

Enzyme Name	Sequence	Count	Cutting Position				
AatI	AGG!CCT	1	2698				
AatII	GACGT!C	4	126	179	262	448	
Acc65I	G!GTACC	0					
AccI	GT!MKAC	0					
AccII	CG!CG	23	61	663	665	756	758
			1211	1364	1436	1454	1526
			1801	2177	2197	2221	2252
			2943	3244	3682	3766	3829
			3898	4171	4752		
AccIII	T!CCGGA	0					
AciI	G!CGG	30	371	404	529	624	648
			717	741	812	1313	1520
			2137	2164	2252	2362	2943
			3007	3108	3111	3391	3446
			3544	3682	3829	3969	4212
			4457	4578	4597	4724	4752
AciI	C!CGC	43	59	87	99	113	280
			508	590	613	661	706
			754	818	1020	1083	1161
			1304	1362	1482	1524	2192
			2195	2209	2299	2547	2559
			2568	2580	2590	2601	2647
			2786	2849	3351	3396	3462
			3488	3613	3616	3764	3896
			4203	4347	4843		
AcyI	GR!CGYC	9	123	176	259	445	610
			703	2877	3579	3658	
AfaI	GT!AC	8	220	245	300	333	384
			541	1000	3183		
AflI	G!GWCC	2	3393	3838			
AflII	C!TTAAG	1	1759				
AflIII	A!CRYGT	3	1247	1442	4795		
AgeI	A!CCGGT	1	786				
AhaI	CC!SGG	7	645	738	1379	2881	3041
			3654	4418			
AhaII	GR!CGYC	9	123	176	259	445	610
			703	2877	3579	3658	
AhaIII	TTT!AAA	3	1579	4021	4040		
AluI	AG!CT	17	567	960	971	1137	1218
			1281	1656	1823	2080	2375
			2663	2983	3441	4238	4495
			4541	4631			
Alw44I	G!TGCAC	2	934	4481			
AlwI	GGATCNNNN!	9	695	788	2745	3056	3302
			3694	4053	4151	4237	
AlwI	!NNNNNGATCC	10	581	666	759	775	987
			1368	3121	3665	4052	4149
AlwNI	CAGNNN!CTG	2	3978	4386			
AocI	CC!TNAGG	2	2357	3989			
AocII	GDGCH!C	11	938	1010	1481	2068	2823
			2916	2990	3180	3242	3952
			4485				
Aor51HI	AGC!GCT	0					
AosI	TGC!GCA	1	2979				
AosII	GR!CGYC	9	123	176	259	445	610
			703	2877	3579	3658	

ApaI	GGGCC!C	0						
ApaLI	G!TGCAC	2	934	4481				
ApoI	R!AATTY	3	1691	1795	1806			
ApyI	CC!WGG	12	91	284	1051	2413	2468	
			2485	3264	3843	3946	4636	
			4649	4770				
AquI	C!YCGRG	1	1063					
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	4	938	2990	3180	4485		
AspI	GACN!NNGTC	1	2995					
AsuI	G!GNCC	16	84	277	642	735	814	
			1068	1198	1261	1326	1381	
			1984	3393	3838	3884	3942	
			3975					
AsuII	TT!CGAA	1	3559					
AvaI	C!YCGRG	1	1063					
AvaII	G!GWCC	2	3393	3838				
AvaIII	ATGCA!T	3	2449	2521	4852			
AviII	TGC!GCA	1	2979					
AvrI	!CYCGRG	1	1062					
AvrII	C!CTAGG	2	2699	3714				
AxyI	CC!TNAGG	2	2357	3989				
BalI	TGG!CCA	1	2959					
BamHI	G!GATCC	1	780					
BanI	G!GYRCC	5	466	1108	2030	2876	2911	
BanII	GRGCY!C	4	1010	2068	3242	3952		
