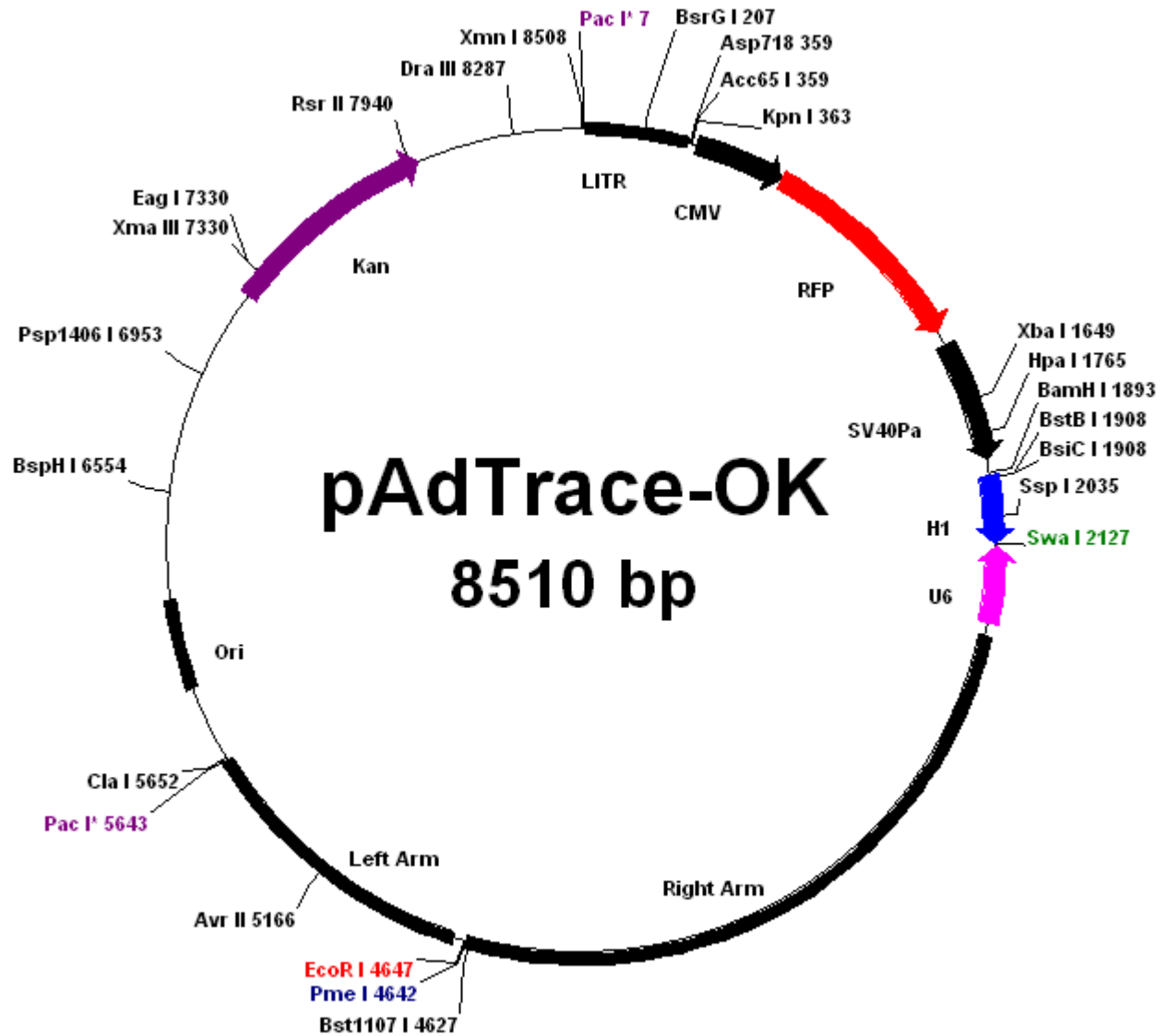


Vector: pAdTrace-OK

Antibiotic Selection: Kan

Creator(S): Fang Deng, Qiang Wei, Molecular Oncology Lab of The University of Chicago Medical Center

Date of Construction: August, 2014



pAdTrace-OK Full-Length Sequence

NNNTAATTAANNNTCCCTTCCAGCTCTCTGCCCTTTTGGATTGAAGCCAATATGATAATGAGGGGGTGGAGTTTGTGACGTGGCGCGGG
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Unique enzymes in pAdTrace-OK

BsrG I	T`GTAC,A	207
Acc65 I	G`GTAC,C	359
Asp718	G`GTAC,C	359
Kpn I	G,GTAC`C	363
Xba I	T`CTAG,A	1649
Hpa I	GTT AAC	1765
BamH I	G`GATC,C	1893
BsiC I	TT`CG,AA	1908
BstB I	TT`CG,AA	1908
Ssp I	AAT ATT	2035
Swa I	ATT AAAT	2127
Bst1107 I	GTA TAC	4627
Xca I	GTA TAC	4627
Pme I	CTTT AAAC	4642
EcoR I	G`AATC,C	4647
Avr II	C`CTAG,G	5166
Cla I	AT`CG,AT	5652
BspH I	T`CATG,A	6554
Psp1406 I	AA`CG,TT	6953
Eag I	C`GGCC,G	7330
Xma III	C`GGCC,G	7330
Rsr II	CG`GWC,CG	7940
Dra III	CAC,NNN`GTG	8287
Xmn I	GAANN NNTTC	8508

Number of enzymes = 24

pAdTrace-OK: sites sorted by name:

Aat II	(6)	489	542	625	811
		982	1921		
Acc I	(2)	3688	4626		
Acc65 I	(1)	359			
Aci I	(112)	88	106	126	151
		233	243	294	422
		450	462	476	643
		734	767	871	892
		953	1067	1090	1114
		1189	1267	1336	1406
		1471	1619	1890	1929
		1944	1963	2459	2462
		2465	2591	2654	2657
		2660	2669	2681	2685
		2687	2719	2750	2754
		2884	2973	2991	3119
		3237	3490	3493	3578
		3582	3599	3695	3775
		3914	3931	3989	4032
		4093	4103	4339	4341
		4383	4459	4464	4473
		4801	5073	5187	5217
		5392	5524	5545	5563
		5674	5700	5717	5760
		5767	5788	5879	5907
		6034	6053	6174	6284
		6419	6428	6790	6881
		7011	7134	7230	7333
		7396	7490	7554	7655
		7658	7898	7938	7943
		7993	8009	8035	8177
		8316	8431	8458	8486
Afl III	(3)	158	4793	5834	
Aha II	(15)	486	539	622	808
		979	1426	1483	1534
		1633	1918	2024	3427

