

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	2	1040	3183			
AatII	GACGT!C	4	126	179	262	448	
Acc65I	G!GTACC	0					
AccI	GT!MKAC	2	1191	1356			
AccII	CG!CG	15	61	2250	2286	2662	2682
			2706	2737	3428	3729	4167
			4251	4314	4383	4656	5237
AccIII	T!CCGGA	1	1316				
AciI	G!CGG	32	371	404	529	749	759
			1071	1113	1412	1415	1430
			1514	1744	2622	2649	2737
			2847	3428	3492	3593	3596
			3876	3931	4029	4167	4314
			4454	4697	4942	5063	5082
			5209	5237			
AciI	C!CGC	39	59	87	99	113	280
			508	590	924	1062	1164
			1434	1521	1669	1945	1990
			2677	2680	2694	2784	3032
			3044	3053	3065	3075	3086
			3132	3271	3334	3836	3881
			3947	3973	4098	4101	4249
			4381	4688	4832	5328	
AcyI	GR!CGYC	9	123	176	259	445	1188
			1979	3362	4064	4143	
AfaI	GT!AC	10	220	245	300	333	384
			541	656	806	1058	3668
AflI	G!GWCC	6	1423	1566	1604	1762	3878
			4323				
AflII	C!TTAAG	0					
AflIII	A!CRYGT	2	2248	5280			
AgeI	A!CCGGT	1	601				
AhaI	CC!SGG	9	1185	1406	1427	1530	1823
			3366	3526	4139	4903	
AhaII	GR!CGYC	9	123	176	259	445	1188
			1979	3362	4064	4143	
AhaIII	TTT!AAA	3	2064	4506	4525		
AluI	AG!CT	20	567	625	652	798	1105
			1334	1340	1708	1897	2141
			2308	2565	2860	3148	3468
			3926	4723	4980	5026	5116
Alw44I	G!TGCAC	2	691	4966			
AlwI	GGATCNNNN!	9	2005	2014	3230	3541	3787
			4179	4538	4636	4722	
AlwI	!NNNNNGATCC	7	581	1146	1992	3606	4150
			4537	4634			
AlwNI	CAGNNN!CTG	3	1278	4463	4871		
AocI	CC!TNAGG	3	870	2842	4474		
AocII	GDGCH!C	17	668	695	728	1305	1336
			1405	1498	1535	1962	2553
			3308	3401	3475	3665	3727
			4437	4970			
Aor51HI	AGC!GCT	1	597				
AosI	TGC!GCA	1	3464				
AosII	GR!CGYC	9	123	176	259	445	1188

			1979	3362	4064	4143	
ApaI	GGGCC!C	1	1535				
ApaLI	G!TGCAC	2	691	4966			
ApoI	R!AATTY	4	1345	2176	2280	2291	
ApyI	CC!WGG	21	91	284	788	836	932
			947	1076	1442	1634	1652
			1766	1869	2898	2953	2970
			3749	4328	4431	5121	5134
			5255				
AquI	C!YCGRG	1	1329				
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	6	695	1336	1962	3475	3665
			4970				
AspI	GACN!NNGTC	1	3480				
AsuI	G!GNCC	21	84	277	761	1004	1096
			1117	1423	1444	1516	1531
			1532	1566	1604	1762	1891
			2469	3878	4323	4369	4427
			4460				
AsuII	TT!CGAA	2	1343	4044			
AvaI	C!YCGRG	1	1329				
AvaII	G!GWCC	6	1423	1566	1604	1762	3878
			4323				
AvaIII	ATGCA!T	3	2934	3006	5337		
AviII	TGC!GCA	1	3464				
AvrI	!CYCGRG	1	1328				
AvrII	C!CTAGG	2	3184	4199			
AxyI	CC!TNAGG	3	870	2842	4474		
BalI	TGG!CCA	2	1271	3444			
BamHI	G!GATCC	1	1997				
BanI	G!GYRCC	8	466	665	725	1400	1495
