

Enzyme Name	Sequence	Count	Cutting Position					
AatI	AGG!CCT	1	2561					
AatII	GACGT!C	4	126	179	262	448		
Acc65I	G!GTACC	1	1360					
AccI	GT!MKAC	1	1355					
AccII	CG!CG	21	61	702	933	972	1257	
			1287	1366	1628	1664	2040	
			2060	2084	2115	2806	3107	
			3545	3629	3692	3761	4034	
			4615					
AccIII	T!CCGGA	0						
AciI	G!CGG	28	371	404	529	621	702	
			1179	1182	1366	2000	2027	
			2115	2225	2806	2870	2971	
			2974	3254	3309	3407	3545	
			3692	3832	4075	4320	4441	
			4460	4587	4615			
AciI	C!CGC	36	59	87	99	113	280	
			508	590	700	931	1255	
			1300	1364	2055	2058	2072	
			2162	2410	2422	2431	2443	
			2453	2464	2510	2649	2712	
			3214	3259	3325	3351	3476	
			3479	3627	3759	4066	4210	
			4706					
AcyI	GR!CGYC	9	123	176	259	445	624	
			813	2740	3442	3521		
AfaI	GT!AC	12	220	245	300	333	384	
			541	833	965	1058	1205	
			1362	3046				
AflI	G!GWCC	2	3256	3701				
AflII	C!TTAAG	0						
AflIII	A!CRYGT	2	1626	4658				
AgeI	A!CCGGT	1	601					
AhaI	CC!SGG	6	1372	1373	2744	2904	3517	
			4281					
AhaII	GR!CGYC	9	123	176	259	445	624	
			813	2740	3442	3521		
AhaIII	TTT!AAA	3	1442	3884	3903			
AluI	AG!CT	21	567	844	864	979	1218	
			1231	1321	1327	1333	1339	
			1519	1686	1943	2238	2526	
			2846	3304	4101	4358	4404	
			4494					
Alw44I	G!TGCAC	2	1190	4344				
AlwI	GGATCNNNN!	9	1383	1392	2608	2919	3165	
			3557	3916	4014	4100		
AlwI	!NNNNNGATCC	7	581	640	1370	2984	3528	
			3915	4012				
AlwNI	CAGNNN!CTG	2	3841	4249				
AocI	CC!TNAGG	2	2220	3852				
AocII	GDGCH!C	15	774	780	788	1194	1200	
			1335	1372	1931	2686	2779	
			2853	3043	3105	3815	4348	
Aor51HI	AGC!GCT	1	597					
AosI	TGC!GCA	1	2842					
AosII	GR!CGYC	9	123	176	259	445	624	

			813	2740	3442	3521	
ApaI	GGGCC!C	1	1372				
ApaLI	G!TGCAC	2	1190	4344			
ApoI	R!AATTY	4	1344	1554	1658	1669	
ApyI	CC!WGG	14	91	284	781	791	989
			2276	2331	2348	3127	3706
			3809	4499	4512	4633	
AquI	C!YCGRG	4	839	1199	1328	1371	
AscI	GG!CGCGCC	0					
AseI	AT!TAAT	1	8				
AsnI	AT!TAAT	1	8				
Asp700I	GAANN!NNTTC	1	851				
Asp718I	G!GTACC	1	1360				
AspHI	GWGCW!C	6	788	1194	1335	2853	3043
			4348				
AspI	GACN!NNGTC	1	2858				
AsuI	G!GNCC	12	84	277	836	1184	1368
			1369	1847	3256	3701	3747
			3805	3838			
AsuII	TT!CGAA	2	1342	3422			
AvaI	C!YCGRG	4	839	1199	1328	1371	
AvaII	G!GWCC	2	3256	3701			
AvaIII	ATGCA!T	3	2312	2384	4715		
AviII	TGC!GCA	1	2842				
AvrI	!CYCGRG	4	838	1198	1327	1370	
AvrII	C!CTAGG	2	2562	3577			
AxyI	CC!TNAGG	2	2220	3852			