The following enzymes do not cut in pAdTrace-OK:

Afl II	Age I	Asc I	Bgl II	Blp I
BsiW I	Bsu36 I	Eco72 I	EcoN I	EcoR V
Esp I	Fse I	HinD III	Mlu I	Nhe I
Not I	Nru I	PaeR7 I	Pml I	Pvu I
Sal I	Sca I	Sfi I	Spe I	Spl I

pAdTrace-OK Vector

Dra I	(9)	1704	2127	2206	2219	HinI I	(15)	486	539	622	808
		2890	4642	4845	6593			979	1426	1483	1534
		6612						1633	1918	2024	3427
Dra III	(1)	8287						4380	7176	7424	
Drd I	(5)	1488	4319	5942	7451	HinP I	(64)	86	200	231	348
		8242						998	1010	1244	1334
Dsa I	(10)	724	961	1381	2530			1426	1534	1607	1609
		2685	3484	4163	4315			1633	1976	1978	1984
		4339	7856					2024	2520	2841	2967
Eae I	(12)	283	1031	1472	1507			2969	2971	3156	3576
		4231	4297	4511	4906			3924	4169	4213	4225
		7330	7504	7895	7922			4227	4241	4271	4291
Eag I	(1)	7330						4356	4358	4380	4426
Ear I	(4)	2375	5718	7768	7978			4461	4698	5207	5565
Eco47 III	(3)	3157	4272	5208				5709	5742	6012	6079
Eco57 I	(10)	1118	1451	1475	1481			6179	6353	6462	6855
		2649	2880	4286	6381			6948	7078	7176	7247
		7569	8001					7416	7424	7488	7525
EcoO109 I	(5)	2365	3009	3212	4175			7791	8051	8435	8443
		4469						8469	8488	8490	8499
EcoR I	(1)	4647				Hpa I	(1)	1765			
EcoR II	(20)	452	645	1131	1285	Hpa II	(39)	203	1531	1630	1644
		1987	3025	3202	3496			2537	2802	2982	3327
		3541	3621	4027	4146			3367	3617	3719	3781
		4233	4299	4498	4530			3809	4218	4277	4444
		5860	5981	5994	7809			4583	4754	4914	4981
Ehe I	(7)	1427	1535	1634	2025			5057	6041	6188	6214
		4381	7177	7425				6404	6808	6842	6909
Fnu4H I	(69)	1067	1115	1268	1332			7031	7158	7329	7406
		1337	1472	1524	1527			7428	7456	7587	7677
		1619	1779	2453	2456			7744	7925	8389	
		2459	2462	2465	2648	Hph I	(19)	120	739	1114	1285
		2657	2660	2669	2672			1336	1557	2058	3052
		2675	2735	2827	3117			4042	4213	4513	4609
		3491	3494	3762	3765			4931	5534	6571	6798
		3778	3835	3838	3841			6972	7603	8289	
		3931	4104	4223	4274	Kas I	(7)	1425	1533	1632	2023
		4342	4361	4383	4464			4379	7175	7423	
		4672	5218	5221	5740						
		5758	5761	5879	6034	Kpn I	(1)	363			
		6177	6242	6245	6451	Mae I	(12)	364	956	1650	1851
		6779	7230	7333	7385			1959	5167	6329	6582
		7396	7486	7491	7528			6917	6983	7125	8438
		7569	7656	7659	7662	Mae II	(32)	81	113	187	486
		7898	7994	8035	8049			498	539	622	703
		8467						808	979	1177	1918
Fok I	(20)	184	258	1125	1182			2074	2255	2260	3031
		1421	2733	3063	3262			3261	3526	5355	5467
		3558	3600	3834	4442			5472	5538	5570	6537
		6693	6874	7046	7229			6953	6996	7540	7727
		7291	7748	7773	8137			8231	8243	8286	8396
Fsp I	(3)	1011	6949	7526		Mae III	(41)	77	109	183	218
Gdi II	(9)	1032	1473	4298	4510			266	337	425	512
		4907	7329	7331	7894			861	1103	1274	1280
		7921						1790	1954	2177	2256
Gsu I	(3)	2548	5176	6798				2694	3041	3434	3632
Hae I	(11)	285	966	1173	1395			4047	4196	4598	4918
		1509	3562	4233	5849			4936	5298	5356	5449
		5860	6312	7506				5468	5473	5517	5539
Hae II	(15)	1429	1537	1636	2027			5571	6190	6253	6369
		3159	3927	4274	4383			6652	7544	7850	8460
		5210	5712	6082	7179			8472			
		7427	8438	8446		Mbo I	(35)	949	1045	1441	1646
Hae III	(44)	285	448	641	966			1659	1893	2387	2835
		1033	1066	1117	1173			2927	3137	3517	3550
		1360	1395	1474	1509			3680	3731	4069	5649
		1618	1948	2366	2536			5653	6400	6475	6486
		2980	3213	3482	3496			6494	6572	6584	6689
		3562	3620	3788	3865			6973	6985	7237	7254
		4146	4233	4299	4349			7265	7284	7595	7673
		4470	4513	4908	5317			7754	7763	7841	
		5849	5860	5878	6312	Mbo II	(17)	1384	1387	2007	2363
		6770	6850	7332	7506			3425	3515	3716	3753
		7897	7924	8137	8279			4274	5706	6495	6568
Hga I	(7)	889	1490	2534	3434			6969	7784	7994	8076
		4283	5944	6522				8426			
HgiA I	(8)	2476	2506	3374	4134	Mme I	(6)	1377	2619	2814	6048
		4509	6152	7537	7727			6232	8263		
HgiE II	(2)	3772	6413			Mnl I	(68)	56	304	905	968
Hha I	(64)	88	202	233	350			976	979	1010	1043
		1000	1012	1246	1336			1049	1055	1070	1158
		1428	1536	1609	1611			1220	1253	1303	1360
		1635	1978	1980	1986			1385	1405	1412	1444
		2026	2522	2843	2969			1481	1571	1576	1607
		2971	2973	3158	3578			1681	1720	1729	2001
		3926	4171	4215	4227			2377	2660	2879	2884
		4229	4243	4273	4293			2999	3083	3333	3448
		4358	4360	4382	4428			3578	3647	4093	4187
		4463	4700	5209	5567			4240	4345	4367	4370
		5711	5744	6014	6081			4382	4402	4575	4703
		6181	6355	6464	6857			4978	5101	5168	5192
		6950	7080	7178	7249			5450	5597	5732	5942
		7418	7426	7490	7527			6015	6266	6666	6747
		7793	8053	8437	8445			6893	7189	7275	7339
		8471	8490	8492	8501			7475	7832	8024	8315
HinC II	(2)	1765	2768			Msc I	(4)	285	1509	4233	7506
Hind II	(2)	1765	2768			Mse I	(37)	5	9	371	1703
Hinf I	(22)	774	1290	1939	2097			1764	1885	1903	2126
		2350	2791	2911	3044			2205	2218	2298	2810
		3332	3888	4440	4984			2889	3241	3826	3874
		4998	5020	5734	5809			4641	4844	4947	5122
		6205	6722	7909	8043			5261	5364	5487	5641
		8214	8236					5645	6540	6592	6597
								6611	6664	6899	6938