			2515	3361	3396		
BanII	GRGCY!C	5	1336	1535	2553	3727	4437
BanIII	AT!CGAT	1	3202				
BbeI	GGCGC!C	1	3365				
BbiIII/AcyI	GR!CGYC	9	123	176	259	445	1188
			1979	3362	4064	4143	
BbrPI	CAC!GTG	0					
BbsI	!GTCTTC	0					
BbsI	!GAAGAC	3	904	1137	1205		
BbvI	GCAGCNNNNNNN!	11	822	1264	1518	1634	1932
			2150	3435	3477	3998	4451
			4951				
BbvI	!NNNNNNNNNNNGCTGC	10	941	1686	1702	2644	3309
			3493	3586	4652	4858	4861
BbvII	GAAGACNN!	3	912	1145	1213		
BbvII	!NNNNNNGTCTTC	0					
BcefI	ACGGCNNNNNNNNNN!	7	823	967	1018	1084	1712
			3788	4275			
BcefI	!NNNNNNNNNNNGCCGT	3	2504	3322	4781		
BcgI	!GCANNNNNNTCG	0					
BcgI	!CGANNNNNNTGC	0					
BclI	T!GATCA	1	2019				
BcnI	CC!SGG	9	1185	1406	1427	1530	1823
			3366	3526	4139	4903	

BfaI	C!TAG	10	593 2629	1290 3185	1993 4200	2010 4534	2211 4787
BfrI	C!TTAAG	0					
BglI	GCCNNNN!NGGC	8	91 1409	213 1442	284 3137	1080	1094
BglII	A!GATCT	1	1325				
BinI	!NNNNNGATCC	7	581 4537	1146 4634	1992	3606	4150
BinI	GGATCNNNN!	9	2005 4179	2014 4538	3230 4636	3541 4722	3787
BlnI	C!CTAGG	2	3184	4199			
BmyI	GDGCH!C	17	668 1405 3308 4437	695 1498 3401 4970	728 1535 3475	1305 1962 3665	1336 2553 3727
BpmI	!CTCCAG	2	943	4159			
BpmI	!CTGGAG	2	1765	4180			
Bpu1102I	GC!TNAGC	0					
BsaAI	YAC!GTR	3	341	2478	3666		
BsaBI	GATNN!NNATC	2	2024	3221			
BsaHI	GR!CGYC	9	123 1979	176 3362	259 4064	445 4143	1188
BsaI	GGTCTCN!	0					
BsaI	!NNNNNGAGACC	4	1043	1232	1856	4351	
BsaJI	C!CNNGG	26	361 930 1854 3126 4199 5120	612 1404 1867 3135 4326	699 1632 2896 3184 4327	834 1650 2968 3525 4429	835 1830 3091 3794 4430
BseAI	T!CCGGA	1	1316				
BsePI	!GCGCGC	0					
BsgI	!GTGCAG	0					
BsgI	!CTGCAC	0					
BsiEI	CGRY!CG	4	753	1434	3271	4946	
BsiWI	C!GTACG	0					
BsiYI	CCNNNNN!NNGG	27	50 1411 2374 4168 4417 5247	722 1541 2700 4200 4418 5265	835 1542 3092 4201 4449	1001 1613 3343 4332 4802	1067 1954 3887 4367 5081
BslI	CCNNNNN!NNGG	27	50 1411 2374 4168 4417 5247	722 1541 2700 4200 4418 5265	835 1542 3092 4201 4449	1001 1613 3343 4332 4802	1067 1954 3887 4367 5081
BsmAI	!GTCTC	1	429				
BsmAI	!GAGAC	7	1048 4228	1237 4356	1861	2791	3209
BsmI	GAATGCN!	1	2113				
BsmI	G!CATTC	1	2206				
Bsp1286I	GDGCH!C	17	668 1405 3308 4437	695 1498 3401 4970	728 1535 3475	1305 1962 3665	1336 2553 3727
BspDI	AT!CGAT	1	3202				
BspEI	T!CCGGA	1	1316				

BspHI	T!CATGA	2	2788	4560			
BspMI	ACCTGCNNNN!	3	1507	3630	4080		
BspMI	!NNNNNNNNGCAGGT	1	3249				
BspMII	T!CCGGA	1	1316				
BspWI	GCNNNNN!NNGC	39	91	213	245	284	377
			401	649	1080	1094	1409
			1442	1520	1628	1684	1705
