

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	2	1489	2946			
AatII	GACGT!C	4	126	179	262	448	
Acc65I	G!GTACC	1	646				
AccI	GT!MKAC	2	641	1640			
AccII	CG!CG	15	61	1774	2049	2425	2445
			2469	2500	3191	3492	3930
			4014	4077	4146	4419	5000
AccIII	T!CCGGA	0					
AciI	G!CGG	29	371	404	529	709	734
			1208	1520	1562	1768	2385
			2412	2500	2610	3191	3255
			3356	3359	3639	3694	3792
			3930	4077	4217	4460	4705
			4826	4845	4972	5000	
AciI	C!CGC	39	59	87	99	113	280
			508	590	650	683	756
			894	1373	1511	1613	1772
			2440	2443	2457	2547	2795
			2807	2816	2828	2838	2849
			2895	3034	3097	3599	3644
			3710	3736	3861	3864	4012
			4144	4451	4595	5091	
AcyI	GR!CGYC	8	123	176	259	445	1637
			3125	3827	3906		
AfaI	GT!AC	14	220	245	300	333	384
			541	648	784	1002	1021
			1105	1255	1507	3431	
AflI	G!GWCC	3	801	3641	4086		
AflIII	C!TTAAG	1	2007				
AflIII	A!CRYGT	1	5043				
AgeI	A!CCGGT	1	1041				
AhaI	CC!SGG	5	1634	3129	3289	3902	4666
AhaII	GR!CGYC	8	123	176	259	445	1637
			3125	3827	3906		
AhaIII	TTT!AAA	3	1827	4269	4288		
AluI	AG!CT	19	567	619	625	767	1074
			1101	1247	1554	1904	2071
			2328	2623	2911	3231	3689
			4486	4743	4789	4879	
Alw44I	G!TGCAC	2	1140	4729			
AlwI	GGATCNNNN!	8	1043	2993	3304	3550	3942
			4301	4399	4485		
AlwI	!NNNNNGATCC	8	581	933	1030	1595	3369
			3913	4300	4397		
AlwNI	CAGNNN!CTG	3	1727	4226	4634		
AocI	CC!TNAGG	3	1319	2605	4237		
AocII	GDGCH!C	14	621	700	1117	1144	1177
			1754	2316	3071	3164	3238
			3428	3490	4200	4733	
Aor51HI	AGC!GCT	3	597	916	1029		
AosI	TGC!GCA	1	3227				
AosII	GR!CGYC	8	123	176	259	445	1637
			3125	3827	3906		
ApaI	GGGCC!C	0					
ApaLI	G!TGCAC	2	1140	4729			
ApoI	R!AATY	5	630	846	1939	2043	2054

ApyI	CC!WGG	19	91 1237 2661 4194	284 1285 2716 4884	898 1381 2733 4897	933 1396 3512 5018	968 1525 4091
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	1	646				
AspHI	GWGCW!C	5	621	1144	3238	3428	4733
AspI	GACN!NNGTC	1	3243				
AsuI	G!GNCC	13	84 1545 4132	277 1566 4190	801 2232 4223	1210 3641	1453 4086
AsuII	TT!CGAA	2	628	3807			
AvaI	C!YCGRG	1	614				
AvaII	G!GWCC	3	801	3641	4086		
AvaIII	ATGCA!T	3	2697	2769	5100		
AviII	TGC!GCA	1	3227				
AvrI	!CYCGRG	1	613				
AvrII	C!CTAGG	2	2947	3962			
AxyI	CC!TNAGG	3	1319	2605	4237		
BalI	TGG!CCA	3	979	1720	3207		
BamHI	G!GATCC	1	1035				
BanI	G!GYRCC	7	466 3124	646 3159	1114	1174	2278
BanII	GRGCY!C	5	621	700	2316	3490	4200
BanIII	AT!CGAT	1	2965				