BanIII	AT!CGAT	1	2717					
BbeI	GGCGC!C	1	2880					
BbiIII/AcyI	GR!CGYC	9	123	176	259	445	610	
			703	2877	3579	3658		
BbrPI	CAC!GTG	1	1443					
BbsI	!GTCTTC	0						
BbsI	!GAAGAC	0						
BbvI	GCAGCNNNNNNNN!	9	914	917	1370	1665	2950	
			2992	3513	3966	4466		
BbvI	!NNNNNNNNNNNGCTGC	14	602	605	608	695	698	
			701	947	2159	2824	3008	
			3101	4167	4373	4376		
BbvII	GAAGACNN!	0						
BbvII	!NNNNNNGTCTTC	0						
BcefI	ACGGCNNNNNNNNNN!	14	876	888	915	936	1017	
			1047	1122	1197	1212	1275	
			1326	1371	3303	3790		
BcefI	!NNNNNNNNNNNGCCGT	5	926	1256	2019	2837	4296	
BcgI	!GCANNNNNNTCG	0						
BcgI	!CGANNNNNNTGC	0						
BclI	T!GATCA	0						
BcnI	CC!SGG	7	645	738	1379	2881	3041	
			3654	4418				
BfaI	C!TAG	8	593	1532	1726	2144	2700	
			3715	4049	4302			
BfrI	C!TTAAG	1	1759					
BglI	GCCNNNN!NGGC	4	91	213	284	2652		
BglII	A!GATCT	0						
BinI	!NNNNNGATCC	10	581	666	759	775	987	

			1368	3121	3665	4052	4149
BinI	GGATCNNNN!	9	695	788	2745	3056	3302
			3694	4053	4151	4237	
BlnI	C!CTAGG	2	2699	3714			
BmyI	GDGCH!C	11	938	1010	1481	2068	2823
			2916	2990	3180	3242	3952
			4485				
BpmI	!CTCCAG	1	3674				
BpmI	!CTGGAG	1	3695				
Bpu1102I	GC!TNAGC	0					
BsaAI	YAC!GTR	4	341	1443	1993	3181	
BsaBI	GATNN!NNATC	2	1539	2736			
BsaHI	GR!CGYC	9	123	176	259	445	610
			703	2877	3579	3658	
BsaI	GGTCTCN!	0					
BsaI	!NNNNNGAGACC	3	943	1234	3866		
BsaJI	C!CNNGG	21	361	797	918	1049	1064
			1257	1377	2411	2483	2606
			2641	2650	2699	3040	3309
			3714	3841	3842	3944	3945
			4635				
BseAI	T!CCGGA	0					
BsePI	!GCGCGC	2	662	755			
BsgI	!GTGCAG	0					
BsgI	!CTGCAC	0					
BsiEI	CGRY!CG	5	791	1083	1524	2786	4461
BsiWI	C!GTACG	0					
BsiYI	CCNNNNN!NNGG	24	50	681	774	971	1221
			1323	1455	1889	2215	2607
			2858	3402	3683	3715	3716
			3847	3882	3932	3933	3964
			4317	4596	4762	4780	
BslI	CCNNNNN!NNGG	24	50	681	774	971	1221
			1323	1455	1889	2215	2607
			2858	3402	3683	3715	3716
			3847	3882	3932	3933	3964
			4317	4596	4762	4780	
BsmAI	!GTCTC	1	429				
BsmAI	!GAGAC	6	948	1239	2306	2724	3743
			3871				
BsmI	GAATGCN!	1	1628				
BsmI	G!CATTC	1	1721				
Bsp1286I	GDGCH!C	11	938	1010	1481	2068	2823
			2916	2990	3180	3242	3952
			4485				
BspDI	AT!CGAT	1	2717				
BspEI	T!CCGGA	0					
BspHI	T!CATGA	2	2303	4075			
BspMI	ACCTGCNNNN!	3	1156	3145	3595		
BspMI	!NNNNNNNNGCAGGT	1	2764				
BspMII	T!CCGGA	0					
BspWI	GCNNNNN!NNGC	44	91	213	245	284	377
			401	618	621	635	657