			1929	1957	2589	2619	2646
			2688	2690	2929	3001	3052
			3131	3137	3353	3437	3460
			3599	3605	3722	3805	4072
			4259	4463	4662	5234	
BsrFI	R!CCGGY	6	601	1502	2579	3681	3862
			4145				
BsrI	ACTGGN!	5	1215	1302	3306	3507	4753
BsrI	C!CAGT	5	296	2391	3069	4865	4878
BssHII	G!CGCGC	0					
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	2	1343	4044			
BstEII	G!GTNACC	0					
BstI	G!GATCC	1	1997				
BstNI	CC!WGG	21	91	284	788	836	932
			947	1076	1442	1634	1652
			1766	1869	2898	2953	2970
			3749	4328	4431	5121	5134
			5255				
BstPI	G!GTNACC	0					
BstUI	CG!CG	15	61	2250	2286	2662	2682
			2706	2737	3428	3729	4167
			4251	4314	4383	4656	5237
BstXI	CCANNNNN!NTGG	0					
BstYI	R!GATCY	12	586	1151	1325	1997	2006
			3533	3779	4171	4530	4542
			4628	4639			
Bsu36I	CC!TNAGG	3	870	2842	4474		
CbiI	TT!CGAA	2	1343	4044			
CcrI	C!TCGAG	1	1329				
CelII	GC!TNAGC	0					
CfoI	GCG!C	23	598	1785	1953	2628	2636
			2662	2684	2693	2706	2737
			3356	3364	3428	3465	3731
			3991	4167	4253	4656	4765
			4939	5039	5106		
Cfr10I	R!CCGGY	6	601	1502	2579	3681	3862
			4145				
Cfr13I	G!GNCC	21	84	277	761	1004	1096
			1117	1423	1444	1516	1531
			1532	1566	1604	1762	1891
			2469	3878	4323	4369	4427
			4460				
Cfr9I	C!CCGGG	0					
CfrI	Y!GGCCR	6	1269	1431	3268	3442	3833
			3860				
ClaI	AT!CGAT	1	3202				
CpoI	CG!GWCCG	1	3878				
Csp45I	TT!CGAA	2	1343	4044			
CspI	CG!GSCCG	0					
CviJI	RG!CY	116	43	85	94	278	567
			608	625	652	715	762

			791	798	876	942	954
			1005	1032	1040	1074	1088
			1097	1105	1119	1229	1265
			1271	1334	1340	1364	1377
			1433	1445	1488	1509	1517
			1528	1533	1547	1598	1617
			1625	1637	1645	1657	1699
			1708	1754	1838	1892	1897
			2031	2141	2308	2328	2470
			2551	2565	2579	2860	2901
			2973	3096	3125	3131	3140
			3148	3171	3183	3189	3270
			3287	3295	3322	3347	3431
			3440	3444	3468	3506	3582
			3599	3618	3681	3725	3735
			3752	3835	3862	3866	3903
			3926	3942	4122	4149	4210
			4332	4371	4428	4435	4442
			4461	4472	4723	4752	4795
			4806	4871	4950	4975	4980
			5026	5116	5214	5240	5258
			5269				
CvnI	CC!TNAGG	3	870	2842	4474		
DdeI	C!TNAG	12	870	1167	1322	1554	1659
			1722	2842	3144	4025	4474
			4597	5006			
DpnI	GA!TC	25	588	977	1127	1153	1327
			1814	1999	2008	2021	3201
			3205	3224	3535	3613	3694
			3703	3781	4157	4173	4532
			4544	4622	4630	4641	4716
DpnII	!GATC	25	586	975	1125	1151	1325
			1812	1997	2006	2019	3199
			3203	3222	3533	3611	3692
			3701	3779	4155	4171	4530
			4542	4620	4628	4639	4714
DraI	TTT!AAA	3	2064	4506	4525		
DraII	RG!GNCCY	4	1423	1532	1762	4460	