pAdTrace-OK Vector

		8097	8109	8126	8224
		8494			
Msl I	(6)	729	2073	3469	3534
		4728	7861		
Msp I	(39)	203	1531	1630	1644
		2537	2802	2982	3327
		3367	3617	3719	3781
		3809	4218	4277	4444
		4583	4754	4914	4981
		5057	6041	6188	6214
		6404	6808	6842	6909
		7031	7158	7329	7406
		7428	7456	7587	7677
		7744	7925	8389	
MspAl I	(14)	1526	2671	2687	2829
		2991	3737	3764	3840
		4341	5217	6176	6421
		7170	7530		
Mun I	(2)	1752	2785		
Nae I	(3)	4915	7926	8390	
Nar I	(7)	1426	1534	1633	2024
		4380	7176	7424	
Nci I	(18)	1531	2538	2802	2803
		2982	2983	3328	3720
		3810	4219	4445	5057
		5058	6214	6910	7031
		7428	7588		
Nco I	(6)	724	961	1381	2530
		4315	7856		
Nde I	(2)	598	2190		
NgoM I	(3)	4913	7924	8388	
Nla III	(39)	162	668	728	965
		998	1016	1385	1508
		2044	2357	2472	2526
		2534	3066	3116	3339
		3392	3456	3687	3882
		3897	4002	4157	4319
		4546	4580	4727	4733
		4797	4913	4979	5838
		6558	7117	7298	7643
		7829	7860	7886	
Nla IV	(41)	361	831	1022	1082
		1118	1166	1340	1361
		1427	1535	1634	1895
		1948	2025	2562	2986
		3010	3169	3482	3590
		3611	3754	3788	3864
		4036	4176	4381	4471
		4758	5285	5866	5905
		6677	6771	6812	7177
		7425	7460	8326	8347
		8359			
Nsi I	(2)	3280	3458		
Nsp7524 I	(9)	158	2040	2522	3878
		3893	3998	4793	5834
		7825			
NspB II	(14)	1526	2671	2687	2829
		2991	3737	3764	3840
		4341	5217	6176	6421
		7170	7530		
NspH I	(9)	162	2044	2526	3882
		3897	4002	4797	5838
		7829			
Pac I	(2)	7	5643		
Pal I	(44)	285	448	641	966
		1033	1066	1117	1173
		1360	1395	1474	1509
		1618	1948	2366	2536
		2980	3213	3482	3496
		3562	3620	3788	3865
		4146	4233	4299	4349
		4470	4513	4908	5317
		5849	5860	5878	6312
		6770	6850	7332	7506
		7897	7924	8137	8279
PflM I	(2)	1387	3032		
Ple I	(3)	4992	6213	8222	
Pme I	(1)	4642			
PpuM I	(2)	3009	4175		
Psp1406 I	(1)	6953			
PspA I	(3)	2801	2981	5056	
Pst I	(5)	1303	2649	2676	3106
		3839			
Pvu II	(7)	1526	2829	3737	3764
		3840	7170	7530	
Rsa I	(14)	209	361	583	608
		663	696	747	904
		1161	1413	1602	2270
		4116	7730		
Rsr II	(1)	7940			
Sac I	(2)	2506	4509		
Sac II	(2)	2688	4342		
Sap I	(3)	5718	7768	7978	
Sau3A I	(35)	949	1045	1441	1646
		1659	1893	2387	2835
		2927	3137	3517	3550
		3680	3731	4069	5649
		5653	6400	6475	6486
		6494	6572	6584	6689
		6973	6985	7237	7254
		7265	7284	7595	7673
		7754	7763	7841	
Sau96 I	(32)	447	640	1064	1116

		1359	1616	1946	1947
		2365	2534	2979	2985
		3009	3212	3480	3481
		3619	3786	3787	3863
		4034	4144	4175	4253
		4348	4469	5315	6769
		6848	6865	7940	8278
ScrF I	(38)	454	647	1133	1287
		1531	1989	2538	2802
		2803	2982	2983	3027
		3204	3328	3498	3543
		3623	3720	3810	4029
		4148	4219	4235	4301
		4445	4500	4532	5057
		5058	5862	5983	5996
		6214	6910	7031	7428
		7588	7811		
Sec I	(45)	724	961	973	1108
		1132	1168	1225	1285
		1381	1417	1486	1612
		1934	1987	2383	2530
		2537	2685	2801	2981
		3203	3215	3327	3484
		3497	3621	3622	3719
		4147	4163	4234	4315
		4339	4351	4369	4444
		4969	5056	5057	5166
		5994	7029	7233	7587
		7856			
SfaN I	(16)	722	1817	3266	3286
		4071	4463	4544	5672
		5710	5930	7383	7638
		7722	7786	7854	8061
Sfc I	(7)	1299	2645	2672	3102
		3835	6099	6290	
Sma I	(3)	2803	2983	5058	
SnaB I	(2)	704	5473		
Sph I	(3)	2526	4002	7829	
Ssp I	(1)	2035			
Stu I	(2)	1173	1395		
Sty I	(15)	724	961	1108	1168
		1381	1934	2383	2530
		3215	4315	4351	4369
		5166	7233	7856	
Swa I	(1)	2127			
Taq I	(15)	1042	1048	1258	1908
		2163	4068	5276	5652
		5934	7537	7693	7717
		7753	7915	8320	
Tfi I	(10)	1939	2097	2350	2791
		3332	4440	5020	5809
		7909	8043		
Tsp45 I	(24)	77	109	183	218
		1103	1274	1280	1954
		2256	2694	3041	3434
		4047	4196	4598	4918
		4936	5298	5449	5539
		5571	7544	7850	8460
Tth111 I	(2)	2622	7542		
Tth111 II	(9)	2741	2907	2946	4034
		4805	6423	6432	6462
		7860			
Vsp I	(6)	5	371	2298	5122
		5641	6899		
Xba I	(1)	1649			
Xca I	(1)	4627			
Xcm I	(3)	2915	4506	7240	
Xho II	(14)	949	1646	1893	2387
		2835	6475	6486	6572
		6584	6973	6985	7237
		7595	7841		
Xma I	(3)	2801	2981	5056	
Xma III	(1)	7330			
Xmn I	(1)	8508			

