

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	1	2574				
AatII	GACGT!C	4	126	179	262	448	
Acc65I	G!GTACC	1	1373				
AccI	GT!MKAC	1	1368				
AccII	CG!CG	19	61	937	939	1257	1379
			1641	1677	2053	2073	2097
			2128	2819	3120	3558	3642
			3705	3774	4047	4628	
AccIII	T!CCGGA	1	1328				
AciI	G!CGG	25	371	404	529	702	1379
			2013	2040	2128	2238	2819
			2883	2984	2987	3267	3322
			3420	3558	3705	3845	4088
			4333	4454	4473	4600	4628
AciI	C!CGC	36	59	87	99	113	280
			508	590	1114	1287	1290
			1293	1377	2068	2071	2085
			2175	2423	2435	2444	2456
			2466	2477	2523	2662	2725
			3227	3272	3338	3364	3489
			3492	3640	3772	4079	4223
			4719				
AcyI	GR!CGYC	8	123	176	259	445	624
			2753	3455	3534		
AfaI	GT!AC	11	220	245	300	333	384
			541	1040	1064	1322	1375
			3059				
AflI	G!GWCC	4	1111	1240	3269	3714	
AflIII	C!TTAAG	0					
AflIII	A!CRYGT	2	1639	4671			
AgeI	A!CCGGT	1	601				
AhaI	CC!SGG	6	1385	1386	2757	2917	3530
			4294				
AhaII	GR!CGYC	8	123	176	259	445	624
			2753	3455	3534		
AhaIII	TTT!AAA	3	1455	3897	3916		
AluI	AG!CT	22	567	631	664	736	769
			807	985	1144	1318	1346
			1352	1532	1699	1956	2251
			2539	2859	3317	4114	4371
			4417	4507			
Alw44I	G!TGCAC	1	4357				
AlwI	GGATCNNNN!	9	1396	1405	2621	2932	3178
			3570	3929	4027	4113	
AlwI	!NNNNNGATCC	6	581	1383	2997	3541	3928
			4025				
AlwNI	CAGNNN!CTG	2	3854	4262			
AocI	CC!TNAGG	3	879	2233	3865		
AocII	GDGCH!C	12	651	780	1348	1385	1944
			2699	2792	2866	3056	3118
			3828	4361			
Aor51HI	AGC!GCT	1	597				
AosI	TGC!GCA	1	2855				
AosII	GR!CGYC	8	123	176	259	445	624
			2753	3455	3534		
ApaI	GGGCC!C	1	1385				

ApaI	G!TGCAC	1	4357				
ApoI	R!AATTY	4	1357	1567	1671	1682	
ApyI	CC!WGG	17	91	284	781	791	893
			968	1022	1084	2289	2344
			2361	3140	3719	3822	4512
			4525	4646			
AquI	C!YCGRG	3	1274	1341	1384		
AscI	GG!CGCGCC	0					
AseI	AT!TAAT	1	8				
AsnI	AT!TAAT	1	8				
Asp700I	GAANN!NNTTC	0					
Asp718I	G!GTACC	1	1373				
AspHI	GWGCW!C	4	1348	2866	3056	4361	
AspI	GACN!NNGTC	1	2871				
AsuI	G!GNCC	14	84	277	783	1111	1184
			1240	1381	1382	1860	3269
			3714	3760	3818	3851	
AsuII	TT!CGAA	2	1355	3435			
AvaI	C!YCGRG	3	1274	1341	1384		
AvaII	G!GWCC	4	1111	1240	3269	3714	
AvaIII	ATGCA!T	3	2325	2397	4728		
AviII	TGC!GCA	1	2855				
AvrI	!CYCGRG	3	1273	1340	1383		
AvrII	C!CTAGG	2	2575	3590			
AxyI	CC!TNAGG	3	879	2233	3865		
BalI	TGG!CCA	3	684	797	2835		
BamHI	G!GATCC	1	1388				
BanI	G!GYRCC	6	466	623	1373	1906	2752
			2787				
BanII	GRGCY!C	5	1348	1385	1944	3118	3828
BanIII	AT!CGAT	1	2593				
BbeI	GGCGC!C	2	627	2756			
BbiIII/AcyI	GR!CGYC	8	123	176	259	445	624
			2753	3455	3534		
BbrPI	CAC!GTG	0					
BbsI	!GTCTTC	0					
BbsI	!GAAGAC	0					
BbvI	GCAGCNNNNNNNN!	9	862	1146	1153	1541	2826
			2868	3389	3842	4342	