DraIII	CACNNN!GTG	3	1261	1878	2478		
DrdI	GACNNNN!NNGTC	4	1984	2433	3389	5178	
DsaI	C!CRYGG	4	361	612	3091	3794	
DsaV	!CCNGG	30	89	282	786	834	930
			945	1074	1183	1404	1425
			1440	1528	1632	1650	1764
			1821	1867	2896	2951	2968
			3364	3524	3747	4137	4326
			4429	4901	5119	5132	5253
EaeI	Y!GGCCR	6	1269	1431	3268	3442	3833
			3860				
EagI	C!GGCCG	2	1431	3268			
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	3	2833	3710	3920		
Ecl136II	GAG!CTC	1	1334				
EclXI	C!GGCCG	2	1431	3268			
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	4	1043	1232	1856	4351	
Eco47I	G!GWCC	6	1423	1566	1604	1762	3878

			4323					
Eco47III	AGC!GCT	1	597					
Eco52I	C!GGCCG	2	1431	3268				
Eco57I	!CTGAAG	4	1099	1969	3486	3918		
Eco57I	!CTTCAG	1	4752					
Eco81I	CC!TNAGG	3	870	2842	4474			
EcoNI	CCTNN!NNNAGG	0						
EcoO109I	RG!GNCCY	4	1423	1532	1762	4460		
EcoO65I	G!GTNACC	0						
EcoRI	G!AATTC	1	1345					
EcoRII	!CCWGG	21	89	282	786	834	930	
			945	1074	1440	1632	1650	
			1764	1867	2896	2951	2968	
			3747	4326	4429	5119	5132	
			5253					
EcoRV	GAT!ATC	0						
EcoT14I	C!CWWGG	7	361	612	1854	3091	3184	
			3794	4199				
EcoT22I	ATGCA!T	3	2934	3006	5337			
EcoT38I	GRGCY!C	5	1336	1535	2553	3727	4437	
EheI	GGC!GCC	1	3363					
Esp3I	CGTCTCN!	0						
Esp3I	!NNNNNGAGACG	0						
EspI	GC!TNAGC	0						
FbaI	T!GATCA	1	2019					
FinI	!GTCCC	7	1941	1965	1986	2892	2964	
			3028	4323				
FinI	!GGGAC	7	161	312	480	1564	3498	
			4030	4139				
Fnu4HI	GC!NGC	42	760	811	955	1072	1253	
			1413	1431	1434	1507	1515	
			1623	1700	1716	1921	2139	
			2658	2680	2694	3132	3271	
			3323	3334	3424	3429	3466	
			3507	3594	3597	3600	3836	
			3932	3973	3987	4101	4440	
			4455	4666	4872	4875	4940	
			5083	5238				
FokI	GGATGNNNNNNNN!	5	1724	3229	3686	3711	4165	
FokI	!NNNNNNNNNNNNNCATCC	7	682	770	827	982	1652	
			1813	3035				
FspI	TGC!GCA	1	3464					
GdiII	!NNNNNYGGCCG	4	1425	3262	3827	3854		
GdiII	CGGCCRN!	2	1437	3274				
GsuI	!CTCCAG	2	943	4159				
GsuI	!CTGGAG	2	1765	4180				
HaeI	WGG!CCW	9	1040	1271	1598	1754	3183	
			3444	4806	5258	5269		
HaeII	RGCGC!Y	6	599	1786	2629	2637	3365	
			5040					
HaeIII	GG!CC	34	85	278	762	1005	1040	
			1074	1097	1119	1229	1271	
			1433	1445	1517	1533	1598	
			1754	1892	2328	2470	3125	
			3131	3140	3183	3270	3444	
			3835	3862	4371	4428	4461	
			4806	5240	5258	5269		
HapII	C!CGG	25	602	1185	1317	1406	1427	
			1503	1529	1822	2004	2580	

			3267	3344	3366	3394	3525
			3615	3682	3863	4138	4146
			4238	4712	4902	4928	5075
