

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	2	1818	3275			
AatII	GACGT!C	4	126	179	262	448	
Acc65I	G!GTACC	0					
AccI	GT!MKAC	1	1969				
AccII	CG!CG	18	61	636	801	1362	2103
			2378	2754	2774	2798	2829
			3520	3821	4259	4343	4406
			4475	4748	5329		
AccIII	T!CCGGA	0					
AciI	G!CGG	29	371	404	529	636	1362
			1537	1849	1891	2097	2714
			2741	2829	2939	3520	3584
			3685	3688	3968	4023	4121
			4259	4406	4546	4789	5034
			5155	5174	5301	5329	
AciI	C!CGC	38	59	87	99	113	280
			508	590	640	776	799
			1702	1840	1942	2101	2769
			2772	2786	2876	3124	3136
			3145	3157	3167	3178	3224
			3363	3426	3928	3973	4039
			4065	4190	4193	4341	4473
			4780	4924	5420		
AcyI	GR!CGYC	8	123	176	259	445	1966
			3454	4156	4235		
AfaI	GT!AC	11	220	245	300	333	384
			541	1155	1434	1584	1836
			3760				
AflI	G!GWCC	5	1058	1072	1084	3970	4415
AflII	C!TTAAG	1	2336				
AflIII	A!CRYGT	1	5372				
AgeI	A!CCGGT	1	1370				
AhaI	CC!SGG	6	1110	1963	3458	3618	4231
			4995				
AhaII	GR!CGYC	8	123	176	259	445	1966
			3454	4156	4235		
AhaIII	TTT!AAA	3	2156	4598	4617		
AluI	AG!CT	22	567	619	625	923	957
			1125	1310	1403	1430	1576
			1883	2233	2400	2657	2952
			3240	3560	4018	4815	5072
			5118	5208			
Alw44I	G!TGCAC	2	1469	5058			
AlwI	GGATCNNNN!	8	1372	3322	3633	3879	4271
			4630	4728	4814		
AlwI	!NNNNNGATCC	8	581	779	1359	1924	3698
			4242	4629	4726		
AlwNI	CAGNNN!CTG	4	1178	2056	4555	4963	
AocI	CC!TNAGG	3	1648	2934	4566		
AocII	GDGCH!C	15	621	696	917	1446	1473
			1506	2083	2645	3400	3493
			3567	3757	3819	4529	5062
Aor51HI	AGC!GCT	1	597				
AosI	TGC!GCA	1	3556				
AosII	GR!CGYC	8	123	176	259	445	1966
			3454	4156	4235		

ApaI	GGGCC!C	0						
ApaLI	G!TGCAC	2	1469	5058				
ApoI	R!AATY	5	630	1282	2268	2372	2383	
ApyI	CC!WGG	20	91	284	856	951	1196	
			1232	1566	1614	1710	1725	
			1854	2990	3045	3062	3841	
			4420	4523	5213	5226	5347	
AquI	C!YCGRG	3	614	650	711			
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	5	621	1473	3567	3757	5062	
AspI	GACN!NNGTC	1	3572					
AsuI	G!GNCC	17	84	277	944	1058	1072	
			1084	1234	1539	1782	1874	
			1895	2561	3970	4415	4461	
			4519	4552				
AsuII	TT!CGAA	3	628	1120	4136			
AvaI	C!YCGRG	3	614	650	711			
AvaII	G!GWCC	5	1058	1072	1084	3970	4415	
AvaIII	ATGCA!T	3	3026	3098	5429			
AviII	TGC!GCA	1	3556					
AvrI	!CYCGRG	3	613	649	710			
AvrII	C!CTAGG	2	3276	4291				
AxyI	CC!TNAGG	3	1648	2934	4566			
BalI	TGG!CCA	2	2049	3536				
BamHI	G!GATCC	1	1364					
BanI	G!GYRCC	6	466	1443	1503	2607	3453	
