	1379	1381	2095	
			2103	2129	2151	2160	2173	
			2204	2823	2831	2895	2932	
			3198	3458	3634	3720	4123	
			4232	4406	4506	4573		
Hin1I	GR!CGYC	10	123	176	259	445	610	
			716	905	2829	3531	3610	
HinPII	G!CGC	29	663	665	716	905	984	
			1025	1187	1377	1379	2093	
			2101	2127	2149	2158	2171	
			2202	2821	2829	2893	2930	
			3196	3456	3632	3718	4121	
			4230	4404	4504	4571		
HincII	GTY!RAC	1	1592					
HindIII	A!AGCTT	1	1410					
HinfI	G!ANTC	14	411	1116	1431	1480	1872	
			1894	2304	3314	3448	3500	
			3558	3595	4377	4773		

HpaI	GTT!AAC	1	1592					
HpaII	C!CGG	20	645	694	854	1214	2047	
			2734	2811	2833	2861	2992	
			3082	3149	3330	3605	3613	
			3705	4179	4369	4395	4542	
HphI	GGTGANNNNNNNN!	5	377	897	1044	3900	4016	
HphI	!NNNNNNNTCACC	6	984	1125	1203	1336	1946	
			3007					
KasI	G!GCGCC	3	715	904	2828			
KpnI	GGTAC!C	0						
Ksp632I	CTCTTCN!	0						
Ksp632I	!NNNNGAAGAG	4	1177	2296	3173	3383		
KspI	CCGC!GG	1	795					
MaeI	C!TAG	8	593	1484	1678	2096	2652	
			3667	4001	4254			
MaeII	A!CGT	19	123	135	176	259	340	
			445	746	974	1103	1286	
			1316	1889	1901	1944	2054	
			2945	3132	3913	4046		
MaeIII	!GTNAC	15	62	149	498	885	1032	
			1335	1617	2118	2130	2949	
			3255	3945	4211	4327	4390	
MamI	GATNN!NNATC	2	1491	2688				
MboI	!GATC	27	586	671	687	737	1161	
			1196	1208	1242	1341	1444	
			1486	2666	2670	2689	3000	
			3078	3159	3168	3246	3622	
			3638	3997	4009	4087	4095	
			4106	4181				
MboII	GAAGANNNNNNNN!	6	1194	2085	2313	3190	3400	
			4019					
MboII	!NNNNNNNTCTTC	4	3480	3647	3847	4090		
McrI	!CGRYCG	3	694	2734	4409			
MfeI	!CAATTG	1	1578					
MflI	R!GATCY	13	586	671	687	737	1196	
			1444	3000	3246	3638	3997	
			4009	4095	4106			
MluI	A!CGCGT	0						
MmeI	!TCCRAC	1	1896					
MmeI	!GTYGGA	2	4372	4556				
MnlI	CCTCNNNNNNN!	10	1468	1548	1557	2612	2618	
			3430	3636	3950	4320	4644	
MnlI	!NNNNNNNGAGG	18	541	752	992	1022	1507	
			1972	2303	2581	2587	2624	
			2627	2639	2679	2743	2879	
			3236	3666	4569			
MroI	T!CCGGA	0						
MscI	TGG!CCA	4	776	1079	1100	2911		
MseI	T!TAA	16	8	1414	1530	1591	1712	
			1733	1744	1756	1767	1784	
			1882	2153	3972	3986	3991	
			4043					
MspI	C!CGG	20	645	694	854	1214	2047	
			2734	2811	2833	2861	2992	
			3082	3149	3330	3605	3613	
			3705	4179	4369	4395	4542	
MstI	TGC!GCA	1	2931					
MstII	CC!TNAGG	2	2309	3941				
MunI	C!AATTG	1	1579					

MvaI	CC!WGG	14	91 2365 3898	284 2420 4588	873 2437 4601	883 3216 4722	1081 3795
MvnI	CG!CG	22	61 1064 2129 3196 4123	663 1349 2149 3634 4704	665 1379 2173 3718	794 1478 2204 3781	1025 1753 2895 3850
NaeI	GCC!GGC	3	2048	3331	3614		
NarI	GG!CGCC	3	716	905	2829		
NciI	CC!SGG	5	645	2833	2993	3606	4370
NcoI	C!CATGG	3	361	2558	3261		
NdeI	CA!TATG	1	235				
NdeII	!GATC	27	586 1196 1486 3078 3638 4106	671 1208 2666 3159 3997 4181	687 1242 2670 3168 4009	737 1341 2689 3246 4087	1161 1444 3000 3622 4095
NgoMI	G!CCGGC	3	2046	3329	3612		
NheI	G!CTAGC	1	592				
NlaIII	CATG!	19	305 1308 2703 3647	365 2259 3048 4031	600 2399 3234 4751	708 2471 3265 4802	963 2562 3291
NlaIV	GGN!NCC	20	468 930 2208 3713	652 1278 2369 3792	689 1984 2441 3837	717 2005 2830 4680	906 2017 2865 4719
NotI	GC!GGCCGC	0					
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	2401	2473	4804		
Nsp(7524)I	RCATG!Y	6	600 4751	963	2399	2471	3234
Nsp(7524)V	TT!CGAA	1	3511				
NspBII	CMG!CKG	7	615 4164	713 4409	794	2327	2935
NspII	GDGCH!C	13	866 2020 3194	872 2775 3904	880 2868 4437	1286 2942	1292 3132
NspIII	C!YCGRG	2	931	1291			
NspIV	G!GNCC	11	84 1936 3927	277 3345	642 3790	928 3836	1276 3894
NunII	GG!CGCC	3	716	905	2829		
PacI	TTAAT!TAA	0					
Paer71	C!TCGAG	0					
PalI	GG!CC	29	85 1079 1277 2598 3302 4273	278 1100 1475 2607 3329 4707	643 1168 1795 2650 3838 4725	776 1178 1937 2737 3895 4736	929 1240 2592 2911 3928
PflMI	CCANNNN!NTGG	2	681	1107			
PleI	GAGTCNNNN!	2	1880	2312			
PleI	!NNNNNGACTC	6	405 4371	1425	1474	1888	3494
PmaCI	CAC!GTG	3	1104	1287	1317		
PmeI	GTTT!AAAC	0					