HgaI	GACGCNNNNN!	4	527	1987	4072	4151	
HgaI	!NNNNNNNNNNGCGTC	4	1177	2695	4590	5168	
HgiAI	GWGCW!C	6	695	1336	1962	3475	3665
			4970				
HgiEII	!ACCNNNNNNNGGT	1	4692				
HhaI	GCG!C	23	598	1785	1953	2628	2636
			2662	2684	2693	2706	2737
			3356	3364	3428	3465	3731
			3991	4167	4253	4656	4765
			4939	5039	5106		
Hin1I	GR!CGYC	9	123	176	259	445	1188
			1979	3362	4064	4143	
HinP1I	G!CGC	23	596	1783	1951	2626	2634
			2660	2682	2691	2704	2735
			3354	3362	3426	3463	3729
			3989	4165	4251	4654	4763
			4937	5037	5104		
HincII	GTY!RAC	2	1357	2125			
HindIII	A!AGCTT	1	1338				
HinfI	G!ANTC	16	411	741	891	1218	1320
			1850	2405	2427	2837	3847
			3981	4033	4091	4128	4910
			5306				
HpaI	GTT!AAC	1	2125				
HpaII	C!CGG	25	602	1185	1317	1406	1427
			1503	1529	1822	2004	2580
			3267	3344	3366	3394	3525
			3615	3682	3863	4138	4146
			4238	4712	4902	4928	5075
HphI	GGTGANNNNNNNN!	5	377	628	997	4433	4549
HphI	!NNNNNNNTCACC	3	886	2479	3540		
KasI	G!GCGCC	1	3361				
KpnI	GGTAC!C	0					
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	3	2829	3706	3916		
KspI	CCGC!GG	0					
MaeI	C!TAG	10	593	1290	1993	2010	2211
			2629	3185	4200	4534	4787
MaeII	A!CGT	16	123	135	176	259	340
			445	969	1245	2422	2434
			2477	2587	3478	3665	4446
			4579				
MaeIII	!GTNAC	15	62	149	498	892	1726
			1901	2150	2651	2663	3482
			3788	4478	4744	4860	4923
MamI	GATNN!NNATC	2	2024	3221			
MboI	!GATC	25	586	975	1125	1151	1325
			1812	1997	2006	2019	3199
			3203	3222	3533	3611	3692
			3701	3779	4155	4171	4530
			4542	4620	4628	4639	4714
MboII	GAAGANNNNNNNN!	10	917	1030	1150	1218	1816
			2618	2846	3723	3933	4552
MboII	!NNNNNNNTCTTC	5	1372	4013	4180	4380	4623
McrI	!CGRYCG	4	749	1430	3267	4942	
MfeI	!CAATTG	1	2111				

MflI	R!GATCY	12	586 3533 4628	1151 3779 4639	1325 4171	1997 4530	2006 4542
MluI	A!CGCGT	1	2248				
MmeI	!TCCRAC	2	997	2429			
MmeI	!GTYGGA	3	1389	4905	5089		
MnlI	CCTCNNNNNNN!	15	774 1295 3963	953 2081 4169	968 2090 4483	1051 3145 4853	1184 3151 5177
MnlI	!NNNNNNNGAGG	36	541 864 1401 1587 1827 3114 3212 5102	654 973 1410 1674 1851 3120 3276	693 1029 1413 1682 2040 3157 3412	714 1218 1476 1761 2505 3160 3769	747 1251 1548 1824 2836 3172 4199
MroI	T!CCGGA	1	1316				
MscI	TGG!CCA	2	1271	3444			
MseI	T!TAA	18	8 2124 2300 4519	630 2245 2317 4524	855 2266 2415 4576	1311 2277 2686	2063 2289 4505
MspI	C!CGG	25	602 1503 3267 3615 4238	1185 1529 3344 3682 4712	1317 1822 3366 3863 4902	1406 2004 3394 4138 4928	1427 2580 3525 4146 5075
MstI	TGC!GCA	1	3464				
MstII	CC!TNAGG	3	870	2842	4474		
MunI	C!AATTG	1	2112				