			3488					
BanII	GRGCY!C	5	621	917	2645	3819	4529	
BanIII	AT!CGAT	1	3294					
BbeI	GGCGC!C	1	3457					
BbiII/AcyI	GR!CGYC	8	123	176	259	445	1966	
			3454	4156	4235			
BbrPI	CAC!GTG	1	965					
BbsI	!GTCTTC	2	1127	1250				
BbsI	!GAAGAC	3	1682	1915	1983			
BbvI	GCAGCNNNNNNNN!	10	932	1184	1600	2042	2242	
			3527	3569	4090	4543	5043	
BbvI	!NNNNNNNNNNNGCTGC	8	1719	2736	3401	3585	3678	
			4744	4950	4953			
BbvII	GAAGACNN!	3	1690	1923	1991			
BbvII	!NNNNNNGTCTTC	2	1121	1244				
BcefI	ACGGCNNNNNNNNNNN!	6	1601	1745	1796	1862	3880	
			4367					
BcefI	!NNNNNNNNNNNGCCGT	3	2596	3414	4873			
BcgI	!GCANNNNNNTCG	1	641					
BcgI	!CGANNNNNNTGC	0						
BclI	T!GATCA	1	925					
BcnI	CC!SGG	6	1110	1963	3458	3618	4231	
			4995					
BfaI	C!TAG	9	593	2068	2109	2303	2721	
			3277	4292	4626	4879		
BfrI	C!TTAAG	1	2336					
BglI	GCCNNNN!NGGC	5	91	213	284	1858	3229	
BglII	A!GATCT	1	610					
BinI	!NNNNNGATCC	8	581	779	1359	1924	3698	

			4242	4629	4726		
BinI	GGATCNNNN!	8	1372	3322	3633	3879	4271
			4630	4728	4814		
BlnI	C!CTAGG	2	3276	4291			
BmyI	GDGCH!C	15	621	696	917	1446	1473
			1506	2083	2645	3400	3493
			3567	3757	3819	4529	5062
BpmI	!CTCCAG	3	848	1721	4251		
BpmI	!CTGGAG	1	4272				
Bpu1102I	GC!TNAGC	0					
BsaAI	YAC!GTR	4	341	965	2570	3758	
BsaBI	GATNN!NNATC	2	2116	3313			
BsaHI	GR!CGYC	8	123	176	259	445	1966
			3454	4156	4235		
BsaI	GGTCTCN!	0					
BsaI	!NNNNNGAGACC	2	2010	4443			
BsaJI	C!CNNGG	22	361	649	710	1381	1477
			1612	1613	1708	2988	3060
			3183	3218	3227	3276	3617
			3886	4291	4418	4419	4521
			4522	5212			
BseAI	T!CCGGA	0					
BsePI	!GCGCGC	0					
BsgI	!GTGCAG	0					
BsgI	!CTGCAC	0					
BsiEI	CGRY!CG	5	640	1375	2101	3363	5038
BsiWI	C!GTACG	0					
BsiYI	CCNNNNN!NNGG	27	50	674	742	794	911
			1081	1500	1613	1779	1845
			2466	2792	3184	3435	3979
			4260	4292	4293	4424	4459
			4509	4510	4541	4894	5173
			5339	5357			
BslI	CCNNNNN!NNGG	27	50	674	742	794	911
			1081	1500	1613	1779	1845
			2466	2792	3184	3435	3979
			4260	4292	4293	4424	4459
			4509	4510	4541	4894	5173
			5339	5357			
BsmAI	!GTCTC	1	429				
BsmAI	!GAGAC	6	976	2015	2883	3301	4320
			4448				
BsmI	GAATGCN!	1	2205				
BsmI	G!CATTC	1	2298				
Bsp1286I	GDGCH!C	15	621	696	917	1446	1473
			1506	2083	2645	3400	3493
			3567	3757	3819	4529	5062
BspDI	AT!CGAT	1	3294				
BspEI	T!CCGGA	0					
BspHI	T!CATGA	5	781	1000	1264	2880	4652
BspMI	ACCTGCNNNN!	2	3722	4172			
BspMI	!NNNNNNNNGCAGGT	1	3341				
BspMII	T!CCGGA	0					
BspWI	GCNNNNN!NNGC	30	91	213	245	284	377
