

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	2	1952	3409			
AatII	GACGT!C	5	126	179	262	448	1136
Acc65I	G!GTACC	0					
AccI	GT!MKAC	2	641	2103			
AccII	CG!CG	16	61	1039	2237	2512	2888
			2908	2932	2963	3654	3955
			4393	4477	4540	4609	4882
			5463				
AccIII	T!CCGGA	1	1203				
AciI	G!CGG	29	371	404	529	1039	1233
			1671	1983	2025	2231	2848
			2875	2963	3073	3654	3718
			3819	3822	4102	4157	4255
			4393	4540	4680	4923	5168
			5289	5308	5435	5463	
AciI	C!CGC	35	59	87	99	113	280
			508	590	1836	1974	2076
			2235	2903	2906	2920	3010
			3258	3270	3279	3291	3301
			3312	3358	3497	3560	4062
			4107	4173	4199	4324	4327
			4475	4607	4914	5058	5554
AcyI	GR!CGYC	9	123	176	259	445	1133
			2100	3588	4290	4369	
AfaI	GT!AC	12	220	245	300	333	384
			541	668	1493	1568	1718
			1970	3894			
AflI	G!GWCC	2	4104	4549			
AflII	C!TTAAG	1	2470				
AflIII	A!CRYGT	1	5506				
AgeI	A!CCGGT	1	1504				
AhaI	CC!SGG	6	810	2097	3592	3752	4365
			5129				
AhaII	GR!CGYC	9	123	176	259	445	1133
			2100	3588	4290	4369	
AhaIII	TTT!AAA	3	2290	4732	4751		
AluI	AG!CT	23	567	619	625	763	893
			1276	1291	1354	1537	1564
			1710	2017	2367	2534	2791
			3086	3374	3694	4152	4949
			5206	5252	5342		
Alw44I	G!TGCAC	3	818	1603	5192		
AlwI	GGATCNNNN!	8	1506	3456	3767	4013	4405
			4764	4862	4948		
AlwI	!NNNNNGATCC	7	581	1493	2058	3832	4376
			4763	4860			
AlwNI	CAGNNN!CTG	3	2190	4689	5097		
AocI	CC!TNAGG	4	1446	1782	3068	4700	
AocII	GDGCH!C	15	621	822	1356	1580	1607
			1640	2217	2779	3534	3627
			3701	3891	3953	4663	5196
Aor51HI	AGC!GCT	1	597				
AosI	TGC!GCA	1	3690				
AosII	GR!CGYC	9	123	176	259	445	1133
			2100	3588	4290	4369	
ApaI	GGGCC!C	0					

ApaI	G!TGCAC	3	818	1603	5192			
ApoI	R!AATTY	7	630	950	989	1254	2402	
			2506	2517				
ApyI	CC!WGG	22	91	284	842	1049	1074	
			1364	1484	1496	1700	1748	
			1844	1859	1988	3124	3179	
			3196	3975	4554	4657	5347	
			5360	5481				
AquI	C!YCGRG	1	614					
AscI	GG!CGCGCC	0						
AseI	AT!TAAT	1	8					
AsnI	AT!TAAT	1	8					
Asp700I	GAANN!NNTTC	0						
Asp718I	G!GTACC	0						
AspHI	GWGCW!C	7	621	822	1356	1607	3701	
			3891	5196				
AspI	GACN!NNGTC	1	3706					
AsuI	G!GNCC	15	84	277	1045	1145	1186	
			1673	1916	2008	2029	2695	
			4104	4549	4595	4653	4686	
AsuII	TT!CGAA	2	628	4270				
AvaI	C!YCGRG	1	614					
AvaII	G!GWCC	2	4104	4549				
AvaIII	ATGCA!T	3	3160	3232	5563			
AviII	TGC!GCA	1	3690					
AvrI	!CYCGRG	1	613					
AvrII	C!CTAGG	2	3410	4425				
AxyI	CC!TNAGG	4	1446	1782	3068	4700		
BalI	TGG!CCA	2	2183	3670				
BamHI	G!GATCC	1	1498					
BanI	G!GYRCC	6	466	1577	1637	2741	3587	