MvaI	CC!WGG	21	91 947 1766 3749 5255	284 1076 1869 4328	788 1442 2898 4431	836 1634 2953 5121	932 1652 2970 5134
MvnI	CG!CG	15	61 2706 4251	2250 2737 4314	2286 3428 4383	2662 3729 4656	2682 4167 5237
NaeI	GCC!GGC	4	1504	2581	3864	4147	
NarI	GG!CGCC	1	3362				
NciI	CC!SGG	9	1185 3366	1406 3526	1427 4139	1530 4903	1823
NcoI	C!CATGG	4	361	612	3091	3794	
NdeI	CA!TATG	1	235				
NdeII	!GATC	25	586 1812 3203 3701 4542	975 1997 3222 3779 4620	1125 2006 3533 4155 4628	1151 2019 3611 4171 4639	1325 3199 3692 4530 4714
NgoMI	G!CCGGC	4	1502	2579	3862	4145	
NheI	G!CTAGC	1	592				
NlaIII	CATG!	27	305 661 1363 2932 3767 5284	365 739 1514 3004 3798 5335	616 805 1564 3095 3824	643 885 1966 3236 4180	649 1096 2792 3581 4564
NlaIV	GGN!NCC	27	468	667	727	1118	1402

			1425	1497	1518	1533	1534
			1567	1656	1778	1999	2517
			2538	2550	2741	2902	2974
			3363	3398	4246	4325	4370
			5213	5252			
NotI	GC!GGCCGC	1	1431				
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	2934	3006	5337		
Nsp(7524)I	RCATG!Y	5	643	2932	3004	3767	5284
Nsp(7524)V	TT!CGAA	2	1343	4044			
NspBII	CMG!CKG	5	1064	2860	3468	4697	4942
NspII	GDGCH!C	17	668	695	728	1305	1336
			1405	1498	1535	1962	2553
			3308	3401	3475	3665	3727
			4437	4970			
NspIII	C!YCGRG	1	1329				
NspIV	G!GNCC	21	84	277	761	1004	1096
			1117	1423	1444	1516	1531
			1532	1566	1604	1762	1891
			2469	3878	4323	4369	4427
			4460				
NunII	GG!CGCC	1	3362				
PacI	TTAAT!TAA	0					
Paer71	C!TCGAG	1	1329				
PalI	GG!CC	34	85	278	762	1005	1040
			1074	1097	1119	1229	1271
			1433	1445	1517	1533	1598
			1754	1892	2328	2470	3125
			3131	3140	3183	3270	3444
			3835	3862	4371	4428	4461
			4806	5240	5258	5269	
PflMI	CCANNNN!NTGG	0					
PleI	GAGTCNNNN!	3	899	2413	2845		
PleI	!NNNNNGACTC	5	405	1314	2421	4027	4904
PmaCI	CAC!GTG	0					
PmeI	GTTT!AAAC	0					
PmlI	CAC!GTG	0					
PpuMI	RG!GWCCY	2	1423	1762			
PshAI	GACNN!NNGTC	2	1244	1571			
PstI	CTGCA!G	1	1354				
PvuI	CGAT!CG	0					
PvuII	CAG!CTG	2	2860	3468			
RmaI	C!TAG	10	593	1290	1993	2010	2211
			2629	3185	4200	4534	4787
RsaI	GT!AC	10	220	245	300	333	384
			541	656	806	1058	3668
RsrI	G!AATTC	1	1345				
RsrII	CG!GWCCG	1	3878				
SacI	GAGCT!C	1	1336				
SacII	CCGC!GG	0					
SalI	G!TCGAC	1	1355				
Sau3AI	!GATC	25	586	975	1125	1151	1325
			1812	1997	2006	2019	3199
			3203	3222	3533	3611	3692
			3701	3779	4155	4171	4530
			4542	4620	4628	4639	4714
Sau96I	G!GNCC	21	84	277	761	1004	1096
			1117	1423	1444	1516	1531
			1532	1566	1604	1762	1891

			2469	3878	4323	4369	4427
			4460				