Site usage in pAdTrace-OK:

Aat II	G,ACGT`C	6	Acc I	GT`MK,AC	2
Acc65 I	G`GTAC,C	1	Aci I	C`CG,C	112
Afl II	C`TTAA,G	-	Afl III	A`CRYG,T	3
Age I	A`CCGG,T	-	Aha II	GR`CG,YC	15
Ahd I	GACNN,N`NNGTC	3	Alu I	AG CT	44
Alw I	GGATC 8/9	26	AlwN I	CAG,NNN`CTG	6
Apa I	G,GGCC`C	3	Apal I	G`TGCA,C	2
Apo I	R`AATT,Y	7	Asc I	GG`CGCG,CC	-
Ase I	AT`TA,AT	6	Asp718	G`GTAC,C	1
Ava I	C`YCGR,G	5	Ava II	G`GWC,C	7
Avr II	C`CTAG,G	1	BamH I	G`GATC,C	1
Ban I	G`GYRC,C	17	Ban II	G,RCGY`C	10
Bbe I	G,CGCC`C	7	Bbs I	GAAGAC 8/12	2
Bbv I	GCAGC 13/17	19	Bbv II	GAAGAC 7/11	2
Bcl I	T`GATC,A	2	Bcn I	CC,S`GG	18
Bfa I	C`TA,G	12	Bgl I	GCCN,NNN`NGGC	7
Bgl II	A`GATC,T	-	Blp I	GC`TNA,GC	-
Bpm I	CTGGAG 22/20	3	Bsa I	GGTCTC 7/11	6
BsaA I	YAC GTR	8	BsaB I	GATNN NNATC	3
BsaH I	GR`CG,YC	15	BsaJ I	C`C`NNG,G	45
BsaW I	W`CCGG,W	8	BseR I	GAGGAG 16/14	5
Bsg I	GTGCAG 22/20	5	BsiC I	TT`CG,AA	1
BsiE I	CG,RY`CG	6	BsiHKA I	G,WGCM`C	8
BsiW I	C`GTAC,G	-	Bsm I	GAATG,C 7	4
BsmA I	GTCTC`/9	12	BsmB I	CGTCTC 7/11	2
BsmF I	GGGAC 15/19	11	BsoF I	GC`N,GC	69