BbvI	!NNNNNNNNNNNGCTGC	10	756	1179	1182	2035	2700
			2884	2977	4043	4249	4252
BbvII	GAAGACNN!	0					
BbvII	!NNNNNNGTCTTC	0					
BcefI	ACGGCNNNNNNNNNN!	6	745	826	937	1321	3179
			3666				
BcefI	!NNNNNNNNNNNGCCGT	3	1895	2713	4172		
BcgI	!GCANNNNNNTCG	0					
BcgI	!CGANNNNNNTGC	1	711				
BclI	T!GATCA	1	1410				
BcnI	CC!SGG	6	1385	1386	2757	2917	3530
			4294				
BfaI	C!TAG	8	593	1401	1602	2020	2576
			3591	3925	4178		
BfrI	C!TTAAG	0					
BglI	GCCNNN!NGGC	5	91	213	284	1079	2528
BglII	A!GATCT	1	1337				
BinI	!NNNNNGATCC	6	581	1383	2997	3541	3928
			4025				
BinI	GGATCNNNN!	9	1396	1405	2621	2932	3178

			3570	3929	4027	4113	
BlnI	C!CTAGG	2	2575	3590			
BmyI	GDGCH!C	12	651	780	1348	1385	1944
			2699	2792	2866	3056	3118
			3828	4361			
BpmI	!CTCCAG	1	3550				
BpmI	!CTGGAG	1	3571				
Bpu1102I	GC!TNAGC	0					
BsaAI	YAC!GTR	3	341	1869	3057		
BsaBI	GATNN!NNATC	2	1415	2612			
BsaHI	GR!CGYC	8	123	176	259	445	624
			2753	3455	3534		
BsaI	GGTCTCN!	0					
BsaI	!NNNNNGAGACC	1	3742				
BsaJI	C!CNNGG	24	361	779	789	942	966
			1021	1236	1303	1377	1384
			2287	2359	2482	2517	2526
			2575	2916	3185	3590	3717
			3718	3820	3821	4511	
BseAI	T!CCGGA	1	1328				
BsePI	!GCGCGC	1	936				
BsgI	!GTGCAG	2	814	1138			
BsgI	!CTGCAC	1	753				
BsiEI	CGRY!CG	3	606	2662	4337		
BsiWI	C!GTACG	0					
BsiYI	CCNNNNN!NNGG	20	50	1237	1309	1765	2091
			2483	2734	3278	3559	3591
			3592	3723	3758	3808	3809
			3840	4193	4472	4638	4656
BslI	CCNNNNN!NNGG	20	50	1237	1309	1765	2091
			2483	2734	3278	3559	3591
			3592	3723	3758	3808	3809
			3840	4193	4472	4638	4656
BsmAI	!GTCTC	1	429				
BsmAI	!GAGAC	4	2182	2600	3619	3747	
BsmI	GAATGCN!	1	1504				
BsmI	G!CATTC	1	1597				
Bsp1286I	GDGCH!C	12	651	780	1348	1385	1944
			2699	2792	2866	3056	3118
			3828	4361			
BspDI	AT!CGAT	1	2593				
BspEI	T!CCGGA	1	1328				
BspHI	T!CATGA	2	2179	3951			
BspMI	ACCTGCNNNN!	2	3021	3471			
BspMI	!NNNNNNNNGCAGGT	1	2640				
BspMII	T!CCGGA	1	1328				
BspWI	GCNNNNN!NNGC	35	91	213	245	284	377
			401	702	708	762	775
			819	828	1079	1980	2010
			2037	2079	2081	2320	2392
			2443	2522	2528	2744	2828
			2851	2990	2996	3113	3196
			3463	3650	3854	4053	4625
BsrFI	R!CCGGY	8	601	638	761	989	1970
			3072	3253	3536		
BsrI	ACTGGN!	3	2697	2898	4144		
BsrI	C!CAGT	5	296	1782	2460	4256	4269
BssHII	G!CGCGC	1	937				
Bst1107I	GTA!TAC	0					

BstBI	TT!CGAA	2	1355	3435				
BstEII	G!GTNACC	0						
BstI	G!GATCC	1	1388					
BstNI	CC!WGG	17	91	284	781	791	893	
			968	1022	1084	2289	2344	
			2361	3140	3719	3822	4512	
			4525	4646				
BstPI	G!GTNACC	0						
BstUI	CG!CG	19	61	937	939	1257	1379	
			1641	1677	2053	2073	2097	
			2128	2819	3120	3558	3642	
			3705	3774	4047	4628		
BstXI	CCANNNNN!NTGG	1	793					
BstYI	R!GATCY	11	586	1337	1388	1397	2924	