			401	1427	1858	2681	2711
			2738	2780	2782	3021	3093
			3144	3223	3229	3445	3529
			3552	3691	3697	3814	3897
			4164	4351	4555	4754	5326

BsrFI	R!CCGGY	6	941 4237	1370	2671	3773	3954
BsrI	ACTGGN!	4	1993	3398	3599	4845	
BsrI	C!CAGT	6	296 4970	1356	2483	3161	4957
BssHII	G!CGCGC	0					
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	3	628	1120	4136		
BstEII	G!GTNACC	0					
BstI	G!GATCC	1	1364				
BstNI	CC!WGG	20	91 1232 1854 4420	284 1566 2990 4523	856 1614 3045 5213	951 1710 3062 5226	1196 1725 3841 5347
BstPI	G!GTNACC	0					
BstUI	CG!CG	18	61 2378 3520 4475	636 2754 3821 4748	801 2774 4259 5329	1362 2798 4343	2103 2829 4406
BstXI	CCANNNNN!NTGG	1	1214				
BstYI	R!GATCY	11	586 3871 4731	610 4263	1364 4622	1929 4634	3625 4720
Bsu36I	CC!TNAGG	3	1648	2934	4566		
CbiI	TT!CGAA	3	628	1120	4136		
CcrI	C!TCGAG	1	614				
CelII	GC!TNAGC	0					
CfoI	GCG!C	21	598 2785 3520 4345 5198	2720 2798 3557 4748	2728 2829 3823 4857	2754 3448 4083 5031	2776 3456 4259 5131
Cfr10I	R!CCGGY	6	941 4237	1370	2671	3773	3954
Cfr13I	G!GNCC	17	84 1084 1895 4519	277 1234 2561 4552	944 1539 3970	1058 1782 4415	1072 1874 4461
Cfr9I	C!CCGGG	0					
CfrI	Y!GGCCR	8	637 3534	796 3925	2047 3952	2098	3360
ClaI	AT!CGAT	1	3294				
CpoI	CG!GWCCG	1	3970				
Csp45I	TT!CGAA	3	628	1120	4136		
CspI	CG!GSCCG	0					
CviJI	RG!CY	110	43 619 828 923 1162 1430 1654 1818 2007 2233 2657 3188 3263 3387	85 625 854 945 1199 1493 1720 1852 2043 2400 2671 3217 3275 3414	94 639 859 957 1235 1540 1732 1875 2049 2420 2952 3223 3281 3439	278 755 908 969 1310 1569 1783 1883 2100 2562 2993 3232 3362 3523	567 798 915 1125 1403 1576 1810 1897 2123 2643 3065 3240 3379 3532

			3536	3560	3598	3674	3691
			3710	3773	3817	3827	3844
			3927	3954	3958	3995	4018
			4034	4214	4241	4302	4424
			4463	4520	4527	4534	4553
			4564	4815	4844	4887	4898
			4963	5042	5067	5072	5118
			5208	5306	5332	5350	5361
CvnI	CC!TNAGG	3	1648	2934	4566		
DdeI	C!TNAG	11	607	1337	1390	1648	1945
			2934	3236	4117	4566	4689
			5098				
DpnI	GA!TC	27	588	612	786	927	1044
			1269	1366	1755	1905	1931
			2113	3293	3297	3316	3627
			3705	3786	3795	3873	4249
			4265	4624	4636	4714	4722
			4733	4808			
DpnII	!GATC	27	586	610	784	925	1042
			1267	1364	1753	1903	1929
			2111	3291	3295	3314	3625
			3703	3784	3793	3871	4247
			4263	4622	4634	4712	4720
			4731	4806			
DraI	TTT!AAA	3	2156	4598	4617		
DraII	RG!GNCCY	2	1072	4552			
DraIII	CACNNN!GTG	2	2039	2570			
DrdI	GACNNNN!NNGTC	3	2525	3481	5270		
DsaI	C!CRYGG	4	361	1381	3183	3886	