			711	714	728	750	953
			1082	1091	1157	1215	1433
			1481	1490	2104	2134	2161
			2203	2205	2444	2516	2567
			2646	2652	2868	2952	2975
			3114	3120	3237	3320	3587

			3774	3978	4177	4749	
BsrFI	R!CCGGY	6	786	952	2094	3196	3377
			3660				
BsrI	ACTGGN!	3	2821	3022	4268		
BsrI	C!CAGT	8	296	655	748	996	1906
			2584	4380	4393		
BssHII	G!CGCGC	2	663	756			
Bst1107I	GTA!TAC	0					
BstBI	TT!CGAA	1	3559				
BstEII	G!GTNACC	0					
BstI	G!GATCC	1	780				
BstNI	CC!WGG	12	91	284	1051	2413	2468
			2485	3264	3843	3946	4636
			4649	4770			
BstPI	G!GTNACC	0					
BstUI	CG!CG	23	61	663	665	756	758
			1211	1364	1436	1454	1526
			1801	2177	2197	2221	2252
			2943	3244	3682	3766	3829
			3898	4171	4752		
BstXI	CCANNNNN!NTGG	0					
BstYI	R!GATCY	13	586	671	687	764	780
			1373	3048	3294	3686	4045
			4057	4143	4154		
Bsu36I	CC!TNAGG	2	2357	3989			
CbiI	TT!CGAA	1	3559				
CcrI	C!TCGAG	0					
CelII	GC!TNAGC	0					
CfoI	GCG!C	31	665	667	758	760	1094
			1211	1277	1366	1436	1454
			1495	2143	2151	2177	2199
			2208	2221	2252	2871	2879
			2943	2980	3246	3506	3682
			3768	4171	4280	4454	4554
			4621				
Cfr10I	R!CCGGY	6	786	952	2094	3196	3377
			3660				
Cfr13I	G!GNCC	16	84	277	642	735	814
			1068	1198	1261	1326	1381
			1984	3393	3838	3884	3942
			3975				
Cfr9I	C!CCGGG	0					
CfrI	Y!GGCCR	7	873	893	1521	2783	2957
			3348	3375			
ClaI	AT!CGAT	1	2717				
CpoI	CG!GWCCG	1	3393				
Csp45I	TT!CGAA	1	3559				
CspI	CG!GSCCG	2	642	735			
CviJI	RG!CY	105	43	85	94	278	567
			629	638	643	651	722
			731	736	744	815	875
			895	960	971	978	1008
			1040	1070	1121	1137	1160
			1184	1191	1199	1218	1262
			1281	1285	1295	1316	1328
			1361	1382	1462	1523	1546
			1656	1823	1843	1985	2066
			2080	2094	2375	2416	2488
			2611	2640	2646	2655	2663

			2686	2698	2704	2785	2802
			2810	2837	2862	2946	2955
			2959	2983	3021	3097	3114
			3133	3196	3240	3250	3267
			3350	3377	3381	3418	3441
			3457	3637	3664	3725	3847
			3886	3943	3950	3957	3976
			3987	4238	4267	4310	4321
			4386	4465	4490	4495	4541
			4631	4729	4755	4773	4784
CvnI	CC!TNAGG	2	2357	3989			
DdeI	C!TNAG	7	1072	2357	2659	3540	3989
			4112	4521			
DpnI	GA!TC	24	588	673	689	766	782
			994	1375	1536	2716	2720
			2739	3050	3128	3209	3218
			3296	3672	3688	4047	4059
			4137	4145	4156	4231	
DpnII	!GATC	24	586	671	687	764	780
			992	1373	1534	2714	2718
			2737	3048	3126	3207	3216
			3294	3670	3686	4045	4057
			4135	4143	4154	4229	