			3170	3562	3921	3933	4019	
			4030					
Bsu36I	CC!TNAGG	3	879	2233	3865			
CbiI	TT!CGAA	2	1355	3435				
CcrI	C!TCGAG	2	1274	1341				
CelII	GC!TNAGC	0						
CfoI	GCG!C	28	598	626	872	900	939	
			941	1229	1257	2019	2027	
			2053	2075	2084	2097	2128	
			2747	2755	2819	2856	3122	
			3382	3558	3644	4047	4156	
			4330	4430	4497			
Cfr10I	R!CCGGY	8	601	638	761	989	1970	
			3072	3253	3536			
Cfr13I	G!GNCC	14	84	277	783	1111	1184	
			1240	1381	1382	1860	3269	
			3714	3760	3818	3851		
Cfr9I	C!CCGGG	1	1384					
CfrI	Y!GGCCR	8	682	795	1071	1146	2659	
			2833	3224	3251			
ClaI	AT!CGAT	1	2593					
CpoI	CG!GWCCG	1	3269					
Csp45I	TT!CGAA	2	1355	3435				
CspI	CG!GSCCG	0						
CviJI	RG!CY	90	43	85	94	278	567	
			631	664	684	736	769	
			784	797	807	885	985	
			1073	1082	1144	1148	1185	
			1318	1346	1352	1383	1422	
			1532	1699	1719	1861	1942	
			1956	1970	2251	2292	2364	
			2487	2516	2522	2531	2539	
			2562	2574	2580	2661	2678	
			2686	2713	2738	2822	2831	
			2835	2859	2897	2973	2990	
			3009	3072	3116	3126	3143	
			3226	3253	3257	3294	3317	
			3333	3513	3540	3601	3723	
			3762	3819	3826	3833	3852	
			3863	4114	4143	4186	4197	
			4262	4341	4366	4371	4417	
			4507	4605	4631	4649	4660	
CvnI	CC!TNAGG	3	879	2233	3865			
DdeI	C!TNAG	9	803	879	1334	2233	2535	
			3416	3865	3988	4397		

DpnI	GA!TC	25	588	659	840	1260	1268
			1339	1390	1399	1412	2592
			2596	2615	2926	3004	3085
			3094	3172	3548	3564	3923
			3935	4013	4021	4032	4107
DpnII	!GATC	25	586	657	838	1258	1266
			1337	1388	1397	1410	2590
			2594	2613	2924	3002	3083
			3092	3170	3546	3562	3921
			3933	4011	4019	4030	4105
DraI	TTT!AAA	3	1455	3897	3916		
DraII	RG!GNCCY	2	1240	3851			
DraIII	CACNNN!GTG	1	1869				
DrdI	GACNNNN!NNGTC	3	1824	2780	4569		
DsaI	C!CRYGG	5	361	1303	1377	2482	3185
DsaV	!CCNGG	23	89	282	779	789	891
			966	1020	1082	1383	1384
			2287	2342	2359	2755	2915
			3138	3528	3717	3820	4292
			4510	4523	4644		
EaeI	Y!GGCCR	8	682	795	1071	1146	2659
			2833	3224	3251		
EagI	C!GGCCG	1	2659				
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	3	2224	3101	3311		
Ecl136II	GAG!CTC	1	1346				
EclXI	C!GGCCG	1	2659				
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	1	3742				
Eco47I	G!GWCC	4	1111	1240	3269	3714	
Eco47III	AGC!GCT	1	597				
Eco52I	C!GGCCG	1	2659				
Eco57I	!CTGAAG	3	742	2877	3309		
Eco57I	!CTTCAG	1	4143				
Eco81I	CC!TNAGG	3	879	2233	3865		
EcoNI	CCTNN!NNNAGG	1	1235				
EcoO109I	RG!GNCCY	2	1240	3851			
EcoO65I	G!GTNACC	0					
EcoRI	G!AATTC	1	1357				
EcoRII	!CCWGG	17	89	282	779	789	891
			966	1020	1082	2287	2342
			2359	3138	3717	3820	4510
			4523	4644			
EcoRV	GAT!ATC	0					
EcoT14I	C!CWGGG	6	361	1236	2482	2575	3185
			3590				
EcoT22I	ATGCA!T	3	2325	2397	4728		
EcoT38I	GRGCY!C	5	1348	1385	1944	3118	3828