DsaV	!CCNGG	26	89	282	854	949	1108
			1194	1230	1564	1612	1708
			1723	1852	1961	2988	3043
			3060	3456	3616	3839	4229
			4418	4521	4993	5211	5224
			5345				
EaeI	Y!GGCCR	8	637	796	2047	2098	3360
			3534	3925	3952		
EagI	C!GGCCG	3	637	2098	3360		
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	1	764				
EarI	!GAAGAG	5	1037	1397	2925	3802	4012
Ecl136II	GAG!CTC	1	619				
EclXI	C!GGCCG	3	637	2098	3360		
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	2	2010	4443			
Eco47I	G!GWCC	5	1058	1072	1084	3970	4415
Eco47III	AGC!GCT	1	597				
Eco52I	C!GGCCG	3	637	2098	3360		
Eco57I	!CTGAAG	3	1877	3578	4010		
Eco57I	!CTTCAG	1	4844				
Eco81I	CC!TNAGG	3	1648	2934	4566		
EcoNI	CCTNN!NNNAGG	0					
EcoO109I	RG!GNCCY	2	1072	4552			
EcoO65I	G!GTNACC	0					
EcoRI	G!AATTC	1	630				
EcoRII	!CCWGG	20	89	282	854	949	1194
			1230	1564	1612	1708	1723
			1852	2988	3043	3060	3839

			4418	4521	5211	5224	5345
EcoRV	GAT!ATC	0					
EcoT14I	C!CWWGG	6	361	1381	3183	3276	3886
			4291				
EcoT22I	ATGCA!T	3	3026	3098	5429		
EcoT38I	GRGCY!C	5	621	917	2645	3819	4529
EheI	GGC!GCC	1	3455				
Esp3I	CGTCTCN!	0					
Esp3I	!NNNNNGAGACG	0					
EspI	GC!TNAGC	0					
FbaI	T!GATCA	1	925				
FinI	!GTCCC	6	1072	1203	2984	3056	3120
			4415				
FinI	!GGGAC	7	161	312	480	652	3590
			4122	4231			
Fnu4HI	GC!NGC	40	637	640	799	921	1173
			1538	1589	1733	1850	2031
			2098	2101	2231	2750	2772
			2786	3224	3363	3415	3426
			3516	3521	3558	3599	3686
			3689	3692	3928	4024	4065
			4079	4193	4532	4547	4758
			4964	4967	5032	5175	5330
FokI	GGATGNNNNNNNNN!	4	3321	3778	3803	4257	
FokI	!NNNNNNNNNNNNNCATCC	9	655	693	888	1275	1460
			1548	1605	1760	3127	
FspI	TGC!GCA	1	3556				
GdiII	!NNNNNYGGCCG	6	631	790	2092	3354	3919
			3946				
GdiII	CGGCCRN!	3	643	2104	3366		
GsuI	!CTCCAG	3	848	1721	4251		
GsuI	!CTGGAG	1	4272				
HaeI	WGG!CCW	9	969	1199	1818	2049	3275
			3536	4898	5350	5361	
HaeII	RGCGC!Y	5	599	2721	2729	3457	5132
HaeIII	GG!CC	34	85	278	639	798	945
			969	1199	1235	1540	1783
			1818	1852	1875	1897	2007
			2049	2100	2420	2562	3217
			3223	3232	3275	3362	3536
			3927	3954	4463	4520	4553
			4898	5332	5350	5361	
HapII	C!CGG	23	602	905	942	974	1109
			1371	1963	2672	3359	3436
			3458	3486	3617	3707	3774
			3955	4230	4238	4330	4804
			4994	5020	5167		
HgaI	GACGCNNNNN!	3	527	4164	4243		
HgaI	!NNNNNNNNNNGCGTC	5	1164	1955	2787	4682	5260
HgiAI	GWGCW!C	5	621	1473	3567	3757	5062
HgiEII	!ACCNNNNNNGGT	1	4784				
HhaI	GCG!C	21	598	2720	2728	2754	2776