			3622					
BanII	GRGCY!C	5	621	1356	2779	3953	4663	
BanIII	AT!CGAT	1	3428					
BbeI	GGCGC!C	1	3591					
BbiII/AcyI	GR!CGYC	9	123	176	259	445	1133	
			2100	3588	4290	4369		
BbrPI	CAC!GTG	0						
BbsI	!GTCTTC	0						
BbsI	!GAAGAC	4	1477	1816	2049	2117		
BbvI	GCAGCNNNNNNNN!	10	788	902	1734	2176	2376	
			3661	3703	4224	4677	5177	
BbvI	!NNNNNNNNNNNGCTGC	9	870	1853	2870	3535	3719	
			3812	4878	5084	5087		
BbvII	GAAGACNN!	4	1485	1824	2057	2125		
BbvII	!NNNNNNGTCTTC	0						
BcefI	ACGCGCNNNNNNNNNN!	6	1735	1879	1930	1996	4014	
			4501					
BcefI	!NNNNNNNNNNNGCCGT	4	642	2730	3548	5007		
BcgI	!GCANNNNNNTCG	0						
BcgI	!CGANNNNNNTGC	0						
BclI	T!GATCA	1	1078					
BcnI	CC!SGG	6	810	2097	3592	3752	4365	
			5129					
BfaI	C!TAG	10	593	851	2202	2243	2437	
			2855	3411	4426	4760	5013	
BfrI	C!TTAAG	1	2470					
BglI	GCCNNNN!NGGC	6	91	213	284	653	1992	
			3363					

BglII	A!GATCT	2	610	1083				
BinI	!NNNNNGATCC	7	581	1493	2058	3832	4376	
			4763	4860				
BinI	GGATCNNNN!	8	1506	3456	3767	4013	4405	
			4764	4862	4948			
BlnI	C!CTAGG	2	3410	4425				
BmyI	GDGCH!C	15	621	822	1356	1580	1607	
			1640	2217	2779	3534	3627	
			3701	3891	3953	4663	5196	
BpmI	!CTCCAG	2	1855	4385				
BpmI	!CTGGAG	2	1306	4406				
Bpu1102I	GC!TNAGC	1	1175					
BsaAI	YAC!GTR	3	341	2704	3892			
BsaBI	GATNN!NNATC	2	2250	3447				
BsaHI	GR!CGYC	9	123	176	259	445	1133	
			2100	3588	4290	4369		
BsaI	GGTCTCN!	0						
BsaI	!NNNNNGAGACC	2	2144	4577				
BsaJI	C!CNNGG	23	361	651	808	1072	1515	
			1611	1746	1747	1842	3122	
			3194	3317	3352	3361	3410	
			3751	4020	4425	4552	4553	
			4655	4656	5346			
BseAI	T!CCGGA	1	1203					
BsePI	!GCGCGC	0						
BsgI	!GTGCAG	0						
BsgI	!CTGCAC	1	733					
BsiEI	CGRY!CG	4	1509	2235	3497	5172		
BsiWI	C!GTACG	0						
BsiYI	CCNNNNN!NNGG	27	50	793	1033	1073	1170	
			1276	1634	1747	1913	1979	
			2600	2926	3318	3569	4113	
			4394	4426	4427	4558	4593	
			4643	4644	4675	5028	5307	
			5473	5491				
BslI	CCNNNNN!NNGG	27	50	793	1033	1073	1170	
			1276	1634	1747	1913	1979	
			2600	2926	3318	3569	4113	
			4394	4426	4427	4558	4593	
			4643	4644	4675	5028	5307	
			5473	5491				
BsmAI	!GTCTC	1	429					
BsmAI	!GAGAC	6	1240	2149	3017	3435	4454	
			4582					
BsmI	GAATGCN!	1	2339					
BsmI	G!CATTC	2	1197	2432				
Bsp1286I	GDGCH!C	15	621	822	1356	1580	1607	
			1640	2217	2779	3534	3627	
			3701	3891	3953	4663	5196	
BspDI	AT!CGAT	1	3428					
BspEI	T!CCGGA	1	1203					
BspHI	T!CATGA	2	3014	4786				
BspMI	ACCTGCNNNN!	2	3856	4306				
BspMI	!NNNNNNNNGCAGGT	1	3475					
BspMII	T!CCGGA	1	1203					
BspWI	GCNNNNN!NNGC	34	91	213	245	284	377	