EheI	GGC!GCC	2	625	2754			
Esp3I	CGTCTCN!	0					
Esp3I	!NNNNNGAGACG	0					
EspI	GC!TNAGC	0					
FbaI	T!GATCA	1	1410				
FinI	!GTCCC	4	2283	2355	2419	3714	
FinI	!GGGAC	6	161	312	480	2889	3421
			3530				
Fnu4HI	GC!NGC	37	703	770	851	1135	1142

			1193	1196	1290	1293	1530
			2049	2071	2085	2523	2662
			2714	2725	2815	2820	2857
			2898	2985	2988	2991	3227
			3323	3364	3378	3492	3831
			3846	4057	4263	4266	4331
			4474	4629			
FokI	GGATGNNNNNNNNNN!	8	929	1022	1142	1325	2620
			3077	3102	3556		
FokI	!NNNNNNNNNNNNNCATCC	4	638	875	1004	2426	
FspI	TGC!GCA	1	2855				
GdiIII	!NNNNNYGGCCG	5	1065	1140	2653	3218	3245
GdiIII	CGGCCRN!	1	2665				
GsuI	!CTCCAG	1	3550				
GsuI	!CTGGAG	1	3571				
HaeI	WGG!CCW	8	684	797	1082	2574	2835
			4197	4649	4660		
HaeII	RGCGC!Y	8	599	627	873	1230	2020
			2028	2756	4431		
HaeIII	GG!CC	27	85	278	684	784	797
			1073	1082	1148	1185	1383
			1719	1861	2516	2522	2531
			2574	2661	2835	3226	3253
			3762	3819	3852	4197	4631
			4649	4660			
HapII	C!CGG	23	602	639	762	990	1329
			1385	1395	1971	2658	2735
			2757	2785	2916	3006	3073
			3254	3529	3537	3629	4103
			4293	4319	4466		
HgaI	GACGCNNNNN!	4	527	943	3463	3542	
HgaI	!NNNNNNNNNNNGCGTC	3	2086	3981	4559		
HgiAI	GWGCW!C	4	1348	2866	3056	4361	
HgiEIII	!ACCNNNNNNNGGT	1	4083				
HhaI	GCG!C	28	598	626	872	900	939
			941	1229	1257	2019	2027
			2053	2075	2084	2097	2128
			2747	2755	2819	2856	3122
			3382	3558	3644	4047	4156
			4330	4430	4497		
HinI	GR!CGYC	8	123	176	259	445	624
			2753	3455	3534		
HinPII	G!CGC	28	596	624	870	898	937
			939	1227	1255	2017	2025
			2051	2073	2082	2095	2126
			2745	2753	2817	2854	3120
			3380	3556	3642	4045	4154
			4328	4428	4495		
HincII	GTY!RAC	2	1369	1516			
HindIII	A!AGCTT	1	1350				
HinfI	G!ANTC	13	411	973	1332	1796	1818
			2228	3238	3372	3424	3482
			3519	4301	4697		
HpaI	GTT!AAC	1	1516				
HpaII	C!CGG	23	602	639	762	990	1329
			1385	1395	1971	2658	2735
			2757	2785	2916	3006	3073
			3254	3529	3537	3629	4103
			4293	4319	4466		

HphI	GGTGANNNNNNNN!	6	377 3940	958	982	1111	3824
HphI	!NNNNNNNTCACC	4	628	1291	1870	2931	
KasI	G!GCGCC	2	623	2752			
KpnI	GGTAC!C	1	1377				
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	3	2220	3097	3307		
KspI	CCGC!GG	1	1380				
MaeI	C!TAG	8	593 3591	1401 3925	1602 4178	2020	2576
MaeII	A!CGT	14	123 445 2869	135 1813 3056	176 1825 3837	259 1868 3970	340 1978
MaeIII	!GTNAC	13	62 2042 4135	149 2054 4251	498 2873 4314	1282 3179	1541 3869
MamI	GATNN!NNATC	2	1415	2612			
MboI	!GATC	25	586 1337 2594 3092 3933	657 1388 2613 3170 4011	838 1397 2924 3546 4019	1258 1410 3002 3562 4030	1266 2590 3083 3921 4105
MboII	GAAGANNNNNNNN!	5	2009	2237	3114	3324	3943
MboII	!NNNNNNNTCTTC	7	853 3771	898 4014	901	3404	3571