BalI	TGG!CCA	4	684	987	1008	2822	
BamHI	G!GATCC	1	1375				
BanI	G!GYRCC	7	466	623	812	1360	1893
			2739	2774			
BanII	GRGCY!C	5	1335	1372	1931	3105	3815
BanIII	AT!CGAT	1	2580				
BbeI	GGCGC!C	3	627	816	2743		
BbiII/AcyI	GR!CGYC	9	123	176	259	445	624
			813	2740	3442	3521	
BbrPI	CAC!GTG	3	1012	1195	1225		
BbsI	!GTCTTC	0					
BbsI	!GAAGAC	0					
BbvI	GCAGCNNNNNNNN!	9	630	969	1137	1528	2813
			2855	3376	3829	4329	
BbvI	!NNNNNNNNNNNGCTGC	8	851	2022	2687	2871	2964
			4030	4236	4239		
BbvII	GAAGACNN!	0					
BbvII	!NNNNNNGTCTTC	0					
BcefI	ACGGCNNNNNNNNNN!	7	655	730	826	850	1099
			3166	3653			
BcefI	!NNNNNNNNNNNGCCGT	4	1275	1882	2700	4159	
BcgI	!GCANNNNNTCG	0					
BcgI	!CGANNNNNTGC	0					
BclI	T!GATCA	1	1397				
BcnI	CC!SGG	6	1372	1373	2744	2904	3517
			4281				
BfaI	C!TAG	8	593	1388	1589	2007	2563
			3578	3912	4165		
BfrI	C!TTAAG	0					
BglI	GCCNNNN!NGGC	4	91	213	284	2515	
BglIII	A!GATCT	1	1104				
BinI	!NNNNNGATCC	7	581	640	1370	2984	3528

			3915	4012				
BinI	GGATCNNNN!	9	1383	1392	2608	2919	3165	
			3557	3916	4014	4100		
BlnI	C!CTAGG	2	2562	3577				
BmyI	GDGCH!C	15	774	780	788	1194	1200	
			1335	1372	1931	2686	2779	
			2853	3043	3105	3815	4348	
BpmI	!CTCCAG	1	3537					
BpmI	!CTGGAG	3	780	1027	3558			
Bpu1102I	GC!TNAGC	1	1232					
BsaAI	YAC!GTR	8	341	655	883	1012	1195	
			1225	1856	3044			
BsaBI	GATNN!NNATC	2	1402	2599				
BsaHI	GR!CGYC	9	123	176	259	445	624	
			813	2740	3442	3521		
BsaI	GGTCTCN!	0						
BsaI	!NNNNNGAGACC	2	1271	3729				
BsaJI	C!CNNGG	24	361	637	700	779	789	
			936	988	1290	1364	1371	
			2274	2346	2469	2504	2513	
			2562	2903	3172	3577	3704	
			3705	3807	3808	4498		
BseAI	T!CCGGA	0						
BsePI	!GCGCGC	1	1284					
BsgI	!GTGCAG	1	883					
BsgI	!CTGCAC	1	753					
BsiEI	CGRY!CG	3	606	2649	4324			
BsiWI	C!GTACG	0						
BsiYI	CCNNNNN!NNGG	24	50	655	706	780	937	
			1015	1178	1752	2078	2470	
			2721	3265	3546	3578	3579	
			3710	3745	3795	3796	3827	
			4180	4459	4625	4643		
BslI	CCNNNNN!NNGG	24	50	655	706	780	937	
			1015	1178	1752	2078	2470	
			2721	3265	3546	3578	3579	
			3710	3745	3795	3796	3827	
			4180	4459	4625	4643		
BsmAI	!GTCTC	1	429					
BsmAI	!GAGAC	5	1276	2169	2587	3606	3734	
BsmI	GAATGCN!	1	1491					
BsmI	G!CATTC	1	1584					
Bsp1286I	GDGCH!C	15	774	780	788	1194	1200	
			1335	1372	1931	2686	2779	
			2853	3043	3105	3815	4348	
BspDI	AT!CGAT	1	2580					
BspEI	T!CCGGA	0						
BspHI	T!CATGA	2	2166	3938				
BspMI	ACCTGCNNNN!	2	3008	3458				