BbeI	GGCGC!C	1	3128				
BbiII/AcyI	GR!CGYC	8	123 3125	176 3827	259 3906	445	1637
BbrPI	CAC!GTG	0					
BbsI	!GTCTTC	0					
BbsI	!GAAGAC	3	1353	1586	1654		
BbvI	GCAGCNNNNNNNN!	9	764 3240	1271 3761	1713 4214	1913 4714	3198
BbvI	!NNNNNNNNNNNGCTGC	10	947 3256	950 3349	1390 4415	2407 4621	3072 4624
BbvII	GAAGACNN!	3	1361	1594	1662		
BbvII	!NNNNNNGTCTTC	0					
BcefI	ACGGCNNNNNNNNNN!	7	962 3551	1272 4038	1416	1467	1533
BcefI	!NNNNNNNNNNNGCCGT	5	935	974	2267	3085	4544
BcgI	!GCANNNNNNTCG	0					
BcgI	!CGANNNNNNTGC	0					
BclI	T!GATCA	0					
BcnI	CC!SGG	5	1634	3129	3289	3902	4666
BfaI	C!TAG	9	593 2948	1739 3963	1780 4297	1974 4550	2392
BfrI	C!TTAAG	1	2007				
BglI	GCCNNNN!NGGC	5	91	213	284	1529	2900
BglII	A!GATCT	1	610				
BinI	!NNNNNGATCC	8	581 3913	933 4300	1030 4397	1595	3369
BinI	GGATCNNNN!	8	1043 4301	2993 4399	3304 4485	3550	3942
BlnI	C!CTAGG	2	2947	3962			
BmyI	GDGCH!C	14	621	700	1117	1144	1177

			1754	2316	3071	3164	3238
			3428	3490	4200	4733	
BpmI	!CTCCAG	3	929	1392	3922		
BpmI	!CTGGAG	1	3943				
Bpu1102I	GC!TNAGC	0					
BsaAI	YAC!GTR	3	341	2241	3429		
BsaBI	GATNN!NNATC	2	1787	2984			
BsaHI	GR!CGYC	8	123	176	259	445	1637
			3125	3827	3906		
BsaI	GGTCTCN!	0					
BsaI	!NNNNNGAGACC	2	1681	4114			
BsaJI	C!CNNGG	25	361	827	833	932	967
			995	1052	1148	1283	1284
			1379	2659	2731	2854	2889
			2898	2947	3288	3557	3962
			4089	4090	4192	4193	4883
BseAI	T!CCGGA	0					
BsePI	!GCGCGC	0					
BsgI	!GTGCAG	0					
BsgI	!CTGCAC	0					
BsiEI	CGRY!CG	5	950	1046	1772	3034	4709
BsiWI	C!GTACG	0					
BsiYI	CCNNNNN!NNGG	25	50	693	694	814	1171
			1284	1450	1516	2137	2463
			2855	3106	3650	3931	3963
			3964	4095	4130	4180	4181
			4212	4565	4844	5010	5028
BsII	CCNNNNN!NNGG	25	50	693	694	814	1171
			1284	1450	1516	2137	2463
			2855	3106	3650	3931	3963
			3964	4095	4130	4180	4181
			4212	4565	4844	5010	5028
BsmAI	!GTCTC	1	429				
BsmAI	!GAGAC	5	1686	2554	2972	3991	4119
BsmI	GAATGCN!	1	1876				
BsmI	G!CATTC	1	1969				
Bsp1286I	GDGCH!C	14	621	700	1117	1144	1177
			1754	2316	3071	3164	3238
			3428	3490	4200	4733	
BspDI	AT!CGAT	1	2965				
BspEI	T!CCGGA	0					
BspHI	T!CATGA	3	842	2551	4323		
BspMI	ACCTGCNNNN!	2	3393	3843			
BspMI	!NNNNNNNNGCAGGT	3	789	871	3012		
BspMII	T!CCGGA	0					
BspWI	GCNNNNN!NNGC	37	91	213	245	284	377
			401	676	743	752	764
			886	960	985	1098	1529
			2352	2382	2409	2451	2453
			2692	2764	2815	2894	2900
			3116	3200	3223	3362	3368