McrI	!CGRYCG	3	602	2658	4333		
MfeI	!CAATTG	1	1502				
MflI	R!GATCY	11	586 3170 4030	1337 3562	1388 3921	1397 3933	2924 4019
MluI	A!CGCGT	1	1639				
MmeI	!TCCRAC	1	1820				
MmeI	!GTYGGA	2	4296	4480			
MnlI	CCTCNNNNNNN!	10	1283 3354	1472 3560	1481 3874	2536 4244	2542 4568
MnlI	!NNNNNNNGAGG	22	541 936 2227 2563 3590	699 948 2505 2603 4493	705 999 2511 2667	873 1431 2548 2803	906 1896 2551 3160
MroI	T!CCGGA	1	1328				
MscI	TGG!CCA	3	684	797	2835		
MseI	T!TAA	15	8 1668 2077	1454 1680 3896	1515 1691 3910	1636 1708 3915	1657 1806 3967
MspI	C!CGG	23	602 1385 2757 3254 4293	639 1395 2785 3529 4319	762 1971 2916 3537 4466	990 2658 3006 3629	1329 2735 3073 4103
MstI	TGC!GCA	1	2855				
MstII	CC!TNAGG	3	879	2233	3865		
MunI	C!AATTG	1	1503				
MvaI	CC!WGG	17	91 968 2361 4525	284 1022 3140 4646	781 1084 3719	791 2289 3822	893 2344 4512
MvnI	CG!CG	19	61 1641	937 1677	939 2053	1257 2073	1379 2097

			2128	2819	3120	3558	3642
			3705	3774	4047	4628	
NaeI	GCC!GGC	3	1972	3255	3538		
NarI	GG!CGCC	2	624	2753			
NciI	CC!SGG	6	1385	1386	2757	2917	3530
			4294				
NcoI	C!CATGG	3	361	2482	3185		
NdeI	CA!TATG	1	235				
NdeII	!GATC	25	586	657	838	1258	1266
			1337	1388	1397	1410	2590
			2594	2613	2924	3002	3083
			3092	3170	3546	3562	3921
			3933	4011	4019	4030	4105
NgoMI	G!CCGGC	3	1970	3253	3536		
NheI	G!CTAGC	1	592				
NlaIII	CATG!	20	305	365	616	847	877
			1267	1312	2183	2323	2395
			2486	2627	2972	3158	3189
			3215	3571	3955	4675	4726
NlaIV	GGN!NCC	19	468	625	1242	1375	1383
			1390	1908	1929	1941	2132
			2293	2365	2754	2789	3637
			3716	3761	4604	4643	
NotI	GC!GGCCGC	0					
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	2325	2397	4728		
Nsp(7524)I	RCATG!Y	4	2323	2395	3158	4675	
Nsp(7524)V	TT!CGAA	2	1355	3435			
NspBII	CMG!CKG	6	1144	1379	2251	2859	4088
			4333				
NspII	GDGCH!C	12	651	780	1348	1385	1944
			2699	2792	2866	3056	3118
			3828	4361			
NspIII	C!YCGRG	3	1274	1341	1384		
NspIV	G!GNCC	14	84	277	783	1111	1184
			1240	1381	1382	1860	3269
			3714	3760	3818	3851	
NunII	GG!CGCC	2	624	2753			
PacI	TTAAT!TAA	0					
Paer71	C!TCGAG	2	1274	1341			
PalI	GG!CC	27	85	278	684	784	797
			1073	1082	1148	1185	1383
			1719	1861	2516	2522	2531
			2574	2661	2835	3226	3253
			3762	3819	3852	4197	4631
			4649	4660			
PflMI	CCANNNN!NTGG	1	1309				
PleI	GAGTCNNNN!	3	981	1804	2236		
PleI	!NNNNNGACTC	5	405	1326	1812	3418	4295
PmaCI	CAC!GTG	0					
PmeI	GTTT!AAAC	0					
PmlI	CAC!GTG	0					
PpuMI	RG!GWCCY	1	1240				
PshAI	GACNN!NNGTC	0					
PstI	CTGCA!G	1	1366				
PvuI	CGAT!CG	0					
PvuII	CAG!CTG	3	1144	2251	2859		
RmaI	C!TAG	8	593	1401	1602	2020	2576
			3591	3925	4178		

RsaI	GT!AC	11	220 541 3059	245 1040	300 1064	333 1322	384 1375
RsrI	G!AATTC	1	1357				
RsrII	CG!GWCCG	1	3269				