			401	653	1039	1176	1407	
			1561	1992	2815	2845	2872	
			2914	2916	3155	3227	3278	

			3357	3363	3579	3663	3686
			3825	3831	3948	4031	4298
			4485	4689	4888	5460	
BsrFI	R!CCGGY	6	1170	1504	2805	3907	4088
			4371				
BsrI	ACTGGN!	4	2127	3532	3733	4979	
BsrI	C!CAGT	7	296	684	787	2617	3295
			5091	5104			
BssHII	G!CGCGC	0					
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	2	628	4270			
BstEII	G!GTNACC	0					
BstI	G!GATCC	1	1498				
BstNI	CC!WGG	22	91	284	842	1049	1074
			1364	1484	1496	1700	1748
			1844	1859	1988	3124	3179
			3196	3975	4554	4657	5347
			5360	5481			
BstPI	G!GTNACC	0					
BstUI	CG!CG	16	61	1039	2237	2512	2888
			2908	2932	2963	3654	3955
			4393	4477	4540	4609	4882
			5463				
BstXI	CCANNNNN!NTGG	0					
BstYI	R!GATCY	12	586	610	1083	1498	2063
			3759	4005	4397	4756	4768
			4854	4865			
Bsu36I	CC!TNAGG	4	1446	1782	3068	4700	
CbiI	TT!CGAA	2	628	4270			
CcrI	C!TCGAG	1	614				
CelII	GC!TNAGC	1	1175				
CfoI	GCG!C	23	598	1039	1060	2854	2862
			2888	2910	2919	2932	2963
			3582	3590	3654	3691	3957
			4217	4393	4479	4882	4991
			5165	5265	5332		
Cfr10I	R!CCGGY	6	1170	1504	2805	3907	4088
			4371				
Cfr13I	G!GNCC	15	84	277	1045	1145	1186
			1673	1916	2008	2029	2695
			4104	4549	4595	4653	4686
Cfr9I	C!CCGGG	0					
CfrI	Y!GGCCR	9	654	1168	1234	2181	2232
			3494	3668	4059	4086	
ClaI	AT!CGAT	1	3428				
CpoI	CG!GWCCG	1	4104				
Csp45I	TT!CGAA	2	628	4270			
CspI	CG!GSCCG	0					
CviJI	RG!CY	111	43	85	94	278	567
			619	625	656	763	779
			813	845	893	1047	1146
			1170	1174	1187	1223	1236
			1276	1291	1306	1354	1386
			1537	1564	1627	1674	1703
			1710	1788	1854	1866	1917
			1944	1952	1986	2009	2017
			2031	2141	2177	2183	2234
			2257	2367	2534	2554	2696
			2777	2791	2805	3086	3127

			3199	3322	3351	3357	3366
			3374	3397	3409	3415	3496
			3513	3521	3548	3573	3657
			3666	3670	3694	3732	3808
			3825	3844	3907	3951	3961
			3978	4061	4088	4092	4129
			4152	4168	4348	4375	4436
			4558	4597	4654	4661	4668
			4687	4698	4949	4978	5021
			5032	5097	5176	5201	5206
			5252	5342	5440	5466	5484
			5495				
CvnI	CC!TNAGG	4	1446	1782	3068	4700	
DdeI	C!TNAG	14	607	695	1175	1396	1446
			1524	1782	2079	3068	3370
			4251	4700	4823	5232	
DpnI	GA!TC	25	588	612	1080	1085	1500
			1889	2039	2065	2247	3427
			3431	3450	3761	3839	3920
			3929	4007	4383	4399	4758
			4770	4848	4856	4867	4942
DpnII	!GATC	25	586	610	1078	1083	1498
			1887	2037	2063	2245	3425
			3429	3448	3759	3837	3918
			3927	4005	4381	4397	4756
			4768	4846	4854	4865	4940
DraI	TTT!AAA	3	2290	4732	4751		