BspMI	!NNNNNNNNGCAGGT	1	2627					
BspMII	T!CCGGA	0						
BspWI	GCNNNNN!NNGC	34	91	213	245	284	377	
			401	708	723	762	819	
			1092	1131	1967	1997	2024	
			2066	2068	2307	2379	2430	
			2509	2515	2731	2815	2838	
			2977	2983	3100	3183	3450	
			3637	3841	4040	4612		
BsrFI	R!CCGGY	7	601	761	1121	1957	3059	

			3240	3523			
BsrI	ACTGGN!	3	2684	2885	4131		
BsrI	C!CAGT	7	296	745	817	1769	2447
			4243	4256			
BssHII	G!CGCGC	1	1285				
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	2	1342	3422			
BstEII	G!GTNACC	2	793	940			
BstI	G!GATCC	1	1375				
BstNI	CC!WGG	14	91	284	781	791	989
			2276	2331	2348	3127	3706
			3809	4499	4512	4633	
BstPI	G!GTNACC	2	793	940			
BstUI	CG!CG	21	61	702	933	972	1257
			1287	1366	1628	1664	2040
			2060	2084	2115	2806	3107
			3545	3629	3692	3761	4034
			4615				
BstXI	CCANNNNN!NTGG	0					
BstYI	R!GATCY	12	586	645	1104	1375	1384
			2911	3157	3549	3908	3920
			4006	4017			
Bsu36I	CC!TNAGG	2	2220	3852			
CbiI	TT!CGAA	2	1342	3422			
CcrI	C!TCGAG	1	1328				
CelII	GC!TNAGC	1	1232				
CfoI	GCG!C	28	598	626	815	894	935
			1097	1287	1289	2006	2014
			2040	2062	2071	2084	2115
			2734	2742	2806	2843	3109
			3369	3545	3631	4034	4143
			4317	4417	4484		
Cfr10I	R!CCGGY	7	601	761	1121	1957	3059
			3240	3523			
Cfr13I	G!GNCC	12	84	277	836	1184	1368
			1369	1847	3256	3701	3747
			3805	3838			
Cfr9I	C!CCGGG	1	1371				
CfrI	Y!GGCCR	8	682	985	1006	1146	2646
			2820	3211	3238		
ClaI	AT!CGAT	1	2580				
CpoI	CG!GWCCG	1	3256				
Csp45I	TT!CGAA	2	1342	3422			
CspI	CG!GSCCG	0					
CviJI	RG!CY	93	43	85	94	278	567
			684	711	837	844	864
			879	979	987	993	1008
			1076	1086	1148	1185	1218
			1231	1269	1321	1327	1333
			1339	1370	1409	1519	1686
			1706	1848	1929	1943	1957
			2238	2279	2351	2474	2503
			2509	2518	2526	2549	2561
			2567	2648	2665	2673	2700
			2725	2809	2818	2822	2846
			2884	2960	2977	2996	3059
			3103	3113	3130	3213	3240
			3244	3281	3304	3320	3500
			3527	3588	3710	3749	3806

			3813	3820	3839	3850	4101
			4130	4173	4184	4249	4328
			4353	4358	4404	4494	4592
			4618	4636	4647		
CvnI	CC!TNAGG	2	2220	3852			
DdeI	C!TNAG	7	1232	2220	2522	3403	3852
			3975	4384			
DpnI	GA!TC	26	588	647	1071	1106	1118
			1152	1251	1377	1386	1399
			2579	2583	2602	2913	2991
			3072	3081	3159	3535	3551
			3910	3922	4000	4008	4019
			4094				
DpnII	!GATC	26	586	645	1069	1104	1116
			1150	1249	1375	1384	1397
			2577	2581	2600	2911	2989
			3070	3079	3157	3533	3549