			2785	2798	2829	3448	3456
			3520	3557	3823	4083	4259
			4345	4748	4857	5031	5131
			5198				
Hin1I	GR!CGYC	8	123	176	259	445	1966
			3454	4156	4235		
HinP1I	G!CGC	21	596	2718	2726	2752	2774

			2783	2796	2827	3446	3454
			3518	3555	3821	4081	4257
			4343	4746	4855	5029	5129
			5196				
HincII	GTY↓RAC	2	722	2217			
HindIII	A↓AGCTT	1	623				
HinfI	G↓ANTC	16	411	605	1519	1669	1996
			2105	2497	2519	2929	3939
			4073	4125	4183	4220	5002
			5398				
HpaI	GTT↓AAC	1	2217				
HpaII	C↓CGG	23	602	905	942	974	1109
			1371	1963	2672	3359	3436
			3458	3486	3617	3707	3774
			3955	4230	4238	4330	4804
			4994	5020	5167		
HphI	GGTGANNNNNNNNN↓	5	377	1193	1775	4525	4641
HphI	↓NNNNNNNTCACC	3	1664	2571	3632		
KasI	G↓GCGCC	1	3453				
KpnI	GGTAC↓C	0					
Ksp632I	CTCTTCN↓	1	771				
Ksp632I	↓NNNNGAAGAG	5	1033	1393	2921	3798	4008
KspI	CCGC↓GG	0					
MaeI	C↓TAG	9	593	2068	2109	2303	2721
			3277	4292	4626	4879	
MaeII	A↓CGT	17	123	135	176	259	340
			445	964	1747	2023	2514
			2526	2569	2679	3570	3757
			4538	4671			
MaeIII	↓GTNAC	14	62	149	498	1188	1670
			2242	2743	2755	3574	3880
			4570	4836	4952	5015	
MamI	GATNN↓NNATC	2	2116	3313			
MboI	↓GATC	27	586	610	784	925	1042
			1267	1364	1753	1903	1929
			2111	3291	3295	3314	3625
			3703	3784	3793	3871	4247
			4263	4622	4634	4712	4720
			4731	4806			
MboII	GAAGANNNNNNNN↓	11	1050	1410	1695	1808	1928
			1996	2710	2938	3815	4025
			4644				
MboII	↓NNNNNNNTCTTC	12	758	764	1094	1097	1109
			1121	1157	1244	4105	4272
			4472	4715			
McrI	↓CGRYCG	5	636	1371	2097	3359	5034
MfeI	↓CAATTG	1	2203				
MflI	R↓GATCY	11	586	610	1364	1929	3625
			3871	4263	4622	4634	4720
			4731				
MluI	A↓CGCGT	0					
MmeI	↓TCCRAC	2	1775	2521			
MmeI	↓GTYGGA	2	4997	5181			
MnlI	CCTCNNNNNNN↓	20	659	720	774	798	957
			1552	1731	1746	1829	1962
			2073	2173	2182	3237	3243
			4055	4261	4575	4945	5269
MnlI	↓NNNNNNNGAGG	26	541	667	799	1432	1471
			1492	1525	1642	1751	1807

			1996	2029	2132	2597	2928
			3206	3212	3249	3252	3264
			3304	3368	3504	3861	4291
			5194				
MroI	T!CCGGA	0					
MscI	TGG!CCA	2	2049	3536			
MseI	T!TAA	19	8	1027	1408	1633	2089
			2155	2216	2337	2358	2369
			2381	2392	2409	2507	2778
			4597	4611	4616	4668	
MspI	C!CGG	23	602	905	942	974	1109
			1371	1963	2672	3359	3436
			3458	3486	3617	3707	3774
			3955	4230	4238	4330	4804
			4994	5020	5167		
MstI	TGC!GCA	1	3556				
MstII	CC!TNAGG	3	1648	2934	4566		
MunI	C!AATTG	1	2204				
MvaI	CC!WGG	20	91	284	856	951	1196
			1232	1566	1614	1710	1725