DraII	RG!GNCCY	1	4686				
DraIII	CACNNN!GTG	2	2173	2704			
DrdI	GACNNNN!NNGTC	3	2659	3615	5404		
DsaI	C!CRYGG	5	361	651	1515	3317	4020
DsaV	!CCNGG	28	89	282	808	840	1047
			1072	1362	1482	1494	1698
			1746	1842	1857	1986	2095
			3122	3177	3194	3590	3750
			3973	4363	4552	4655	5127
			5345	5358	5479		
EaeI	Y!GGCCR	9	654	1168	1234	2181	2232
			3494	3668	4059	4086	
EagI	C!GGCCG	2	2232	3494			
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	1	802				
EarI	!GAAGAG	5	1462	1531	3059	3936	4146
Ecl136II	GAG!CTC	2	619	1354			
EclXI	C!GGCCG	2	2232	3494			
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	2	2144	4577			
Eco47I	G!GWCC	2	4104	4549			
Eco47IIII	AGC!GCT	1	597				
Eco52I	C!GGCCG	2	2232	3494			
Eco57I	!CTGAAG	3	2011	3712	4144		
Eco57I	!CTTCAG	2	1329	4978			
Eco81I	CC!TNAGG	4	1446	1782	3068	4700	
EcoNI	CCTNN!NNNAGG	1	1071				
EcoO109I	RG!GNCCY	1	4686				
EcoO65I	G!GTNACC	0					
EcoRI	G!AATTC	1	630				
EcoRII	!CCWGG	22	89	282	840	1047	1072

			1362	1482	1494	1698	1746
			1842	1857	1986	3122	3177
			3194	3973	4552	4655	5345
			5358	5479			
EcoRV	GAT!ATC	0					
EcoT14I	C!CWWGG	7	361	651	1515	3317	3410
			4020	4425			
EcoT22I	ATGCA!T	3	3160	3232	5563		
EcoT38I	GRGCY!C	5	621	1356	2779	3953	4663
EheI	GGC!GCC	1	3589				
Esp3I	CGTCTCN!	0					
Esp3I	!NNNNNGAGACG	0					
EspI	GC!TNAGC	1	1175				
FbaI	T!GATCA	1	1078				
FinI	!GTCCC	5	1134	3118	3190	3254	4549
FinI	!GGGAC	6	161	312	480	3724	4256
			4365				
Fnu4HI	GC!NGC	40	777	884	891	1040	1234
			1672	1723	1867	1984	2165
			2232	2235	2365	2884	2906
			2920	3358	3497	3549	3560
			3650	3655	3692	3733	3820
			3823	3826	4062	4158	4199
			4213	4327	4666	4681	4892
			5098	5101	5166	5309	5464
FokI	GGATGNNNNNNNN!	5	1313	3455	3912	3937	4391
FokI	!NNNNNNNNNNNNNCATCC	7	851	1397	1594	1682	1739
			1894	3261			
FspI	TGC!GCA	1	3690				
GdiII	!NNNNNYGGCCG	6	648	1162	2226	3488	4053
			4080				
GdiII	CGGCCRN!	3	1240	2238	3500		
GsuI	!CTCCAG	2	1855	4385			
GsuI	!CTGGAG	2	1306	4406			
HaeI	WGG!CCW	8	845	1952	2183	3409	3670
			5032	5484	5495		
HaeII	RGCGC!Y	6	599	1061	2855	2863	3591
			5266				
HaeIII	GG!CC	35	85	278	656	845	1047
			1146	1170	1187	1236	1674
			1917	1952	1986	2009	2031
			2141	2183	2234	2554	2696
			3351	3357	3366	3409	3496
			3670	4061	4088	4597	4654
			4687	5032	5466	5484	5495
HapII	C!CGG	22	602	810	1171	1204	1505
			2097	2806	3493	3570	3592
			3620	3751	3841	3908	4089
			4364	4372	4464	4938	5128
			5154	5301			
HgaI	GACGCNNNNN!	4	527	1007	4298	4377	
HgaI	!NNNNNNNNNNNGCGTC	4	2089	2921	4816	5394	
HgiAI	GWGCW!C	7	621	822	1356	1607	3701