			3485	3568	3835	4022	4226
			4425	4997			
BsrFI	R!CCGGY	6	814	1041	2342	3444	3625
			3908				
BsrI	ACTGGN!	4	1664	3069	3270	4516	
BsrI	C!CAGT	5	296	2154	2832	4628	4641
BssHII	G!CGCGC	0					
Bst1107I	GTA!TAC	0					

BstBI	TT!CGAA	2	628	3807				
BstEII	G!GTNACC	0						
BstI	G!GATCC	1	1035					
BstNI	CC!WGG	19	91	284	898	933	968	
			1237	1285	1381	1396	1525	
			2661	2716	2733	3512	4091	
			4194	4884	4897	5018		
BstPI	G!GTNACC	0						
BstUI	CG!CG	15	61	1774	2049	2425	2445	
			2469	2500	3191	3492	3930	
			4014	4077	4146	4419	5000	
BstXI	CCANNNNN!NTGG	1	834					
BstYI	R!GATCY	12	586	610	938	1035	1600	
			3296	3542	3934	4293	4305	
			4391	4402				
Bsu36I	CC!TNAGG	3	1319	2605	4237			
CbiI	TT!CGAA	2	628	3807				
CcrI	C!TCGAG	1	614					
CelII	GC!TNAGC	0						
CfoI	GCG!C	29	598	723	746	752	875	
			903	917	956	1030	2391	
			2399	2425	2447	2456	2469	
			2500	3119	3127	3191	3228	
			3494	3754	3930	4016	4419	
			4528	4702	4802	4869		
Cfr10I	R!CCGGY	6	814	1041	2342	3444	3625	
			3908					
Cfr13I	G!GNCC	13	84	277	801	1210	1453	
			1545	1566	2232	3641	4086	
			4132	4190	4223			
Cfr9I	C!CCGGG	0						
CfrI	Y!GGCCR	8	947	977	1718	1769	3031	
			3205	3596	3623			
ClaI	AT!CGAT	1	2965					
CpoI	CG!GWCCG	1	3641					
Csp45I	TT!CGAA	2	628	3807				
CspI	CG!GSCCG	0						
CviJI	RG!CY	102	43	85	94	278	567	
			619	625	666	698	755	
			767	832	889	949	979	
			1009	1074	1101	1164	1211	
			1240	1247	1325	1391	1403	
			1454	1481	1489	1523	1546	
			1554	1568	1678	1714	1720	
			1771	1794	1904	2071	2091	
			2233	2314	2328	2342	2623	
			2664	2736	2859	2888	2894	
			2903	2911	2934	2946	2952	
			3033	3050	3058	3085	3110	
			3194	3203	3207	3231	3269	
			3345	3362	3381	3444	3488	
			3498	3515	3598	3625	3629	
			3666	3689	3705	3885	3912	
			3973	4095	4134	4191	4198	
			4205	4224	4235	4486	4515	
			4558	4569	4634	4713	4738	
			4743	4789	4879	4977	5003	
			5021	5032				
CvnI	CC!TNAGG	3	1319	2605	4237			

DdeI	C!TNAG	13	607 1319 4237	716 1616 4360	1004 2605 4769	1031 2907	1061 3788
DpnI	GA!TC	24	588 1576 2987 3544 4385	612 1602 3298 3920 4393	940 1784 3376 3936 4404	1037 2964 3457 4295 4479	1426 2968 3466 4307
DpnII	!GATC	24	586 1574 2985 3542 4383	610 1600 3296 3918 4391	938 1782 3374 3934 4402	1035 2962 3455 4293 4477	1424 2966 3464 4305
DraI	TTT!AAA	3	1827	4269	4288		
DraII	RG!GNCCY	1	4223				
DraIII	CACNNN!GTG	3	990	1710	2241		
DrdI	GACNNNN!NNGTC	3	2196	3152	4941		