Bsp120 I	G`GGCC,C	3	Bsp1286 I	G,DGCH`C	2
BspH I	T`CATG,A	1	BspM I	ACCTGC 10/14	8
BspM II	T`CCGG,A	2	Bsr I	ACT,GG`	20
BsrB I	GAG CGG	6	BsrD I	GCAATG, 8	3
BsrG I	T`GTAC,A	1	BssH II	G`CGCG,C	7
BssS I	C`TCGT,G	5	Bst1107 I	GTA TAC	1
BstB I	TT`CG,AA	1	BstE II	G`GTNAC,C	4
BstN I	CC`W,GG	20	BstU I	CG CG	39
BstX I	CCAN,NNNN`NTGG	2	BstY I	R`GATC,Y	14
Bsu36 I	CC`TNA,GG	-	Cac8 I	GCN NGC	52
Cfr10 I	R`CCGG,Y	10	Cla I	AT`CG,AT	1
Csp6 I	G`TA,C	14	Dde I	C`TNA,G	9
Dpn I	GA TC	35	DpnII	`GATC,	35
Dra I	TTT AAA	9	Dra III	CAC,NNN`GTG	1
Drd I	GACNN,NN`NNGTC	5	Dsa I	C`CRYG,G	10
Eae I	Y`GGCC,R	12	Eag I	C`GGCC,C	1
Ear I	CTCTTC 7/10	4	Eco47 III	AGC GCT	3
Eco57 I	CTGAAG 21/19	10	Eco72 I	CAC GTG	-
EcoN I	CCTNN`N,NNAGG	-	Eco109 I	RG`GNC,CY	5
EcoR I	G`AATT,C	1	EcoR II	`CCWGG,	20
EcoR V	GAT ATC	-	Ehe I	GGC GCC	7
Esp I	GC`TNA,GC	-	Fnu4H I	GC`N,GC	69
Fok I	GGATG 14/18	20	Fse I	GG,CCGG`CC	-
Fsp I	TGC GCA	3	Gdi II	`YGGC,CG	9
Gsu I	CTGGAG 21/19	3	Hae I	WGG CCW	11
Hae II	R,GCGC`Y	15	Hae III	GG CC	44
Hga I	GACGC 9/14	7	HgiA I	G,WGCW`C	8
HgiE II	ACCNNNNNNGGT -1/132	-	Hha I	G,CG`C	64
Hinc II	GTY CAC	2	Hind II	GTY CAC	2
Hind III	A`AGCT,T	-	Hinf I	G`ANT,C	22
HinI I	GR`CG,YC	15	HinP I	G`CG,C	64
Hpa I	GTT AAC	1	Hpa II	C`CG,G	39
Hph I	GGTGA 12/11	19	Kas I	G`GCGC,C	7
Kpn I	G,GTAC`C	1	Mae I	C`TA,G	12
Mae II	A`CG,T	32	Mae III	`GTNAC,	41
Mbo I	`GATC,	35	Mbo III	GAAAGA 12/11	17
Mlu I	A`CGCG,T	-	Mme I	TCCRAC 25/23	6
Mnl I	CCTC 10/10	68	Msc I	TGG CCA	4
Mse I	T`TA,A	37	Msl I	CAYNN NNRTG	6
Msp I	C`CG,G	39	Msp1 I	CMG CKG	14
Mun I	C`AATT,G	2	Nae I	GCC GGC	3
Nar I	GG`CG,CC	7	Nci I	CC`S,GG	18
Nco I	C`CATG,G	6	Nde I	CA`TA,TG	2
NgoM I	G`CCGG,C	3	Nhe I	G`CTAG,C	-
Nla III	,CATG`	39	Nla IV	GGN NCC	41
Not I	GC`GGCC,GC	-	Nru I	TCG CGA	-
Nsi I	A,TGCA`T	2	Nsp7524 I	R`CATG,Y	9
NspB II	CMG CRG	14	NspH I	R,CATG`Y	9
Pac I	TTA`AT,TAA	2	Paer7 I	C`TCGA,G	-
Pal I	GG CC	44	PflM I	CCAN,NNN`NTGG	2
Ple I	GAGTC 9/10	3	Pme I	CTTT AAAC	1
Pml I	CAC GTG	-	PpuM I	RG`GWC,CY	2
Psp1406 I	AA`CG,TT	1	PspA I	C`CCGG,C	3
Pst I	C,TGCA`G	5	Pvu I	CG,AT`CG	-
Pvu II	CAG CTG	7	Rsa I	GT AC	14
Rsr II	CG`GWC,CG	1	Sac I	G,AGCT`C	2
Rsc II	CC,GC`GG	2	Sal I	G`TCGA,C	-
Sap I	GCTCTTC 8/11	3	Sau3A I	`GATC,	35
Sau96 I	G`GNC,C	32	Sca I	AGT ACT	-
ScaF I	CC`N,GG	38	Sec I	C`CNNG,G	45
SfaN I	GCATC 9/13	16	Sfc I	C`TRVA,G	7
Sfi I	GGCCN,NNN`NGGCC	-	Sma I	CCC GGG	3
SnaB I	TAC GTA	2	Spe I	A`CTAG,T	-
Sph I	G,CATG`C	3	Spl I	C`GTAC,G	-
Srf I	GCCC GGGC	-	Ssp I	AAT ATT	1
Stu I	AGG CCT	2	Sty I	C`CWGG,G	15
Swa I	ATTT AAAT	1	Taq I	T`CG,A	15
Tfi I	G`AWT,C	10	Tsp45 I	`GTSAC,	24
Tth111 I	GACN`N,NGTC	2	Tth111 II	CAARCA 16/14	9
Vsp I	AT`TA,AT	6	Xba I	T`CTAG,A	1
Xca I	GTA TAC	1	Xcm I	CCANNNN,N`NNNTGG3	-
Xho I	C`TCGA,G	-	Xho II	R`GATC,Y	14
Xma I	C`CCGG,G	3	Xma III	C`GGCC,G	-
BsrG I	T`GTAC,A	207			
Acc65 I	G`GTAC,C	359			
Asp718	G`GTAC,C	359			
Kpn I	G,GTAC`C	363			
Hpa I	GTT AAC	1765			
Mlu I	A`CGCG,T	1888			
BsiC I	TT`CG,AA	1933			
BstB I	TT`CG,AA	1933			
Ssp I	AAT ATT	2060			
Swa I	ATTT AAAT	2152			
Bst1107 I	GTA TAC	4652			
Xca I	GTA TAC	4652			
Pme I	CTTT AAAC	4667			
Avr II	C`CTAG,G	5191			
Cla I	AT`CG,AT	5677			
BspH I	T`CATG,A	6579			
Psp1406 I	AA`CG,TT	6978			
Eag I	C`GGCC,G	7355			
Xma III	C`GGCC,G	7355			
Rsr II	CG`GWC,CG	7965			
Dra III	CAC,NNN`GTG	8312			
Xmn I	GAANN NNTTC	8533			

The following enzymes do not cut in pAdTrace-OK:

Afl II	Age I	Asc I	BamH I	Bgl II
Blp I	BsiW I	Bsu36 I	Eco72 I	EcoN I
EcoR V	Esp I	Fse I	HinD III	Nhe I
Not I	Nru I	Paer7 I	Pml I	Pvu I
Sal I	Sca I	Sfi I	Spe I	Spl I

pAdTrace-OK: sites sorted by name:

Aat II	(6)	489	542	625	811
		982	1946		
Acc I	(2)	3713	4651		
Acc65 I	(1)	359			
AcI I	(111)	88	106	126	151
		233	243	294	422
		450	462	476	643
		734	767	871	892
		953	1067	1090	1114
		1189	1267	1336	1406
		1471	1619	1954	1969
		1988	2484	2487	2490
		2616	2679	2682	2685
		2694	2706	2710	2712
		2744	2775	2779	2909
		2998	3016	3144	3262
		3515	3518	3603	3607
		3624	3720	3800	3939
		3956	4014	4057	4118
		4128	4364	4366	4408
		4484	4489	4498	4826
		5098	5212	5242	5417
		5549	5570	5588	5699
		5725	5742	5785	5792
		5813	5904	5932	6059
		6078	6199	6309	6444
		6453	6815	6906	7036
		7159	7255	7358	7421
		7515	7579	7680	7683
		7923	7963	7968	8018
		8034	8060	8202	8341
		8456	8483	8511	
Afl III	(4)	158	1888	4818	5859
Aha II	(15)	486	539	622	808
		979	1426	1483	1534
		1633	1943	2049	3452
		4405	7201	7449	
Ahd I	(3)	1142	3016	6752	
Alu I	(45)	25	930	1097	1214
		1331	1460	1526	1556
		1781	1917	1926	2529
		2762	2854	3134	3220
		3270	3420	3762	3789
		3859	3865	4068	4388
		4445	4532	4776	4991
		5315	5801	6027	6117
		6163	6420	6941	7070
		7123	7149	7195	7286
		7555	8013	8142	8360
Alw I	(24)	945	1653	2419	2867
		2959	3158	3549	3712
		4090	5674	5681	6421
		6507	6507	6604	6605
		7005	7006	7269	7286
		7316	7627	7694	7873
AlwN I	(6)	2696	2854	2939	4660
		6275	7073		
Apa I	(3)	1975	3509	3815	
ApaL I	(2)	3395	6173		
Apo I	(8)	257	1816	1909	2153
		3406	4672	8114	8125
		5	371	2323	5147
Ase I	(6)	5666	6924		
Asp718	(1)	359			
Ava I	(5)	1224	1416	2826	3006
		5081			
Ava II	(7)	3010	3034	4059	4200
		4278	6890	7965	
Avr II	(1)	5191			
Ban I	(17)	359	829	1080	1338
		1425	1533	1632	2048
		3192	3613	4404	5308
		6700	7200	7448	7483
		8349			
Ban II	(10)	1023	1975	2531	2588
		2744	3509	3815	4534
		7814	8387		
Bbe I	(7)	1429	1537	1636	2052
		4408	7204	7452	
Bbs I	(2)	1383	4298		
Bbv I	(19)	1535	1790	2489	2492
		2684	2711	2771	2863
		3798	3814	3874	4397
		5257	6278	6281	6487
		7522	7564	8085	
Bbv II	(2)	1382	4299		
Bcl I	(2)	1659	7290		
Bcn I	(18)	1532	2564	2828	2829
		3008	3009	3354	3746
		3836	4245	4471	5083
		5084	6240	6936	7057