			3891	5196			
HgiEII	!ACCNNNNNNGGT	1	4918				
HhaI	GCG!C	23	598	1039	1060	2854	2862
			2888	2910	2919	2932	2963
			3582	3590	3654	3691	3957
			4217	4393	4479	4882	4991

			5165	5265	5332		
Hin1I	GR!CGYC	9	123	176	259	445	1133
			2100	3588	4290	4369	
HinPII	G!CGC	23	596	1037	1058	2852	2860
			2886	2908	2917	2930	2961
			3580	3588	3652	3689	3955
			4215	4391	4477	4880	4989
			5163	5263	5330		
HincII	GTY!RAC	4	642	724	913	2351	
HindIII	A!AGCTT	1	623				
HinfI	G!ANTC	20	411	605	697	728	1210
			1434	1653	1803	2130	2239
			2631	2653	3063	4073	4207
			4259	4317	4354	5136	5532
HpaI	GTT!AAC	1	2351				
HpaII	C!CGG	22	602	810	1171	1204	1505
			2097	2806	3493	3570	3592
			3620	3751	3841	3908	4089
			4364	4372	4464	4938	5128
			5154	5301			
HphI	GGTGANNNNNNNN!	5	377	1088	1909	4659	4775
HphI	!NNNNNNNTCACC	3	1798	2705	3766		
KasI	G!GCGCC	1	3587				
KpnI	GGTAC!C	0					
Ksp632I	CTCTTCN!	1	809				
Ksp632I	!NNNNGAAGAG	5	1458	1527	3055	3932	4142
KspI	CCGC!GG	0					
MaeI	C!TAG	10	593	851	2202	2243	2437
			2855	3411	4426	4760	5013
MaeII	A!CGT	19	123	135	176	259	340
			445	1053	1133	1318	1881
			2157	2648	2660	2703	2813
			3704	3891	4672	4805	
MaeIII	!GTNAC	16	62	149	498	679	1319
			1343	1804	2376	2877	2889
			3708	4014	4704	4970	5086
			5149				
MamI	GATNN!NNATC	2	2250	3447			
MboI	!GATC	25	586	610	1078	1083	1498
			1887	2037	2063	2245	3425
			3429	3448	3759	3837	3918
			3927	4005	4381	4397	4756
			4768	4846	4854	4865	4940
MboII	GAAGANNNNNNNN!	14	913	1219	1475	1490	1544
			1829	1942	2062	2130	2844
			3072	3949	4159	4778	
MboII	!NNNNNNNTCTTC	6	796	1078	4239	4406	4606
			4849				
McrI	!CGRYCG	4	1505	2231	3493	5168	
MfeI	!CAATTG	1	2337				
MflI	R!GATCY	12	586	610	1083	1498	2063
			3759	4005	4397	4756	4768
			4854	4865			
MluI	A!CGCGT	0					
MmeI	!TCCRAC	2	1909	2655			
MmeI	!GTYGGA	2	5131	5315			
MnlI	CCTCNNNNNNNN!	21	749	1038	1074	1174	1224
			1424	1686	1865	1880	1963
			2096	2207	2307	2316	3371

			3377 5403	4189	4395	4709	5079
MnlI	!NNNNNNNGAGG	28	541	818	1121	1419	1440
			1566	1605	1626	1659	1776
			1885	1941	2130	2163	2266
			2731	3062	3340	3346	3383
			3386	3398	3438	3502	3638
			3995	4425	5328		
MroI	T!CCGGA	1	1203				
MscI	TGG!CCA	2	2183	3670			
MseI	T!TAA	19	8	984	1542	1767	2223
			2289	2350	2471	2492	2503
			2515	2526	2543	2641	2912
			4731	4745	4750	4802	
MspI	C!CGG	22	602	810	1171	1204	1505
			2097	2806	3493	3570	3592
			3620	3751	3841	3908	4089
			4364	4372	4464	4938	5128
			5154	5301			
MstI	TGC!GCA	1	3690				
MstII	CC!TNAGG	4	1446	1782	3068	4700	
MunI	C!AATTG	1	2338				