DraI	TTT!AAA	3	1579	4021	4040		
DraII	RG!GNCCY	2	1068	3975			
DraIII	CACNNN!GTG	4	942	1392	1476	1993	
DrdI	GACNNNN!NNGTC	3	1948	2904	4693		
DsaI	C!CRYGG	6	361	797	918	1257	2606
			3309				
DsaV	!CCNGG	19	89	282	643	736	1049
			1377	2411	2466	2483	2879
			3039	3262	3652	3841	3944
			4416	4634	4647	4768	
EaeI	Y!GGCCR	7	873	893	1521	2783	2957
			3348	3375			
EagI	C!GGCCG	2	1521	2783			
Eam1105I	GACNNN!NNGTC	0					
EarI	!CTCTTC	0					
EarI	!GAAGAG	3	2348	3225	3435		
Ecl136II	GAG!CTC	0					
EclXI	C!GGCCG	2	1521	2783			
Eco105I	TAC!GTA	1	341				
Eco31I	GGTCTCN!	0					
Eco31I	!NNNNNGAGACC	3	943	1234	3866		
Eco47I	G!GWCC	2	3393	3838			
Eco47III	AGC!GCT	0					
Eco52I	C!GGCCG	2	1521	2783			
Eco57I	!CTGAAG	2	3001	3433			
Eco57I	!CTTCAG	1	4267				
Eco81I	CC!TNAGG	2	2357	3989			
EcoNI	CCTNN!NNNAGG	0					
EcoO109I	RG!GNCCY	2	1068	3975			
EcoO65I	G!GTNACC	0					
EcoRI	G!AATTC	0					
EcoRII	!CCWGG	12	89	282	1049	2411	2466
			2483	3262	3841	3944	4634
			4647	4768			
EcoRV	GAT!ATC	0					
EcoT14I	C!CWGGG	6	361	797	2606	2699	3309

			3714					
EcoT22I	ATGCA!T	3	2449	2521	4852			
EcoT38I	GRGCY!C	4	1010	2068	3242	3952		
EheI	GGC!GCC	1	2878					
Esp3I	CGTCTCN!	0						
Esp3I	!NNNNNGAGACG	0						
EspI	GC!TNAGC	0						
FbaI	T!GATCA	0						
FinI	!GTCCC	4	2407	2479	2543	3838		
FinI	!GGGAC	7	161	312	480	1417	3013	
			3545	3654				
Fnu4HI	GC!NGC	50	613	616	619	622	649	
			661	706	709	712	715	
			742	754	813	903	906	
			961	1161	1314	1359	1362	
			1521	1524	1654	2173	2195	
			2209	2647	2786	2838	2849	
			2939	2944	2981	3022	3109	
			3112	3115	3351	3447	3488	
			3502	3616	3955	3970	4181	
			4387	4390	4455	4598	4753	
FokI	GGATGNNNNNNNNN!	4	2744	3201	3226	3680		
FokI	!NNNNNNNNNNNNNCATCC	2	1213	2550				
FspI	TGC!GCA	1	2979					
GdiII	!NNNNNYGGCCG	5	887	1515	2777	3342	3369	
GdiII	CGGCCRN!	3	879	1527	2789			
GsuI	!CTCCAG	1	3674					
GsuI	!CTGGAG	1	3695					
HaeI	WGG!CCW	7	1285	1462	2698	2959	4321	
			4773	4784				
HaeII	RGCGC!Y	5	1496	2144	2152	2880	4555	
HaeIII	GG!CC	33	85	278	643	736	815	
			875	895	1070	1199	1262	
			1285	1316	1328	1382	1462	
			1523	1843	1985	2640	2646	
			2655	2698	2785	2959	3350	
			3377	3886	3943	3976	4321	
			4755	4773	4784			
HapII	C!CGG	21	645	738	787	953	1379	
			2095	2782	2859	2881	2909	
			3040	3130	3197	3378	3653	
			3661	3753	4227	4417	4443	
			4590					
HgaI	GACGCNNNNN!	5	527	618	711	3587	3666	
HgaI	!NNNNNNNNNNNGCGTC	3	2210	4105	4683			