			3908	3920	3998	4006	4017
			4092				
DraI	TTT!AAA	3	1442	3884	3903		
DraII	RG!GNCCY	1	3838				
DraIII	CACNNN!GTG	1	1856				
DrdI	GACNNNN!NNGTC	4	1312	1811	2767	4556	
DsaI	C!CRYGG	7	361	637	700	1290	1364
			2469	3172			
DsaV	!CCNGG	20	89	282	779	789	987
			1370	1371	2274	2329	2346
			2742	2902	3125	3515	3704
			3807	4279	4497	4510	4631
EaeI	Y!GGCCR	8	682	985	1006	1146	2646
			2820	3211	3238		
EagI	C!GGCCG	1	2646				
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	4	1089	2211	3088	3298	
Ecl136II	GAG!CTC	1	1333				
EclXI	C!GGCCG	1	2646				
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	2	1271	3729			
Eco47I	G!GWCC	2	3256	3701			
Eco47III	AGC!GCT	1	597				
Eco52I	C!GGCCG	1	2646				
Eco57I	!CTGAAG	4	844	1087	2864	3296	
Eco57I	!CTTCAG	2	690	4130			
Eco81I	CC!TNAGG	2	2220	3852			
EcoNI	CCTNN!NNNAGG	0					
EcoO109I	RG!GNCCY	1	3838				
EcoO65I	G!GTNACC	2	793	940			
EcoRI	G!AATTC	1	1344				
EcoRII	!CCWGG	14	89	282	779	789	987
			2274	2329	2346	3125	3704
			3807	4497	4510	4631	
EcoRV	GAT!ATC	0					
EcoT14I	C!CWGGG	5	361	2469	2562	3172	3577
EcoT22I	ATGCA!T	3	2312	2384	4715		
EcoT38I	GRGCY!C	5	1335	1372	1931	3105	3815
EheI	GGC!GCC	3	625	814	2741		
Esp3I	CGTCTCN!	0					

Esp3I	!NNNNNGAGACG	0						
EspI	GC!TNAGC	1	1232					
FbaI	T!GATCA	1	1397					
FinI	!GTCCC	4	2270	2342	2406	3701		
FinI	!GGGAC	6	161	312	480	2876	3408	
			3517					
Fnu4HI	GC!NGC	36	619	622	865	958	1126	
			1180	1183	1300	1517	2036	
			2058	2072	2510	2649	2701	
			2712	2802	2807	2844	2885	
			2972	2975	2978	3214	3310	
			3351	3365	3479	3818	3833	
			4044	4250	4253	4318	4461	
			4616					
FokI	GGATGNNNNNNNNN!	8	692	752	1016	1253	2607	
			3064	3089	3543			
FokI	!NNNNNNNNNNNNNNCATCC	2	683	2413				
FspI	TGC!GCA	1	2842					
GdiII	!NNNNNYGGCCG	4	1140	2640	3205	3232		
GdiII	CGGCCRN!	1	2652					
GsuI	!CTCCAG	1	3537					
GsuI	!CTGGAG	3	780	1027	3558			
HaeI	WGG!CCW	9	684	987	1008	1076	2561	
			2822	4184	4636	4647		
HaeII	RGCGC!Y	8	599	627	816	1098	2007	
			2015	2743	4418			
HaeIII	GG!CC	28	85	278	684	837	987	
			1008	1076	1086	1148	1185	
			1370	1706	1848	2503	2509	
			2518	2561	2648	2822	3213	
			3240	3749	3806	3839	4184	
			4618	4636	4647			
HapII	C!CGG	21	602	762	1122	1372	1382	
			1958	2645	2722	2744	2772	
			2903	2993	3060	3241	3516	
			3524	3616	4090	4280	4306	
			4453					
HgaI	GACGCNNNNN!	3	527	3450	3529			
HgaI	!NNNNNNNNNNNGCGTC	3	2073	3968	4546			
HgiAI	GWGCW!C	6	788	1194	1335	2853	3043	