SacI	GAGCT!C	1	1348				
SacII	CCGC!GG	1	1380				
SalI	G!TCGAC	1	1367				
Sau3AI	!GATC	25	586 1337 2594 3092 3933	657 1388 2613 3170 4011	838 1397 2924 3546 4019	1258 1410 3002 3562 4030	1266 2590 3083 3921 4105
Sau96I	G!GNCC	14	84 1240 3714	277 1381 3760	783 1382 3818	1111 1860 3851	1184 3269
SauI	CC!TNAGG	3	879	2233	3865		
ScaI	AGT!ACT	0					
ScrFI	CC!NGG	23	91 968 2289 3140 4512	284 1022 2344 3530 4525	781 1084 2361 3719 4646	791 1385 2757 3822	893 1386 2917 4294
SduI	GDGCH!C	12	651 2699 3828	780 2792 4361	1348 2866	1385 3056	1944 3118
SecI	C!CNNGG	24	361 1021 2287 2575 3718	779 1236 2359 2916 3820	789 1303 2482 3185 3821	942 1377 2517 3590 4511	966 1384 2526 3717
SexI	!CTCGAG	2	1273	1340			
SfaNI	GCATCNNNNN!	9	651 2404	987 3052	1101 3116	1569 3391	2332
SfaNI	!NNNNNNNNNGATGC	6	358 4574	709	2711	2966	3182
SfcI	C!TRYAG	4	1362	2088	4215	4406	
SfiI	GGCCNNNN!NGGCC	2	1079	2528			
SfuI	TT!CGAA	2	1355	3435			
SgrAI	CR!CCGGYG	0					
SinI	G!GWCC	4	1111	1240	3269	3714	
SmaI	CCC!GGG	1	1386				
SnaBI	TAC!GTA	1	341				
SnaI	!GTATAC	0					
SnoI	G!TGCAC	1	4357				
SpeI	A!CTAGT	0					
SphI	GCATG!C	3	2323	2395	3158		
SplI	C!GTACG	0					
SrfI	GCCC!GGGC	0					
Sse8387I	CCTGCA!GG	0					
SspI	AAT!ATT	2	1661	2214			
SstI	GAGCT!C	1	1348				
SstII	CCGC!GG	1	1380				
SstIII	!ACGT	14	122 444 2868	134 1812 3055	175 1824 3836	258 1867 3969	339 1977
StuI	AGG!CCT	1	2574				
StyI	C!CWGG	6	361 3590	1236	2482	2575	3185

StySJI	!GAGNNNNNNNGTRC	0						
StySJI	!GYACNNNNNNNCTC	0						
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	19	660	912	954	981	1275	
			1342	1355	1368	1902	2593	
			2866	3022	3046	3082	3244	
			3435	3480	3735	4573		
TaqII	GACCGANNNNNNNNNNN!	3	1773	2790	3458			
TaqII	!NNNNNNNNNTCGGTC	1	3702					
TaqII	CACCCANNNNNNNNNNN!	2	1235	1316				
TaqII	!NNNNNNNNNTGGGTG	3	173	2657	3249			
TfiI	G!AWTC	5	3238	3372	3482	3519	4697	
ThaI	CG!CG	19	61	937	939	1257	1379	
			1641	1677	2053	2073	2097	
			2128	2819	3120	3558	3642	
			3705	3774	4047	4628		
Tsp45I	!GTSAC	4	1282	2042	2873	3179		
TspEI	!AATT	11	19	1357	1503	1567	1645	
			1671	1682	2329	2401	2493	
			3911					
Tth111I	GACN!NNGTC	1	2871					
Tth111II	CAARCANNNNNNNNNNN!	1	4080					
Tth111III	!NNNNNNNNNTGYTTG	3	3188	4048	4087			
TthHB8I	T!CGA	19	660	912	954	981	1275	
			1342	1355	1368	1902	2593	
			2866	3022	3046	3082	3244	
			3435	3480	3735	4573		
Van91I	CCANNNN!NTGG	1	1309					
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	1400					
XcmI	!CCANNNNNNNNNNTCG	2	1055	3028				
XcyI	C!CCGGG	1	1384					
XhoI	C!TCGAG	2	1274	1341				
XhoII	R!GATCY	11	586	1337	1388	1397	2924	
			3170	3562	3921	3933	4019	
			4030					
XmaI	C!CCGGG	1	1384					
XmaIII	C!GGCCG	1	2659					
XmnI	GAANN!NNTTC	0						
XorII	CGAT!CG	0						