Number of enzymes = 22

pAdTrace-OK Vector

Bfa I	(13)	7454 364 1898 6607 8463	7614 956 1984 6942	1650 5192 7008	1851 6354 7150	BstX I	(2)	1505 3496				
Bgl I	(7)	454 4939	576 6872	647 8532	2888	BstY I	(13)	949 6500 6511 6597 6609 6998 7010 7262 7620 7866	1646 2412 2860			
Bpm I	(3)	2572	5200	6822		Cac8 I	(52)	450 2003 2710 2744 2752	643 2007 2549 2616	1609 2748 2752	1971 2616	
Bsa I	(6)	2136 4512	2636 6813	3024	4000			2779 3218 3236 3605 3953	2870 2994 2996	2994 3605 3953	2996 3953	
BsaA I	(8)	704 5498	1178 7022	2286 7753	3287 8312			4025 4390 4493 4940 5102 5421 5790 5876 5913	4252 4493 4940 5102	4256 4940 5102	4383 5102 5913	
BsaB I	(3)	1664	2951	7308				6473 7176 7193 7343 7529 7748 7814 7820 7848 7852 7893 7897 7951 8415 8458 8472 8515	6864 7101 7125	7101 7125	7125 7529 7848 7951 8515	
BsaH I	(15)	486 979 1633 4405	539 1426 1483 1534	622 808				7748 7852 7893 7897 7951 8415 8458 8472 8515	7101 7125	7101 7125	7125 7529 7848 7951 8515	
BsaJ I	(45)	724 1132 1381 1417 1599 2562 2710 2826 3006 3228 3240 3352 3509 3522 3646 3647 3744 4172 4188 4259 4340 4364 4376 4394 4469 4994 5081 5082 5191 6019 7054 7258 7612	961 1168 1225 1285	973 1108					202 1629 3805 4301	202 1629 3805 4301	202 1629 3805 4301	202 1629 3805 4301
BsaW I	(8)	1643 6065	3391 6212	4778 7182	5005 7480			202 4607 7949	1629 3805 4301	202 1629 3805 4301	202 1629 3805 4301	
BseR I	(5)	959 4983	1283	4098	4414			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
Bsg I	(5)	1542 5368	2778	3388	4443			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsiC I	(1)	1933						951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsiE I	(6)	3026 7358	5072 8345	5775	6199			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsiHKA I	(8)	2501 4534	2531 6177	3399 7562	4159 7752			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
Bsm I	(4)	1753	1846	3481	4512			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsmA I	(12)	797 3023 4511	2137 3107 5104	2637 3444 6814	2661 4001 7292			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsmB I	(2)	3108	3443					951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsmF I	(11)	539 1149 4045	690 2049 4629	858 3023 7600	1130 3892			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsoF I	(69)	1067 1337 1619 2484 2487 2682 2700 2760 3516 3803 3956 4367 4697 5783 6202 6267 6270 6476 6804 7421 7511 7516 7553 7594 7681 7684 7687 7923 8019 8060 8074 8492	1115 1472 1524 1527	1268 1332					951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648
Bsp120 I	(3)	1971	3505	3811				951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
Bsp1286 I	(2)	2501	7752					951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BspH I	(1)	6579						951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BspM I	(8)	2863 4759	3725 7107	3765 7336	4085 7717			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BspM II	(2)	4778	5005					951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
Bsr I	(20)	660 3847 6279 6960 7242	1158 4962 6393 7032 7392	1976 5267 6266 6917 7158 8226	3167 6266 6917 7158 8226			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsrB I	(6)	1406 8062	1956 8456	5214	5792			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsrD I	(3)	3507	6813	7682				951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BsrG I	(1)	207						951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BssH II	(7)	1607 4250	2001 4381	2992 8513	2994			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BssS I	(5)	1035 8041	4411	4518	6032			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
Bst1107 I	(1)	4652						951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BstB I	(1)	1933						951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BstE II	(4)	1103	1274	4072	4961			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BstN I	(20)	454 2014 3568 4260 5887	647 3052 3648 4326 6008	1133 3229 4054 4525 6021	1287 3523 4173 4557 7836			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
BstU I	(39)	88 350 1609 2005 2994 4118 4383 5423 6817 8496	231 424 1890 2545 2998 4252 4453 5590 7515 8513	233 1246 1969 2712 2998 4318 4486 5906 7816 8515	294 1336 2003 2781 3603 4366 4723 6487 8204			951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	951 1047 1443 1648	
								184 1421 3583 6718 7316	258 2758 3088 3859 4467	1125 3088 3859 4467	1182 3287 4467	
								1011 1032 4932 7946	6974 1473 7354 7356	7551 4323 7356	7551 4535 7919	
								2573	5201	6823		