PmlI	CAC!GTG	3	1104	1287	1317			
PpuMI	RG!GWCCY	0						
PshAI	GACNN!NNGTC	0						
PstI	CTGCA!G	0						
PvuI	CGAT!CG	0						
PvuII	CAG!CTG	2	2327	2935				
RmaI	C!TAG	8	593	1484	1678	2096	2652	
			3667	4001	4254			
RsaI	GT!AC	11	220	245	300	333	384	
			541	925	1057	1150	1297	
			3135					
RsrI	G!AATTC	0						
RsrII	CG!GWCCG	1	3345					
SacI	GAGCT!C	0						
SacII	CCGC!GG	1	795					
SalI	G!TCGAC	0						
Sau3AI	!GATC	27	586	671	687	737	1161	
			1196	1208	1242	1341	1444	
			1486	2666	2670	2689	3000	
			3078	3159	3168	3246	3622	
			3638	3997	4009	4087	4095	
			4106	4181				
Sau96I	G!GNCC	11	84	277	642	928	1276	
			1936	3345	3790	3836	3894	
			3927					
SauI	CC!TNAGG	2	2309	3941				
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	19	91	284	645	873	883	
			1081	2365	2420	2437	2833	
			2993	3216	3606	3795	3898	
			4370	4588	4601	4722		
SduI	GDGCH!C	13	866	872	880	1286	1292	
			2020	2775	2868	2942	3132	
			3194	3904	4437			
SecI	C!CNNGG	22	361	729	792	871	881	
			1028	1080	1382	2363	2435	
			2558	2593	2602	2651	2992	
			3261	3666	3793	3794	3896	
			3897	4587				
SexI	!CTCGAG	0						
SfaNI	GCATCNNNNN!	7	797	1645	2408	2480	3128	
			3192	3467				
SfaNI	!NNNNNNNNNGATGC	7	358	801	822	2787	3042	
			3258	4650				
SfcI	C!TRYAG	3	2164	4291	4482			
SfiI	GGCCNNNN!NGGCC	1	2604					
SfuI	TT!CGAA	1	3511					
SgrAI	CR!CCGGYG	1	853					
SinI	G!GWCC	2	3345	3790				
SmaI	CCC!GGG	0						
SnaBI	TAC!GTA	1	341					
SnaI	!GTATAC	0						
SnoI	G!TGCAC	2	1282	4433				
SpeI	A!CTAGT	0						
SphI	GCATG!C	4	963	2399	2471	3234		
SplI	C!GTACG	0						
SrfI	GCCC!GGGC	0						
Sse8387I	CCTGCA!GG	0						
SspI	AAT!ATT	2	1737	2290				

SstI	GAGCT!C	0						
SstII	CCGC!GG	1	795					
SstIII	!ACGT	19	122	134	175	258	339	
			444	745	973	1102	1285	
			1315	1888	1900	1943	2053	
			2944	3131	3912	4045		
StuI	AGG!CCT	1	2650					
StyI	C!CWGG	5	361	2558	2651	3261	3666	
StySJI	!GAGNNNNNNGTRC	0						
StySJI	!GYACNNNNNNCTC	0						
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	13	998	1040	1978	2669	2942	
			3098	3122	3158	3320	3511	
			3556	3811	4649			
TaqII	GACCGANNNNNNNNNN!	4	1354	1849	2866	3534		
TaqII	!NNNNNNNNNTCGGTC	1	3778					
TaqII	CACCCANNNNNNNNNN!	1	1264					
TaqII	!NNNNNNNNNTGGGTG	3	173	2733	3325			
TfiI	G!AWTC	6	1116	3314	3448	3558	3595	
			4773					
ThaI	CG!CG	22	61	663	665	794	1025	
			1064	1349	1379	1478	1753	
			2129	2149	2173	2204	2895	
			3196	3634	3718	3781	3850	
			4123	4704				
Tsp45I	!GTSAC	6	885	1032	1335	2118	2949	
			3255					
TspEI	!AATT	13	19	1011	1128	1415	1579	
			1643	1721	1747	1758	2405	
			2477	2569	3987			
Tth111I	GACN!NNGTC	1	2947					
Tth111II	CAARCANNNNNNNNNN!	1	4156					
Tth111III	!NNNNNNNNNTGYTTG	3	3264	4124	4163			
TthHB8I	T!CGA	13	998	1040	1978	2669	2942	
			3098	3122	3158	3320	3511	
			3556	3811	4649			
Van91I	CCANNNN!NTGG	2	681	1107				
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	1483					
XcmI	!CCANNNNNNNNNTCG	1	3104					
XcyI	C!CCGGG	0						
XhoI	C!TCGAG	0						
XhoII	R!GATCY	13	586	671	687	737	1196	
			1444	3000	3246	3638	3997	
			4009	4095	4106			
XmaI	C!CCGGG	0						
XmaIII	C!GGCCG	1	2735					
XmnI	GAANN!NNTTC	1	943					
XorII	CGAT!CG	0						