DsaI	C!CRYGG	5	361	833	1052	2854	3557
DsaV	!CCNGG	24	89 1235 1632 3287 4664	282 1283 2659 3510 4882	896 1379 2714 3900 4895	931 1394 2731 4089 5016	966 1523 3127 4192
EaeI	Y!GGCCR	8	947 3205	977 3596	1718 3623	1769	3031
EagI	C!GGCCG	3	947	1769	3031		
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	4	1068	2596	3473	3683	
Ecl136II	GAG!CTC	1	619				
EclXI	C!GGCCG	3	947	1769	3031		
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	2	1681	4114			
Eco47I	G!GWCC	3	801	3641	4086		
Eco47III	AGC!GCT	3	597	916	1029		
Eco52I	C!GGCCG	3	947	1769	3031		
Eco57I	!CTGAAG	4	792	1548	3249	3681	
Eco57I	!CTTCAG	1	4515				
Eco81I	CC!TNAGG	3	1319	2605	4237		
EcoNI	CCTNN!NNNAGG	0					
EcoO109I	RG!GNCCY	1	4223				
EcoO65I	G!GTNACC	0					
EcoRI	G!AATTC	2	630	846			
EcoRII	!CCWGG	19	89 1235 2659 4192	282 1283 2714 4882	896 1379 2731 4895	931 1394 3510 5016	966 1523 4089
EcoRV	GAT!ATC	0					
EcoT14I	C!CWWGG	8	361 2947	827 3557	833 3962	1052	2854
EcoT22I	ATGCA!T	3	2697	2769	5100		
EcoT38I	GRGCY!C	5	621	700	2316	3490	4200
EheI	GGC!GCC	1	3126				
Esp3I	CGTCTCN!	0					
Esp3I	!NNNNNGAGACG	0					
EspI	GC!TNAGC	0					
FbaI	T!GATCA	0					
FinI	!GTCCC	4	2655	2727	2791	4086	

FinI	!GGGAC	6	161 3902	312	480	3261	3793
Fnu4HI	GC!NGC	40	735 1209 1769 2457 3187 3360 3750 4635	753 1260 1772 2895 3192 3363 3864 4638	756 1404 1902 3034 3229 3599 4203 4703	961 1521 2421 3086 3270 3695 4218 4846	964 1702 2443 3097 3357 3736 4429 5001
FokI	GGATGNNNNNNNNN!	4	2992	3449	3474	3928	
FokI	!NNNNNNNNNNNNNCATCC	5	1131	1219	1276	1431	2798
FspI	TGC!GCA	1	3227				
GdiII	!NNNNNYGGCCG	5	941	1763	3025	3590	3617
GdiII	CGGCCRN!	3	953	1775	3037		
GsuI	!CTCCAG	3	929	1392	3922		
GsuI	!CTGGAG	1	3943				
HaeI	WGG!CCW	10	832 2946	979 3207	1009 4569	1489 5021	1720 5032
HaeII	RGCGC!Y	7	599 3128	918 4803	1031	2392	2400
HaeIII	GG!CC	32	85 1009 1546 2091 2946 4134 5021	278 1211 1568 2233 3033 4191 5032	832 1454 1678 2888 3207 4224	949 1489 1720 2894 3598 4569	979 1523 1771 2903 3625 5003
HapII	C!CGG	20	602 3030 3378 4001	815 3107 3445 4475	1042 3129 3626 4665	1634 3157 3901 4691	2343 3288 3909 4838
HgaI	GACGCNNNNN!	3	527	3835	3914		
HgaI	!NNNNNNNNNNGCGTC	4	1626	2458	4353	4931	
HgiAI	GWGCW!C	5	621	1144	3238	3428	4733
HgiEII	!ACCNNNNNNGGT	1	4455				
HhaI	GCG!C	29	598 903 2399 2500 3494 4528	723 917 2425 3119 3754 4702	746 956 2447 3127 3930 4802	752 1030 2456 3191 4016 4869	875 2391 2469 3228 4419