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Hae I	(11)	285	966	1173	1395	2719	3066	3459	3657		
		1509	3587	4258	5874	4072	4221	4623	4943		
		5885	6337	7531		4961	5323	5381	5474		
Hae II	(15)	1429	1537	1636	2052	5493	5498	5542	5564		
		3184	3952	4299	4408	5596	6215	6278	6394		
		5235	5737	6107	7204	6677	7569	7875	8485		
		7452	8463	8471		8497					
Hae III	(44)	285	448	641	966	Mbo I	(34)	949	1045	1441	1646
		1033	1066	1117	1173			1659	2412	2860	2952
		1360	1395	1474	1509			3162	3542	3575	3705
		1618	1973	2391	2561			3756	4094	5674	5678
		3005	3238	3507	3521			6425	6500	6511	6519
		3587	3645	3813	3890			6597	6609	6714	6998
		4171	4258	4324	4374			7010	7262	7279	7290
		4495	4538	4933	5342			7309	7620	7698	7779
		5874	5885	5903	6337			7788	7866		
		6795	6875	7357	7531	Mbo II	(17)	1384	1387	2032	2388
		7922	7949	8162	8304			3450	3540	3741	3778
Hga I	(7)	889	1490	2559	3459			4299	5731	6520	6593
		4308	5969	6547				6994	7809	8019	8101
								8451			
HgiA I	(8)	2501	2531	3399	4159	Mlu I	(1)	1888			
		4534	6177	7562	7752	Mme I	(6)	1377	2644	2839	6073
HgiE II	(2)	3797	6438					6257	8288		
Hha I	(64)	88	202	233	350	Mnl I	(68)	56	304	905	968
		1000	1012	1246	1336			976	979	1010	1043
		1428	1536	1609	1611			1049	1055	1070	1158
		1635	2003	2005	2011			1220	1253	1303	1360
		2051	2547	2868	2994			1385	1405	1412	1444
		2996	2998	3183	3603			1481	1571	1576	1607
		3951	4196	4240	4252			1681	1720	1729	2026
		4254	4268	4298	4318			2402	2685	2904	2909
		4383	4385	4407	4453			3024	3108	3358	3473
		4488	4725	5234	5592			3603	3672	4118	4212
		5736	5769	6039	6106			4265	4370	4392	4395
		6206	6380	6489	6882			4407	4427	4600	4728
		6975	7105	7203	7274			5003	5126	5193	5217
		7443	7451	7515	7552			5475	5622	5757	5967
		7818	8078	8462	8470			6040	6291	6691	6772
		8496	8515	8517	8526			6918	7214	7300	7364
HinC II	(2)	1765	2793					7500	7857	8049	8340
Hind II	(2)	1765	2793			Msc I	(4)	285	1509	4258	7531
Hinf I	(22)	774	1290	1964	2122	Mse I	(37)	5	9	371	1703
		2375	2816	2936	3069			1764	1885	1928	2151
		3357	3913	4465	5009			2230	2243	2323	2835
		5023	5045	5759	5834			2914	3266	3851	3899
		6230	6747	7934	8068			4666	4869	4972	5147
		8239	8261					5286	5389	5512	5666
HinI I	(15)	486	539	622	808			5670	6565	6617	6622
		979	1426	1483	1534			6636	6689	6924	6963
		1633	1943	2049	3452			8122	8134	8151	8249
		4405	7201	7449				8519			
HinP I	(64)	86	200	231	348	Msl I	(6)	729	2098	3494	3559
		998	1010	1244	1334			4753	7886		
		1426	1534	1607	1609	Msp I	(39)	203	1531	1630	1644
		1633	2001	2003	2009			2562	2827	3007	3352
		2049	2545	2866	2992			3392	3642	3744	3806
		2994	2996	3181	3601			3834	4243	4302	4469
		3949	4194	4238	4250			4608	4779	4939	5006
		4252	4266	4296	4316			5082	6066	6213	6239
		4381	4383	4405	4451			6429	6833	6867	6934
		4486	4723	5232	5590			7056	7183	7354	7431
		5734	5767	6037	6104			7453	7481	7612	7702
		6204	6378	6487	6880			7769	7950	8414	
		6973	7103	7201	7272	MspAI I	(14)	1526	2696	2712	2854
		7441	7449	7513	7550			3016	3762	3789	3865
		7816	8076	8460	8468			4366	5242	6201	6446
		8494	8513	8515	8524			7195	7555		
Hpa I	(1)	1765				Mun I	(2)	1752	2810		
Hpa II	(39)	203	1531	1630	1644	Nae I	(3)	4940	7951	8415	
		2562	2827	3007	3352	Nar I	(7)	1426	1534	1633	2049
		3392	3642	3744	3806			4405	7201	7449	
		3834	4243	4302	4469	Nci I	(18)	1531	2563	2827	2828
		4608	4779	4939	5006			3007	3008	3353	3745
		5082	6066	6213	6239			3835	4244	4470	5082
		6429	6833	6867	6934			5083	6239	6935	7056
		7056	7183	7354	7431			7453	7613		
		7453	7481	7612	7702	Nco I	(6)	724	961	1381	2555
		7769	7950	8414				4340	7881		
Hph I	(19)	120	739	1114	1285	Nde I	(2)	598	2215		
		1336	1557	2083	3077	NgoM I	(3)	4938	7949	8413	
		4067	4238	4538	4634	Nla III	(39)	162	668	728	965
		4956	5559	6596	6823			998	1016	1385	1508
		6997	7628	8314				2069	2382	2497	2551
Kas I	(7)	1425	1533	1632	2048			2559	3091	3141	3364
		4404	7200	7448				3417	3481	3712	3907
Kpn I	(1)	363						3922	4027	4182	4344
Mae I	(13)	364	956	1650	1851			4571	4605	4752	4758
		1898	1984	5192	6354			4822	4938	5004	5863
		6607	6942	7008	7150			6583	7142	7323	7668
		8463						7854	7885	7911	
Mae II	(33)	81	113	187	486	Nla IV	(40)	361	831	1022	1082
		498	539	622	703			1118	1166	1340	1361
		808	979	1177	1902			1427	1535	1634	1973
		1943	2099	2280	2285			2050	2587	3011	3035
		3056	3286	3551	5380			3194	3507	3615	3636
		5492	5497	5563	5595			3779	3813	3889	4061
		6562	6978	7021	7565			4201	4406	4496	4783
		7752	8256	8268	8311			5310	5891	5930	6702
		8421						6796	6837	7202	7450
Mae III	(41)	77	109	183	218			7485	8351	8372	8384
		266	337	425	512	Nsi I	(2)	3305	3483		
		861	1103	1274	1280	Nsp7524 I	(9)	158	2065	2547	3903
		1790	1979	2202	2281						