			4348					
HgiEII	!ACCNNNNNNGGT	1	4070					
HhaI	GCG!C	28	598	626	815	894	935	
			1097	1287	1289	2006	2014	
			2040	2062	2071	2084	2115	
			2734	2742	2806	2843	3109	
			3369	3545	3631	4034	4143	
			4317	4417	4484			
Hin1I	GR!CGYC	9	123	176	259	445	624	
			813	2740	3442	3521		
HinP1I	G!CGC	28	596	624	813	892	933	
			1095	1285	1287	2004	2012	
			2038	2060	2069	2082	2113	
			2732	2740	2804	2841	3107	
			3367	3543	3629	4032	4141	
			4315	4415	4482			
HincII	GTY!RAC	2	1356	1503				
HindIII	A!AGCTT	1	1337					
HinfI	G!ANTC	12	411	1024	1783	1805	2215	

			3225	3359	3411	3469	3506
			4288	4684			
HpaI	GTT!AAC	1	1503				
HpaII	C!CGG	21	602	762	1122	1372	1382
			1958	2645	2722	2744	2772
			2903	2993	3060	3241	3516
			3524	3616	4090	4280	4306
			4453				
HphI	GGTGANNNNNNNN!	5	377	805	952	3811	3927
HphI	!NNNNNNNTCACC	5	892	1033	1244	1857	2918
KasI	G!GCGCC	3	623	812	2739		
KpnI	GGTAC!C	1	1364				
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	4	1085	2207	3084	3294	
KspI	CCGC!GG	2	703	1367			
MaeI	C!TAG	8	593	1388	1589	2007	2563
			3578	3912	4165		
MaeII	A!CGT	19	123	135	176	259	340
			445	654	882	1011	1194
			1224	1800	1812	1855	1965
			2856	3043	3824	3957	
MaeIII	!GTNAC	15	62	149	498	793	940
			1243	1528	2029	2041	2860
			3166	3856	4122	4238	4301
MamI	GATNN!NNATC	2	1402	2599			
MboI	!GATC	26	586	645	1069	1104	1116
			1150	1249	1375	1384	1397
			2577	2581	2600	2911	2989
			3070	3079	3157	3533	3549
			3908	3920	3998	4006	4017
			4092				
MboII	GAAGANNNNNNNN!	6	1102	1996	2224	3101	3311
			3930				
MboII	!NNNNNNNTCTTC	4	3391	3558	3758	4001	
McrI	!CGRYCG	3	602	2645	4320		
MfeI	!CAATTG	1	1489				
MflI	R!GATCY	12	586	645	1104	1375	1384
			2911	3157	3549	3908	3920
			4006	4017			
MluI	A!CGCGT	1	1626				
MmeI	!TCCRAC	1	1807				
MmeI	!GTYGGA	2	4283	4467			
MnlI	CCTCNNNNNNN!	9	1459	1468	2523	2529	3341
			3547	3861	4231	4555	
MnlI	!NNNNNNNGAGG	18	541	660	900	930	1418
			1883	2214	2492	2498	2535
			2538	2550	2590	2654	2790
			3147	3577	4480		
MroI	T!CCGGA	0					
MscI	TGG!CCA	4	684	987	1008	2822	
MseI	T!TAA	15	8	1441	1502	1623	1644
			1655	1667	1678	1695	1793
			2064	3883	3897	3902	3954
MspI	C!CGG	21	602	762	1122	1372	1382
			1958	2645	2722	2744	2772
			2903	2993	3060	3241	3516
			3524	3616	4090	4280	4306
			4453				
MstI	TGC!GCA	1	2842				

MstII	CC!TNAGG	2	2220	3852				
MunI	C!AATTG	1	1490					
MvaI	CC!WGG	14	91	284	781	791	989	
			2276	2331	2348	3127	3706	
			3809	4499	4512	4633		