Hin1I	GR!CGYC	8	123 3125	176 3827	259 3906	445	1637
HinPII	G!CGC	29	596 901 2397 2498 3492 4526	721 915 2423 3117 3752 4700	744 954 2445 3125 3928 4800	750 1028 2454 3189 4014 4867	873 2389 2467 3226 4417
HincII	GTY!RAC	2	642	1888			
HindIII	A!AGCTT	1	623				
HinfI	G!ANTC	16	411 1776 3744 5069	605 2168 3796	1190 2190 3854	1340 2600 3891	1667 3610 4673
HpaI	GTT!AAC	1	1888				
HpaII	C!CGG	20	602 3030	815 3107	1042 3129	1634 3157	2343 3288

			3378	3445	3626	3901	3909
			4001	4475	4665	4691	4838
HphI	GGTGANNNNNNNN!	6	377	723	895	1446	4196
			4312				
HphI	!NNNNNNNTCACC	5	918	1005	1335	2242	3303
KasI	G!GCGCC	1	3124				
KpnI	GGTAC!C	1	650				
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	4	1064	2592	3469	3679	
KspI	CCGC!GG	0					
MaeI	C!TAG	9	593	1739	1780	1974	2392
			2948	3963	4297	4550	
MaeII	A!CGT	16	123	135	176	259	340
			445	1418	1694	2185	2197
			2240	2350	3241	3428	4209
			4342				
MaeIII	!GTNAC	14	62	149	498	711	1341
			1913	2414	2426	3245	3551
			4241	4507	4623	4686	
MamI	GATNN!NNATC	2	1787	2984			
MboI	!GATC	24	586	610	938	1035	1424
			1574	1600	1782	2962	2966
			2985	3296	3374	3455	3464
			3542	3918	3934	4293	4305
			4383	4391	4402	4477	
MboII	GAAGANNNNNNN!	11	738	1081	1366	1479	1599
			1667	2381	2609	3486	3696
			4315				
MboII	!NNNNNNNTCTTC	4	3776	3943	4143	4386	
McrI	!CGRYCG	5	946	1042	1768	3030	4705
MfeI	!CAATG	1	1874				
MflI	R!GATCY	12	586	610	938	1035	1600
			3296	3542	3934	4293	4305
			4391	4402			
MluI	A!CGCGT	0					
MmeI	!TCCRAC	2	1446	2192			
MmeI	!GTYGGA	2	4668	4852			
MnlI	CCTCNNNNNN!	16	939	1223	1402	1417	1500
			1633	1744	1844	1853	2908
			2914	3726	3932	4246	4616
			4940				
MnlI	!NNNNNNNGAGG	26	541	878	989	1103	1142
			1163	1196	1313	1422	1478
			1667	1700	1803	2268	2599
			2877	2883	2920	2923	2935
			2975	3039	3175	3532	3962
			4865				
MroI	T!CCGGA	0					
MscI	TGG!CCA	3	979	1720	3207		
MseI	T!TAA	18	8	1079	1304	1760	1826
			1887	2008	2029	2040	2052
			2063	2080	2178	2449	4268
			4282	4287	4339		
MspI	C!CGG	20	602	815	1042	1634	2343
			3030	3107	3129	3157	3288
			3378	3445	3626	3901	3909
			4001	4475	4665	4691	4838
MstI	TGC!GCA	1	3227				
MstII	CC!TNAGG	3	1319	2605	4237		

MunI	C!AATTG	1	1875					
MvaI	CC!WGG	19	91	284	898	933	968	
			1237	1285	1381	1396	1525	
			2661	2716	2733	3512	4091	
			4194	4884	4897	5018		
MvnI	CG!CG	15	61	1774	2049	2425	2445	
			2469	2500	3191	3492	3930	