			1854	2990	3045	3062	3841
			4420	4523	5213	5226	5347
MvnI	CG!CG	18	61	636	801	1362	2103
			2378	2754	2774	2798	2829
			3520	3821	4259	4343	4406
			4475	4748	5329		
NaeI	GCC!GGC	4	943	2673	3956	4239	
NarI	GG!CGCC	1	3454				
NciI	CC!SGG	6	1110	1963	3458	3618	4231
			4995				
NcoI	C!CATGG	4	361	1381	3183	3886	
NdeI	CA!TATG	1	235				
NdeII	!GATC	27	586	610	784	925	1042
			1267	1364	1753	1903	1929
			2111	3291	3295	3314	3625
			3703	3784	3793	3871	4247
			4263	4622	4634	4712	4720
			4731	4806			
NgoMI	G!CCGGC	4	941	2671	3954	4237	
NheI	G!CTAGC	1	592				
NlaIII	CATG!	30	305	365	785	932	1004
			1136	1154	1172	1268	1385
			1421	1427	1439	1517	1583
			1663	1874	2884	3024	3096
			3187	3328	3673	3859	3890
			3916	4272	4656	5376	5427
NlaIV	GGN!NCC	20	468	1073	1074	1366	1445
			1505	1896	2609	2630	2642
			2833	2994	3066	3455	3490
			4338	4417	4462	5305	5344
NotI	GC!GGCCGC	2	637	2098			
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	3026	3098	5429		
Nsp(7524)I	RCATG!Y	5	1421	3024	3096	3859	5376
Nsp(7524)V	TT!CGAA	3	628	1120	4136		
NspBII	CMG!CKG	6	923	1842	2952	3560	4789
			5034				
NspII	GDGCH!C	15	621	696	917	1446	1473
			1506	2083	2645	3400	3493

			3567	3757	3819	4529	5062
NspIII	C!YCGRG	3	614	650	711		
NspIV	G!GNCC	17	84	277	944	1058	1072
			1084	1234	1539	1782	1874
			1895	2561	3970	4415	4461
			4519	4552			
NunII	GG!CGCC	1	3454				
PacI	TTAAT!TAA	0					
Paer71	C!TCGAG	1	614				
PalI	GG!CC	34	85	278	639	798	945
			969	1199	1235	1540	1783
			1818	1852	1875	1897	2007
			2049	2100	2420	2562	3217
			3223	3232	3275	3362	3536
			3927	3954	4463	4520	4553
			4898	5332	5350	5361	
PflMI	CCANNNN!NTGG	0					
PleI	GAGTCNNNN!	3	1677	2505	2937		
PleI	!NNNNNGACTC	6	405	599	2099	2513	4119
			4996				
PmaCI	CAC!GTG	1	965				
PmeI	GTTT!AAAC	0					
PmlI	CAC!GTG	1	965				
PpuMI	RG!GWCCY	1	1072				
PshAI	GACNN!NNGTC	1	2022				
PstI	CTGCA!G	1	922				
PvuI	CGAT!CG	0					
PvuII	CAG!CTG	3	923	2952	3560		
RmaI	C!TAG	9	593	2068	2109	2303	2721
			3277	4292	4626	4879	
RsaI	GT!AC	11	220	245	300	333	384
			541	1155	1434	1584	1836
			3760				
RsrI	G!AATTC	1	630				
RsrII	CG!GWCCG	1	3970				
SacI	GAGCT!C	1	621				
SacII	CCGC!GG	0					
SalI	G!TCGAC	0					
Sau3AI	!GATC	27	586	610	784	925	1042
			1267	1364	1753	1903	1929
			2111	3291	3295	3314	3625
			3703	3784	3793	3871	4247
			4263	4622	4634	4712	4720
			4731	4806			
Sau96I	G!GNCC	17	84	277	944	1058	1072
			1084	1234	1539	1782	1874
			1895	2561	3970	4415	4461
			4519	4552			
SauI	CC!TNAGG	3	1648	2934	4566		