HgiAI	GWGCW!C	4	938	2990	3180	4485		
HgiEII	!ACNNNNNNNGGT	1	4207					
HhaI	GCG!C	31	665	667	758	760	1094	
			1211	1277	1366	1436	1454	
			1495	2143	2151	2177	2199	
			2208	2221	2252	2871	2879	
			2943	2980	3246	3506	3682	
			3768	4171	4280	4454	4554	
			4621					
Hin1I	GR!CGYC	9	123	176	259	445	610	
			703	2877	3579	3658		
HinP1I	G!CGC	31	663	665	756	758	1092	
			1209	1275	1364	1434	1452	
			1493	2141	2149	2175	2197	

			2206	2219	2250	2869	2877
			2941	2978	3244	3504	3680
			3766	4169	4278	4452	4552
			4619				
HincII	GTY!RAC	1	1640				
HindIII	A!AGCTT	0					
HinfI	G!ANTC	12	411	1528	1920	1942	2352
			3362	3496	3548	3606	3643
			4425	4821			
HpaI	GTT!AAC	1	1640				
HpaII	C!CGG	21	645	738	787	953	1379
			2095	2782	2859	2881	2909
			3040	3130	3197	3378	3653
			3661	3753	4227	4417	4443
			4590				
HphI	GGTGANNNNNNNN!	5	377	879	1167	3948	4064
HphI	!NNNNNNNTCACC	7	876	1158	1290	1395	1479
			1994	3055			
KasI	G!GCGCC	1	2876				
KpnI	GGTAC!C	0					
Ksp632I	CTCTTCN!	0					
Ksp632I	!NNNNGAAGAG	3	2344	3221	3431		
KspI	CCGC!GG	0					
MaeI	C!TAG	8	593	1532	1726	2144	2700
			3715	4049	4302		
MaeII	A!CGT	16	123	135	176	259	340
			445	932	1442	1937	1949
			1992	2102	2993	3180	3961
			4094				
MaeIII	!GTNAC	13	62	149	498	1392	1665
			2166	2178	2997	3303	3993
			4259	4375	4438		
MamI	GATNN!NNATC	2	1539	2736			
MboI	!GATC	24	586	671	687	764	780
			992	1373	1534	2714	2718
			2737	3048	3126	3207	3216
			3294	3670	3686	4045	4057
			4135	4143	4154	4229	
MboII	GAAGANNNNNNNN!	5	2133	2361	3238	3448	4067
MboII	!NNNNNNNTCTTC	6	843	1005	3528	3695	3895
			4138				
McrI	!CGRYCG	5	787	1079	1520	2782	4457
MfeI	!CAATTG	1	1626				
MflI	R!GATCY	13	586	671	687	764	780
			1373	3048	3294	3686	4045
			4057	4143	4154		
MluI	A!CGCGT	0					
MmeI	!TCCRAC	1	1944				
MmeI	!GTYGGA	2	4420	4604			
MnlI	CCTCNNNNNNN!	9	1596	1605	2660	2666	3478
			3684	3998	4368	4692	
MnlI	!NNNNNNNGAGG	21	541	800	857	1058	1408
			1448	1502	1555	2020	2351
			2629	2635	2672	2675	2687
			2727	2791	2927	3284	3714
			4617				
MroI	T!CCGGA	0					
MscI	TGG!CCA	1	2959				
MseI	T!TAA	16	8	1515	1578	1639	1760

			1781	1792	1804	1815	1832
			1930	2201	4020	4034	4039
			4091				
MspI	C!CGG	21	645	738	787	953	1379
			2095	2782	2859	2881	2909
			3040	3130	3197	3378	3653
			3661	3753	4227	4417	4443
			4590				
MstI	TGC!GCA	1	2979				
MstII	CC!TNAGG	2	2357	3989			
MunI	C!AATTG	1	1627				