			4014	4077	4146	4419	5000	
NaeI	GCC!GGC	3	2344	3627	3910			
NarI	GG!CGCC	1	3125					
NciI	CC!SGG	5	1634	3129	3289	3902	4666	
NcoI	C!CATGG	5	361	833	1052	2854	3557	
NdeI	CA!TATG	1	235					
NdeII	!GATC	24	586	610	938	1035	1424	
			1574	1600	1782	2962	2966	
			2985	3296	3374	3455	3464	
			3542	3918	3934	4293	4305	
			4383	4391	4402	4477		
NgoMI	G!CCGGC	3	2342	3625	3908			
NheI	G!CTAGC	1	592					
NlaIII	CATG!	26	305	365	660	837	846	
			1056	1092	1098	1110	1188	
			1254	1334	1545	2555	2695	
			2767	2858	2999	3344	3530	
			3561	3587	3943	4327	5047	
			5098					
NlaIV	GGN!NCC	20	468	648	699	1037	1116	
			1176	1567	2280	2301	2313	
			2504	2665	2737	3126	3161	
			4009	4088	4133	4976	5015	
NotI	GC!GGCCGC	1	1769					
NruI	TCG!CGA	0						
NsiI	ATGCA!T	3	2697	2769	5100			
Nsp(7524)I	RCATG!Y	5	1092	2695	2767	3530	5047	
Nsp(7524)V	TT!CGAA	2	628	3807				
NspBII	CMG!CKG	5	1513	2623	3231	4460	4705	
NspII	GDGCH!C	14	621	700	1117	1144	1177	
			1754	2316	3071	3164	3238	
			3428	3490	4200	4733		
NspIII	C!YCGRG	1	614					
NspIV	G!GNCC	13	84	277	801	1210	1453	
			1545	1566	2232	3641	4086	
			4132	4190	4223			
NunII	GG!CGCC	1	3125					
PacI	TTAAT!TAA	0						
Paer71	C!TCGAG	1	614					
PaiI	GG!CC	32	85	278	832	949	979	
			1009	1211	1454	1489	1523	
			1546	1568	1678	1720	1771	
			2091	2233	2888	2894	2903	
			2946	3033	3207	3598	3625	
			4134	4191	4224	4569	5003	
			5021	5032				
PflMI	CCANNNN!NTGG	0						
PleI	GAGTCNNNN!	3	1348	2176	2608			
PleI	!NNNNNGACTC	6	405	599	1770	2184	3790	
			4667					
PmaCI	CAC!GTG	0						
PmeI	GTTT!AAAC	0						

PmlI	CAC!GTG	0						
PpuMI	RG!GWCCY	0						
PshAI	GACNN!NNGTC	1	1693					
PstI	CTGCA!G	1	639					
PvuI	CGAT!CG	0						
PvuII	CAG!CTG	2	2623	3231				
RmaI	C!TAG	9	593	1739	1780	1974	2392	
			2948	3963	4297	4550		
RsaI	GT!AC	14	220	245	300	333	384	
			541	648	784	1002	1021	
			1105	1255	1507	3431		
RsrI	G!AATTC	2	630	846				
RsrII	CG!GWCCG	1	3641					
SacI	GAGCT!C	1	621					
SacII	CCGC!GG	0						
SalI	G!TCGAC	1	640					
Sau3AI	!GATC	24	586	610	938	1035	1424	
			1574	1600	1782	2962	2966	
			2985	3296	3374	3455	3464	
			3542	3918	3934	4293	4305	
			4383	4391	4402	4477		
Sau96I	G!GNCC	13	84	277	801	1210	1453	
			1545	1566	2232	3641	4086	
			4132	4190	4223			
SauI	CC!TNAGG	3	1319	2605	4237			
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	24	91	284	898	933	968	
			1237	1285	1381	1396	1525	
			1634	2661	2716	2733	3129	