MvnI	CG!CG	21	61	702	933	972	1257	
			1287	1366	1628	1664	2040	
			2060	2084	2115	2806	3107	
			3545	3629	3692	3761	4034	
			4615					
NaeI	GCC!GGC	4	1123	1959	3242	3525		
NarI	GG!CGCC	3	624	813	2740			
NciI	CC!SGG	6	1372	1373	2744	2904	3517	
			4281					
NcoI	C!CATGG	3	361	2469	3172			
NdeI	CA!TATG	1	235					
NdeII	!GATC	26	586	645	1069	1104	1116	
			1150	1249	1375	1384	1397	
			2577	2581	2600	2911	2989	
			3070	3079	3157	3533	3549	
			3908	3920	3998	4006	4017	
			4092					
NgoMI	G!CCGGC	4	1121	1957	3240	3523		
NheI	G!CTAGC	1	592					
NlaIII	CATG!	18	305	365	616	871	1216	
			2170	2310	2382	2473	2614	
			2959	3145	3176	3202	3558	
			3942	4662	4713			
NlaIV	GGN!NCC	21	468	625	814	838	1186	
			1362	1370	1377	1895	1916	
			1928	2119	2280	2352	2741	
			2776	3624	3703	3748	4591	
			4630					
NotI	GC!GGCCGC	0						
NruI	TCG!CGA	0						
NsiI	ATGCA!T	3	2312	2384	4715			
Nsp(7524)I	RCATG!Y	5	871	2310	2382	3145	4662	
Nsp(7524)V	TT!CGAA	2	1342	3422				
NspBII	CMG!CKG	7	621	702	1366	2238	2846	
			4075	4320				
NspII	GDGCH!C	15	774	780	788	1194	1200	
			1335	1372	1931	2686	2779	
			2853	3043	3105	3815	4348	
NspIII	C!YCGRG	4	839	1199	1328	1371		
NspIV	G!GNCC	12	84	277	836	1184	1368	
			1369	1847	3256	3701	3747	
			3805	3838				
NunII	GG!CGCC	3	624	813	2740			
PacI	TTAAT!TAA	0						
Paer71	C!TCGAG	1	1328					
PalI	GG!CC	28	85	278	684	837	987	
			1008	1076	1086	1148	1185	
			1370	1706	1848	2503	2509	
			2518	2561	2648	2822	3213	
			3240	3749	3806	3839	4184	
			4618	4636	4647			
PflMI	CCANNNN!NTGG	1	1015					
PleI	GAGTCNNNN!	2	1791	2223				
PleI	!NNNNNGACTC	4	405	1799	3405	4282		

PmaCI	CAC!GTG	3	1012	1195	1225		
PmeI	GTTT!AAAC	0					
PmlI	CAC!GTG	3	1012	1195	1225		
PpuMI	RG!GWCCY	0					
PshAI	GACNN!NNGTC	0					
PstI	CTGCA!G	1	1353				
PvuI	CGAT!CG	0					
PvuII	CAG!CTG	2	2238	2846			
RmaI	C!TAG	8	593	1388	1589	2007	2563
			3578	3912	4165		
RsaI	GT!AC	12	220	245	300	333	384
			541	833	965	1058	1205
			1362	3046			
RsrI	G!AATTC	1	1344				
RsrII	CG!GWCCG	1	3256				
SacI	GAGCT!C	1	1335				
SacII	CCGC!GG	2	703	1367			
SalI	G!TCGAC	1	1354				
Sau3AI	!GATC	26	586	645	1069	1104	1116
			1150	1249	1375	1384	1397
			2577	2581	2600	2911	2989
			3070	3079	3157	3533	3549
			3908	3920	3998	4006	4017
			4092				
Sau96I	G!GNCC	12	84	277	836	1184	1368
			1369	1847	3256	3701	3747
			3805	3838			
SauI	CC!TNAGG	2	2220	3852			
ScaI	AGT!ACT	0					
ScrFI	CC!NGG	20	91	284	781	791	989