MvaI	CC!WGG	12	91	284	1051	2413	2468
			2485	3264	3843	3946	4636
			4649	4770			
MvnI	CG!CG	23	61	663	665	756	758
			1211	1364	1436	1454	1526
			1801	2177	2197	2221	2252
			2943	3244	3682	3766	3829
			3898	4171	4752		
NaeI	GCC!GGC	3	2096	3379	3662		
NarI	GG!CGCC	1	2877				
NciI	CC!SGG	7	645	738	1379	2881	3041
			3654	4418			
NcoI	C!CATGG	4	361	797	2606	3309	
NdeI	CA!TATG	1	235				
NdeII	!GATC	24	586	671	687	764	780
			992	1373	1534	2714	2718
			2737	3048	3126	3207	3216
			3294	3670	3686	4045	4057
			4135	4143	4154	4229	
NgoMI	G!CCGGC	3	2094	3377	3660		
NheI	G!CTAGC	1	592				
NlaIII	CATG!	22	305	365	600	801	840
			969	1116	1209	1251	2307
			2447	2519	2610	2751	3096
			3282	3313	3339	3695	4079
			4799	4850			
NlaIV	GGN!NCC	21	468	652	745	782	816
			1110	1200	1263	2032	2053
			2065	2256	2417	2489	2878
			2913	3761	3840	3885	4728
			4767				
NotI	GC!GGCCGC	1	1521				
NruI	TCG!CGA	0					
NsiI	ATGCA!T	3	2449	2521	4852		
Nsp (7524) I	RCATG!Y	6	600	1251	2447	2519	3282
			4799				
Nsp (7524) V	TT!CGAA	1	3559				
NspBII	CMG!CKG	8	615	708	1218	1281	2375
			2983	4212	4457		
NspII	GDGCH!C	11	938	1010	1481	2068	2823
			2916	2990	3180	3242	3952
			4485				
NspIII	C!YCGRG	1	1063				
NspIV	G!GNCC	16	84	277	642	735	814
			1068	1198	1261	1326	1381
			1984	3393	3838	3884	3942
			3975				
NunII	GG!CGCC	1	2877				

PacI	TTAAT!TAA	0						
Paer71	C!TCGAG	0						
PalI	GG!CC	33	85	278	643	736	815	
			875	895	1070	1199	1262	
			1285	1316	1328	1382	1462	
			1523	1843	1985	2640	2646	
			2655	2698	2785	2959	3350	
			3377	3886	3943	3976	4321	
			4755	4773	4784			
PflMI	CCANNNN!NTGG	4	681	774	971	1221		
PleI	GAGTCNNNN!	2	1928	2360				
PleI	!NNNNNGACTC	5	405	1522	1936	3542	4419	
PmaCI	CAC!GTG	1	1443					
PmeI	GTTT!AAAC	0						
PmlI	CAC!GTG	1	1443					
PpuMI	RG!GWCCY	0						
PshAI	GACNN!NNGTC	0						
PstI	CTGCA!G	0						
PvuI	CGAT!CG	0						
PvuII	CAG!CTG	4	1218	1281	2375	2983		
RmaI	C!TAG	8	593	1532	1726	2144	2700	
			3715	4049	4302			
RsaI	GT!AC	8	220	245	300	333	384	
			541	1000	3183			
RsrI	G!AATTC	0						
RsrII	CG!GWCCG	1	3393					
SacI	GAGCT!C	0						
SacII	CCGC!GG	0						
SalI	G!TCGAC	0						
Sau3AI	!GATC	24	586	671	687	764	780	
			992	1373	1534	2714	2718	
			2737	3048	3126	3207	3216	
			3294	3670	3686	4045	4057	
			4135	4143	4154	4229		
Sau96I	G!GNCC	16	84	277	642	735	814	
			1068	1198	1261	1326	1381	
			1984	3393	3838	3884	3942	
			3975					