			3289	3512	3902	4091	4194	
			4666	4884	4897	5018		
SduI	GDGCH!C	14	621	700	1117	1144	1177	
			1754	2316	3071	3164	3238	
			3428	3490	4200	4733		
SecI	C!CNNGG	25	361	827	833	932	967	
			995	1052	1148	1283	1284	
			1379	2659	2731	2854	2889	
			2898	2947	3288	3557	3962	
			4089	4090	4192	4193	4883	
SexI	!CTCGAG	1	613					
SfaNI	GCATCNNNNN!	9	848	884	1298	1941	2704	
			2776	3424	3488	3763		
SfaNI	!NNNNNNNNGATGC	8	358	1451	1490	1619	3083	
			3338	3554	4946			
SfcI	C!TRYAG	4	635	2460	4587	4778		
SfiI	GGCCNNNN!NGGCC	1	2900					
SfuI	TT!CGAA	2	628	3807				
SgrAI	CR!CCGGYG	0						
SinI	G!GWCC	3	801	3641	4086			
SmaI	CCC!GGG	0						
SnaBI	TAC!GTA	1	341					
SnaI	!GTATAC	0						
SnoI	G!TGCAC	2	1140	4729				
SpeI	A!CTAGT	0						
SphI	GCATG!C	3	2695	2767	3530			
SplI	C!GTACG	0						
SrfI	GCCC!GGGC	0						
Sse8387I	CCTGCA!GG	0						
SspI	AAT!ATT	2	2033	2586				

SstI	GAGCT!C	1	621					
SstII	CCGC!GG	0						
SstIII	!ACGT	16	122	134	175	258	339	
			444	1417	1693	2184	2196	
			2239	2349	3240	3427	4208	
			4341					
StuI	AGG!CCT	2	1489	2946				
StyI	C!CWWGG	8	361	827	833	1052	2854	
			2947	3557	3962			
StySJI	!GAGNNNNNNNGTRC	0						
StySJI	!GYACNNNNNNNCTC	0						
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	19	615	628	641	869	919	
			1202	1229	1697	2274	2965	
			3238	3394	3418	3454	3616	
			3807	3852	4107	4945		
TaqII	GACCGANNNNNNNNNNN!	3	2145	3162	3830			
TaqII	!NNNNNNNNNTCGGTC	1	4074					
TaqII	CACCCANNNNNNNNNNN!	2	1192	1297				
TaqII	!NNNNNNNNNTGGGTG	3	173	3029	3621			
TfiI	G!AWTC	7	1190	1667	3610	3744	3854	
			3891	5069				
ThaI	CG!CG	15	61	1774	2049	2425	2445	
			2469	2500	3191	3492	3930	
			4014	4077	4146	4419	5000	
Tsp45I	!GTSAC	5	711	1341	2414	3245	3551	
TspEI	!AATT	13	19	630	846	1761	1875	
			1939	2017	2043	2054	2701	
			2773	2865	4283			
Tth111I	GACN!NNGTC	1	3243					
Tth111II	CAARCANNNNNNNNNNN!	2	997	4452				
Tth111III	!NNNNNNNNNTGYTTG	3	3560	4420	4459			
TthHB8I	T!CGA	19	615	628	641	869	919	
			1202	1229	1697	2274	2965	
			3238	3394	3418	3454	3616	
			3807	3852	4107	4945		
Van91I	CCANNNN!NTGG	0						
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	1779					
XcmI	!CCANNNNNNNNTCG	1	3400					
XcyI	C!CCGGG	0						
XhoI	C!TCGAG	1	614					
XhoII	R!GATCY	12	586	610	938	1035	1600	
			3296	3542	3934	4293	4305	
			4391	4402				
XmaI	C!CCGGG	0						
XmaIII	C!GGCCG	3	947	1769	3031			
XmnI	GAANN!NNTTC	0						
XorII	CGAT!CG	0						