MvaI	CC!WGG	22	91	284	842	1049	1074
			1364	1484	1496	1700	1748
			1844	1859	1988	3124	3179
			3196	3975	4554	4657	5347
			5360	5481			
MvnI	CG!CG	16	61	1039	2237	2512	2888
			2908	2932	2963	3654	3955
			4393	4477	4540	4609	4882
			5463				
NaeI	GCC!GGC	4	1172	2807	4090	4373	
NarI	GG!CGCC	1	3588				
NciI	CC!SGG	6	810	2097	3592	3752	4365
			5129				
NcoI	C!CATGG	5	361	651	1515	3317	4020
NdeI	CA!TATG	1	235				
NdeII	!GATC	25	586	610	1078	1083	1498
			1887	2037	2063	2245	3425
			3429	3448	3759	3837	3918
			3927	4005	4381	4397	4756
			4768	4846	4854	4865	4940
NgoMI	G!CCGGC	4	1170	2805	4088	4371	
NheI	G!CTAGC	1	592				
NlaIII	CATG!	32	305	365	655	707	720
			796	802	860	1159	1241
			1373	1519	1555	1561	1573
			1651	1717	1797	2008	3018
			3158	3230	3321	3462	3807
			3993	4024	4050	4406	4790
			5510	5561			
NlaIV	GGN!NCC	19	468	1188	1500	1579	1639
			2030	2743	2764	2776	2967
			3128	3200	3589	3624	4472
			4551	4596	5439	5478	
NotI	GC!GGCCGC	1	2232				
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	3160	3232	5563		
Nsp(7524)I	RCATG!Y	7	802	1159	1555	3158	3230

			3993	5510				
Nsp(7524)V	TT!CGAA	2	628	4270				
NspBII	CMG!CKG	6	1276	1976	3086	3694	4923	
			5168					
NspII	GDGCH!C	15	621	822	1356	1580	1607	
			1640	2217	2779	3534	3627	
			3701	3891	3953	4663	5196	
NspIII	C!YCGRG	1	614					
NspIV	G!GNCC	15	84	277	1045	1145	1186	
			1673	1916	2008	2029	2695	
			4104	4549	4595	4653	4686	
NunII	GG!CGCC	1	3588					
PacI	TTAAT!TAA	0						
Paer71	C!TCGAG	1	614					
PalI	GG!CC	35	85	278	656	845	1047	
			1146	1170	1187	1236	1674	
			1917	1952	1986	2009	2031	
			2141	2183	2234	2554	2696	
			3351	3357	3366	3409	3496	
			3670	4061	4088	4597	4654	
			4687	5032	5466	5484	5495	
PflMI	CCANNNN!NTGG	2	793	1276				
PleI	GAGTCNNNN!	4	705	1811	2639	3071		
PleI	!NNNNNGACTC	7	405	599	722	2233	2647	
			4253	5130				
PmaCI	CAC!GTG	0						
PmeI	GTTT!AAAC	0						
PmlI	CAC!GTG	0						
PpuMI	RG!GWCCY	0						
PshAI	GACNN!NNGTC	1	2156					
PstI	CTGCA!G	1	639					
PvuI	CGAT!CG	0						
PvuII	CAG!CTG	3	1276	3086	3694			
RmaI	C!TAG	10	593	851	2202	2243	2437	
			2855	3411	4426	4760	5013	
RsaI	GT!AC	12	220	245	300	333	384	
			541	668	1493	1568	1718	
			1970	3894				
RsrI	G!AATTC	1	630					
RsrII	CG!GWCCG	1	4104					
SacI	GAGCT!C	2	621	1356				
SacII	CCGC!GG	0						
SalI	G!TCGAC	1	640					
Sau3AI	!GATC	25	586	610	1078	1083	1498	
			1887	2037	2063	2245	3425	
			3429	3448	3759	3837	3918	
			3927	4005	4381	4397	4756	
			4768	4846	4854	4865	4940	
Sau96I	G!GNCC	15	84	277	1045	1145	1186	
			1673	1916	2008	2029	2695	
			4104	4549	4595	4653	4686	