SauI	CC!TNAGG	2	2357	3989				
ScaI	AGT!ACT	0						
ScrFI	CC!NGG	19	91	284	645	738	1051	
			1379	2413	2468	2485	2881	
			3041	3264	3654	3843	3946	
			4418	4636	4649	4770		
SduI	GDGCH!C	11	938	1010	1481	2068	2823	
			2916	2990	3180	3242	3952	
			4485					
SecI	C!CNNGG	21	361	797	918	1049	1064	
			1257	1377	2411	2483	2606	
			2641	2650	2699	3040	3309	
			3714	3841	3842	3944	3945	
			4635					
SexI	!CTCGAG	0						
SfaNI	GCATCNNNNN!	10	1043	1085	1172	1493	1693	
			2456	2528	3176	3240	3515	
SfaNI	!NNNNNNNNNGATGC	5	358	2835	3090	3306	4698	
SfcI	C!TRYAG	3	2212	4339	4530			
SfiI	GGCCNNNN!NGGCC	1	2652					
SfuI	TT!CGAA	1	3559					

SgrAI	CR!CCGGYG	0						
SinI	G!GWCC	2	3393	3838				
SmaI	CCC!GGG	0						
SnaBI	TAC!GTA	1	341					
SnaI	!GTATAC	0						
SnoI	G!TGCAC	2	934	4481				
SpeI	A!CTAGT	0						
SphI	GCATG!C	3	2447	2519	3282			
SplI	C!GTACG	0						
SrfI	GCCC!GGGC	0						
Sse8387I	CCTGCA!GG	0						
SspI	AAT!ATT	2	1785	2338				
SstI	GAGCT!C	0						
SstII	CCGC!GG	0						
SstIII	!ACGT	16	122	134	175	258	339	
			444	931	1441	1936	1948	
			1991	2101	2992	3179	3960	
			4093					
StuI	AGG!CCT	1	2698					
StyI	C!CWGGG	6	361	797	2606	2699	3309	
			3714					
StySJI	!GAGNNNNNGTRC	0						
StySJI	!GYACNNNNNCTC	0						
SwaI	ATTT!AAAT	0						
TaqI	T!CGA	16	857	1079	1097	1334	1370	
			2026	2717	2990	3146	3170	
			3206	3368	3559	3604	3859	
			4697					
TaqII	GACCGANNNNNNNNNN!	3	1897	2914	3582			
TaqII	!NNNNNNNNNTCGGTC	1	3826					
TaqII	CACCCANNNNNNNNNN!	0						
TaqII	!NNNNNNNNNTGGGTG	3	173	2781	3373			
TfiI	G!AWTC	5	3362	3496	3606	3643	4821	
ThaI	CG!CG	23	61	663	665	756	758	
			1211	1364	1436	1454	1526	
			1801	2177	2197	2221	2252	
			2943	3244	3682	3766	3829	
			3898	4171	4752			
Tsp45I	!GTSAC	4	1392	2166	2997	3303		
TspEI	!AATT	10	19	1627	1691	1769	1795	
			1806	2453	2525	2617	4035	
Tth111I	GACN!NNGTC	1	2995					
Tth111II	CAARCANNNNNNNNNN!	2	1423	4204				
Tth111III	!NNNNNNNNNTGYTTG	3	3312	4172	4211			
TthHB8I	T!CGA	16	857	1079	1097	1334	1370	
			2026	2717	2990	3146	3170	
			3206	3368	3559	3604	3859	
			4697					
Van91I	CCANNNN!NTGG	4	681	774	971	1221		
VspI	AT!TAAT	1	8					
XbaI	T!CTAGA	1	1531					
XcmI	!CCANNNNNNNNTCG	1	3152					
XcyI	C!CCGGG	0						
XhoI	C!TCGAG	0						
XhoII	R!GATCY	13	586	671	687	764	780	
			1373	3048	3294	3686	4045	
			4057	4143	4154			
XmaI	C!CCGGG	0						
XmaIII	C!GGCCG	2	1521	2783				

XmnI	GAANN!NNTTC	0	
XorII	CGAT!CG	0	