			1372	1373	2276	2331	2348
			2744	2904	3127	3517	3706
			3809	4281	4499	4512	4633
SduI	GDGCH!C	15	774	780	788	1194	1200
			1335	1372	1931	2686	2779
			2853	3043	3105	3815	4348
SecI	C!CNNGG	24	361	637	700	779	789
			936	988	1290	1364	1371
			2274	2346	2469	2504	2513
			2562	2903	3172	3577	3704
			3705	3807	3808	4498	
SexI	!CTCGAG	1	1327				
SfaNI	GCATCNNNNN!	7	705	1556	2319	2391	3039
			3103	3378			
SfaNI	!NNNNNNNNGATGC	7	358	709	730	2698	2953
			3169	4561			
SfcI	C!TRYAG	5	1322	1349	2075	4202	4393
SfiI	GGCCNNNN!NGGCC	1	2515				
SfuI	TT!CGAA	2	1342	3422			
SgrAI	CR!CCGGYG	1	761				
SinI	G!GWCC	2	3256	3701			
SmaI	CCC!GGG	1	1373				
SnaBI	TAC!GTA	1	341				
SnaI	!GTATAC	0					
SnoI	G!TGCAC	2	1190	4344			
SpeI	A!CTAGT	0					
SphI	GCATG!C	4	871	2310	2382	3145	
SplI	C!GTACG	0					
SrfI	GCCC!GGGC	0					

Sse8387I	CCTGCA!GG	0						
SspI	AAT!ATT	2	1648	2201				
SstI	GAGCT!C	1	1335					
SstII	CCGC!GG	2	703	1367				
SstIII	!ACGT	19	122	134	175	258	339	
			444	653	881	1010	1193	
			1223	1799	1811	1854	1964	
			2855	3042	3823	3956		
StuI	AGG!CCT	1	2561					
StyI	C!CWGG	5	361	2469	2562	3172	3577	
StySJI	!GAGNNNNNNGTRC	0						
StySJI	!GYACNNNNNNCTC	0						
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	16	906	948	1329	1342	1355	
			1889	2580	2853	3009	3033	
			3069	3231	3422	3467	3722	
			4560					
TaqII	GACCGANNNNNNNNNNN!	4	1262	1760	2777	3445		
TaqII	!NNNNNNNNNTCGGTC	1	3689					
TaqII	CACCCANNNNNNNNNNN!	1	1172					
TaqII	!NNNNNNNNNTGGGTG	3	173	2644	3236			
TfiI	G!AWTC	6	1024	3225	3359	3469	3506	
			4684					
ThaI	CG!CG	21	61	702	933	972	1257	
			1287	1366	1628	1664	2040	
			2060	2084	2115	2806	3107	
			3545	3629	3692	3761	4034	
			4615					
Tsp45I	!GTSAC	6	793	940	1243	2029	2860	
			3166					
TspEI	!AATT	13	19	919	1036	1344	1490	
			1554	1632	1658	1669	2316	
			2388	2480	3898			
Tth111I	GACN!NNGTC	1	2858					
Tth111II	CAARCANNNNNNNNNNN!	1	4067					
Tth111III	!NNNNNNNNNTGYTTG	3	3175	4035	4074			
TthHB8I	T!CGA	16	906	948	1329	1342	1355	
			1889	2580	2853	3009	3033	
			3069	3231	3422	3467	3722	
			4560					
Van91I	CCANNNN!NTGG	1	1015					
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	1387					
XcmI	!CCANNNNNNNNNNTCG	2	1316	3015				
XcyI	C!CCGGG	1	1371					
XhoI	C!TCGAG	1	1328					
XhoII	R!GATCY	12	586	645	1104	1375	1384	
			2911	3157	3549	3908	3920	
			4006	4017				
XmaI	C!CCGGG	1	1371					
XmaIII	C!GGCCG	1	2646					
XmnI	GAANN!NNTTC	1	851					
XorII	CGAT!CG	0						