ScaI	AGT!ACT	0					
ScrFI	CC!NGG	26	91	284	856	951	1110
			1196	1232	1566	1614	1710
			1725	1854	1963	2990	3045
			3062	3458	3618	3841	4231
			4420	4523	4995	5213	5226
			5347				
SduI	GDGCH!C	15	621	696	917	1446	1473
			1506	2083	2645	3400	3493
			3567	3757	3819	4529	5062

SecI	C!CNNGG	22	361	649	710	1381	1477
			1612	1613	1708	2988	3060
			3183	3218	3227	3276	3617
			3886	4291	4418	4419	4521
			4522	5212			
SexI	!CTCGAG	1	613				
SfaNI	GCATCNNNNN!	11	715	787	1108	1297	1627
			2270	3033	3105	3753	3817
			4092				
SfaNI	!NNNNNNNNNGATGC	8	358	1780	1819	1948	3412
			3667	3883	5275		
SfcI	C!TRYAG	5	699	918	2789	4916	5107
SfiI	GGCCNNNN!NGGCC	1	3229				
SfuI	TT!CGAA	3	628	1120	4136		
SgrAI	CR!CCGGYG	0					
SinI	G!GWCC	5	1058	1072	1084	3970	4415
SmaI	CCC!GGG	0					
SnaBI	TAC!GTA	1	341				
SnaI	!GTATAC	0					
SnoI	G!TGCAC	2	1469	5058			
SpeI	A!CTAGT	0					
SphI	GCATG!C	3	3024	3096	3859		
SplI	C!GTACG	0					
SrfI	GCCC!GGGC	0					
Sse8387I	CCTGCA!GG	0					
SspI	AAT!ATT	2	2362	2915			
SstI	GAGCT!C	1	621				
SstII	CCGC!GG	0					
SstIII	!ACGT	17	122	134	175	258	339
			444	963	1746	2022	2513
			2525	2568	2678	3569	3756
			4537	4670			
StuI	AGG!CCT	2	1818	3275			
StyI	C!CWWGG	6	361	1381	3183	3276	3886
			4291				
StySJI	!GAGNNNNNGTRC	0					
StySJI	!GYACNNNNNNCTC	0					
SwaI	ATTT!AAAT	0					
TaqI	T!CGA	19	615	628	1036	1066	1120
			1531	1558	2026	2603	3294
			3567	3723	3747	3783	3945
			4136	4181	4436	5274	
TaqII	GACCGANNNNNNNNNN!	3	2474	3491	4159		
TaqII	!NNNNNNNNNTCGGTC	1	4403				
TaqII	CACCCANNNNNNNNNN!	2	1521	1626			
TaqII	!NNNNNNNNNTGGGTG	3	173	3358	3950		
TfiI	G!AWTC	7	1519	1996	3939	4073	4183
			4220	5398			
ThaI	CG!CG	18	61	636	801	1362	2103
			2378	2754	2774	2798	2829
			3520	3821	4259	4343	4406
			4475	4748	5329		
Tsp45I	!GTSAC	4	1670	2743	3574	3880	
TspEI	!AATT	13	19	630	1282	2090	2204
			2268	2346	2372	2383	3030
			3102	3194	4612		
Tth111I	GACN!NNGTC	1	3572				
Tth111II	CAARCANNNNNNNNNN!	2	741	4781			
Tth111II	!NNNNNNNNNTGYTTG	4	826	3889	4749	4788	

TthHB8I	T!CGA	19	615	628	1036	1066	1120
			1531	1558	2026	2603	3294
			3567	3723	3747	3783	3945
			4136	4181	4436	5274	
Van91I	CCANNNN!NTGG	0					
VspI	AT!TAAT	1	8				
XbaI	T!CTAGA	1	2108				
XcmI	!CCANNNNNNNNNTCG	2	1347	3729			
XcyI	C!CCGGG	0					
XhoI	C!TCGAG	1	614				
XhoII	R!GATCY	11	586	610	1364	1929	3625
			3871	4263	4622	4634	4720
			4731				
XmaI	C!CCGGG	0					
XmaIII	C!GGCCG	3	637	2098	3360		
XmnI	GAANN!NNTTC	0					
XorII	CGAT!CG	0					