SauI	CC!TNAGG	3	870	2842	4474		
ScaI	AGT!ACT	0					
ScrFI	CC!NGG	30	91	284	788	836	932
			947	1076	1185	1406	1427
			1442	1530	1634	1652	1766
			1823	1869	2898	2953	2970
			3366	3526	3749	4139	4328
			4431	4903	5121	5134	5255
SduI	GDGCH!C	17	668	695	728	1305	1336
			1405	1498	1535	1962	2553
			3308	3401	3475	3665	3727
			4437	4970			
SecI	C!CNNGG	26	361	612	699	834	835
			930	1404	1632	1650	1830
			1854	1867	2896	2968	3091
			3126	3135	3184	3525	3794
			4199	4326	4327	4429	4430
			5120				
SexI	!CTCGAG	1	1328				
SfaNI	GCATCNNNNN!	10	849	1674	1727	1835	2178
			2941	3013	3661	3725	4000
SfaNI	!NNNNNNNNNGATGC	8	358	1002	1170	1702	3320
			3575	3791	5183		
SfcI	C!TRYAG	4	1350	2697	4824	5015	
SfiI	GGCCNNNN!NGGCC	1	3137				
SfuI	TT!CGAA	2	1343	4044			
SgrAI	CR!CCGGYG	0					
SinI	G!GWCC	6	1423	1566	1604	1762	3878
			4323				
SmaI	CCC!GGG	0					
SnaBI	TAC!GTA	1	341				
SnaI	!GTATAC	0					
SnoI	G!TGCAC	2	691	4966			
SpeI	A!CTAGT	0					
SphI	GCATG!C	3	2932	3004	3767		
SplI	C!GTACG	0					
SrfI	GCCC!GGGC	0					
Sse8387I	CCTGCA!GG	0					
SspI	AAT!ATT	2	2270	2823			
SstI	GAGCT!C	1	1336				
SstII	CCGC!GG	0					
SstIII	!ACGT	16	122	134	175	258	339
			444	968	1244	2421	2433
			2476	2586	3477	3664	4445
			4578				
StuI	AGG!CCT	2	1040	3183			
StyI	C!CWWGG	7	361	612	1854	3091	3184
			3794	4199			
StySJI	!GAGNNNNNGTRC	0					
StySJI	!GYACNNNNNNCTC	0					
SwaI	ATTT!AAAT	0					
TaqI	T!CGA	17	753	780	1248	1330	1343
			1356	2511	3202	3475	3631
			3655	3691	3853	4044	4089
			4344	5182			
TaqII	GACCGANNNNNNNNNN!	3	2382	3399	4067		
TaqII	!NNNNNNNNNTCGGTC	1	4311				

TaqII	CACCCANNNNNNNNNNN!	2	743	848			
TaqII	!NNNNNNNNNTGGGTG	3	173	3266	3858		
TfiI	G!AWTC	8	741	1218	1850	3847	3981
			4091	4128	5306		
ThaI	CG!CG	15	61	2250	2286	2662	2682
			2706	2737	3428	3729	4167
			4251	4314	4383	4656	5237
Tsp45I	!GTSAC	6	892	1726	1901	2651	3482
			3788				
TspEI	!AATT	12	19	1312	1345	2112	2176
			2254	2280	2291	2938	3010
			3102	4520			
Tth111I	GACN!NNGTC	1	3480				
Tth111II	CAARCANNNNNNNNNN!	2	1940	4689			
Tth111III	!NNNNNNNNNTGYTTG	3	3797	4657	4696		
TthHB8I	T!CGA	17	753	780	1248	1330	1343
			1356	2511	3202	3475	3631
			3655	3691	3853	4044	4089
			4344	5182			
Van91I	CCANNNN!NTGG	0					
VspI	AT!TAAT	1	8				
XbaI	T!CTAGA	1	2009				
XcmI	!CCANNNNNNNNTCG	1	3637				
XcyI	C!CCGGG	0					
XhoI	C!TCGAG	1	1329				
XhoII	R!GATCY	12	586	1151	1325	1997	2006
			3533	3779	4171	4530	4542
			4628	4639			
XmaI	C!CCGGG	0					
XmaIII	C!GGCCG	2	1431	3268			
XmnI	GAANN!NNTTC	0					
XorII	CGAT!CG	0					