SauI	CC!TNAGG	4	1446	1782	3068	4700		
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	28	91	284	810	842	1049	
			1074	1364	1484	1496	1700	
			1748	1844	1859	1988	2097	
			3124	3179	3196	3592	3752	
			3975	4365	4554	4657	5129	
			5347	5360	5481			

SduI	GDGCH!C	15	621 1640 3701	822 2217 3891	1356 2779 3953	1580 3534 4663	1607 3627 5196
SecI	C!CNNGG	23	361 1611 3194 3751 4655	651 1746 3317 4020 4656	808 1747 3352 4425 5346	1072 1842 3361 4552	1515 3122 3410 4553
SexI	!CTCGAG	1	613				
SfaNI	GCATCNNNNN!	10	1410 3167	1419 3239	1477 3887	1761 3951	2404 4226
SfaNI	!NNNNNNNNNGATGC	9	358 3546	1237 3801	1914 4017	1953 5409	2082
SfcI	C!TRYAG	5	635	927	2923	5050	5241
SfiI	GGCCNNNN!NGGCC	1	3363				
SfuI	TT!CGAA	2	628	4270			
SgrAI	CR!CCGGYG	0					
SinI	G!GWCC	2	4104	4549			
SmaI	CCC!GGG	0					
SnaBI	TAC!GTA	1	341				
SnaI	!GTATAC	0					
SnoI	G!TGCAC	3	818	1603	5192		
SpeI	A!CTAGT	0					
SphI	GCATG!C	3	3158	3230	3993		
SplI	C!GTACG	0					
SrfI	GCCC!GGGC	0					
Sse8387I	CCTGCA!GG	0					
SspI	AAT!ATT	2	2496	3049			
SstI	GAGCT!C	2	621	1356			
SstII	CCGC!GG	0					
SstIII	!ACGT	19	122 444 2156 3703	134 1052 2647 3890	175 1132 2659 4671	258 1317 2702 4804	339 1880 2812
StuI	AGG!CCT	2	1952	3409			
StyI	C!CWWGG	7	361 4020	651 4425	1515	3317	3410
StySJI	!GAGNNNNNNNGTRC	0					
StySJI	!GYACNNNNNNNCTC	0					
SwaI	ATTT!AAAT	0					
TaqI	T!CGA	18	615 1692 3857 4315	628 2160 3881 4570	641 2737 3917 5408	1350 3428 4079	1665 3701 4270
TaqII	GACCGANNNNNNNNNNN!	3	2608	3625	4293		
TaqII	!NNNNNNNNNTCGGTC	1	4537				
TaqII	CACCCANNNNNNNNNNN!	2	1655	1760			
TaqII	!NNNNNNNNNTGGGTG	4	173	896	3492	4084	
TfiI	G!AWTC	9	1210 4207	1434 4317	1653 4354	2130 5532	4073
ThaI	CG!CG	16	61 2908 4393 5463	1039 2932 4477	2237 2963 4540	2512 3654 4609	2888 3955 4882
Tsp45I	!GTSAC	7	679 3708	1319 4014	1343	1804	2877
TspEI	!AATT	17	19 989 2480	630 1254 2506	935 2224 2517	950 2338 3164	961 2402 3236

			3328	4746				
Tth111I	GACN!NNGTC	1	3706					
Tth111II	CAARCANNNNNNNNNNNN!	2	903	4915				
Tth111III	!NNNNNNNNNTGYTTG	4	701	4023	4883	4922		
TthHB8I	T!CGA	18	615	628	641	1350	1665	
			1692	2160	2737	3428	3701	
			3857	3881	3917	4079	4270	
			4315	4570	5408			
Van91I	CCANNNN!NTGG	2	793	1276				
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	2242					
XcmI	!CCANNNNNNNNTCG	1	3863					
XcyI	C!CCGGG	0						
XhoI	C!TCGAG	1	614					
XhoII	R!GATCY	12	586	610	1083	1498	2063	
			3759	4005	4397	4756	4768	
			4854	4865				
XmaI	C!CCGGG	0						
XmaIII	C!GGCCG	2	2232	3494				
XmnI	GAANN!NNTTC	0						
XorII	CGAT!CG	0						