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		3918	4023	4818	5859
		7850			
NspB II	(14)	1526	2696	2712	2854
		3016	3762	3789	3865
		4366	5242	6201	6446
		7195	7555		
NspH I	(9)	162	2069	2551	3907
		3922	4027	4822	5863
		7854			
PacI	(2)	7	5668		
Pal I	(44)	285	448	641	966
		1033	1066	1117	1173
		1360	1395	1474	1509
		1618	1973	2391	2561
		3005	3238	3507	3521
		3587	3645	3813	3890
		4171	4258	4324	4374
		4495	4538	4933	5342
		5874	5885	5903	6337
		6795	6875	7357	7531
		7922	7949	8162	8304
PflM I	(2)	1387	3057		
Ple I	(3)	5017	6238	8247	
Pme I	(1)	4667			
PpuM I	(2)	3034	4200		
Psp1406 I	(1)	6978			
PspA I	(3)	2826	3006	5081	
Pst I	(5)	1303	2674	2701	3131
		3864			
Pvu II	(7)	1526	2854	3762	3789
		3865	7195	7555	
Rsa I	(14)	209	361	583	608
		663	696	747	904
		1161	1413	1602	2295
		4141	7755		
Rsr II	(1)	7965			
Sac I	(2)	2531	4534		
Sac II	(2)	2713	4367		
Sap I	(3)	5743	7793	8003	
Sau3A I	(34)	949	1045	1441	1646
		1659	2412	2860	2952
		3162	3542	3575	3705
		3756	4094	5674	5678
		6425	6500	6511	6519
		6597	6609	6714	6998
		7010	7262	7279	7290
		7309	7620	7698	7779
		7788	7866		
Sau96 I	(32)	447	640	1064	1116
		1359	1616	1971	1972
		2390	2559	3004	3010
		3034	3237	3505	3506
		3644	3811	3812	3888
		4059	4169	4200	4278
		4373	4494	5340	6794
		6873	6890	7965	8303
ScrF I	(38)	454	647	1133	1287
		1531	2014	2563	2827
		2828	3007	3008	3052
		3229	3353	3523	3568
		3648	3745	3835	4054
		4173	4244	4260	4326
		4470	4525	4557	5082
		5083	5887	6008	6021
		6239	6935	7056	7453
		7613	7836		
Sec I	(45)	724	961	973	1108
		1132	1168	1225	1285
		1381	1417	1486	1612
		1959	2012	2408	2555
		2562	2710	2826	3006
		3228	3240	3352	3509
		3522	3646	3647	3744
		4172	4188	4259	4340
		4364	4376	4394	4469
		4994	5081	5082	5191
		6019	7054	7258	7612
		7881			
SfaN I	(16)	722	1817	3291	3311
		4096	4488	4569	5697
		5735	5955	7408	7663
		7747	7811	7879	8086
Sfc I	(7)	1299	2670	2697	3127
		3860	6124	6315	
Sma I	(3)	2828	3008	5083	
SnaB I	(2)	704	5498		
Sph I	(3)	2551	4027	7854	
Ssp I	(1)	2060			
Stu I	(2)	1173	1395		
Sty I	(15)	724	961	1108	1168
		1381	1959	2408	2555
		3240	4340	4376	4394
		5191	7258	7881	
SwaI	(1)	2152			
Taq I	(16)	1042	1048	1258	1907
		1933	2188	4093	5301
		5677	5959	7562	7718
		7742	7778	7940	8345
Tfi I	(10)	1964	2122	2375	2816
		3357	4465	5045	5834
		7934	8068		
Tsp45 I	(24)	77	109	183	218
		1103	1274	1280	1979

		2281	2719	3066	3459
		4072	4221	4623	4943
		4961	5323	5474	5564
		5596	7569	7875	8485
Tth111 I	(2)	2647	7567		
Tth111 II	(9)	2766	2932	2971	4059
		4830	6448	6457	6487
		7885			
Vsp I	(6)	5	371	2323	5147
		5666	6924		
Xba I	(2)	1649	1897		
Xca I	(1)	4652			
Xcm I	(3)	2940	4531	7265	
Xho II	(13)	949	1646	2412	2860
		6500	6511	6597	6609
		6998	7010	7262	7620
		7866			
Xma I	(3)	2826	3006	5081	
Xma III	(1)	7355			
Xmn I	(1)	8533			

Site usage in pAdTrace-OK:

Aat II	G,ACGT`C	6	Acc I	GT`MK,AC	2
Acc65 I	G`GTAC,C	1	Aci I	C`CG,C	111
Afl II	C`TTAA,G	-	Afl III	A`CRYG,T	4
Age I	A`CCGG,T	-	Aha II	GR`CG,YC	15
Ahd I	GACNN,N`NNGTC	3	Alu I	AG CT	45
Alw I	GGATC 8/9	24	AlwN I	CAG,NNN`CTG	6
Apa I	G,GGCC`C	3	ApaL I	G`TGCA,C	2
Apo I	R`AATT,Y	8	Asc I	GG`CGCG,CC	-
Ase I	AT`TA,AT	6	Asp718	G`GTAC,C	1
Ava I	C`YCGR,G	5	Ava II	G`GWC,C	7
Avr II	C`CTAG,G	1	BamH I	G`GATC,C	-
Ban I	G`GYRC,C	17	Ban II	G,RCGY`C	10
Bbe I	G,CGCC`C	7	Bbs I	GAAGAC 8/12	2
Bbv I	GCAGC 13/17	19	Bbv II	GAAGAC 7/11	2
Bcl I	T`GATC,A	2	Bcn I	CC,S`GG	18
Bfa I	C`TA,G	13	Bgl I	GCCN,NNN`NGGC	7
Bgl II	A`GATC,T	-	Blp I	GC`TNA,GC	-
Bpm I	CTGGAG 22/20	3	Bsa I	GGTCTC 7/11	6
BsaA I	YAC GTR	8	BsaB I	GATNN NNATC	3
BsaH I	GR`CG,YC	15	BsaJ I	C`CNNG,G	45
BsaW I	W`CCGG,W	8	BseR I	GAGGAG 16/14	5
Bsg I	GTGCG 22/20	5	BsiC I	TT`CG,AA	1
BsiE I	CG,RY`CG	6	BsiHKA I	G,WGCW`C	8
BsiW I	C`GTAC,G	-	Bsm I	GAATG,C 7	4
BsmA I	GTCTC`/9	12	BsmB I	CGTCTC 7/11	2
BsmF I	GGGAC 15/19	11	BsoF I	GC`N,GC	69
Bsp120 I	G`GGCC,C	3	Bsp1286 I	G,DGCH`C	2
BspH I	T`CATG,A	1	BspM I	ACCTGC 10/14	8
BspM II	T`CCGG,A	2	Bsr I	ACT,GG`	20
BsrE I	GAG CGG	6	Bsrd I	GCAATG, 8	3
BsrG I	T`GTAC,A	1	Bssh II	G`CGCG,C	7
BssS I	C`TCGT,G	5	Bst1107 I	GTA TAC	1
BstB I	TT`CG,AA	1	BstE II	G`GTNAC,C	4
BstN I	CC`W,GG	20	BstU I	CG CG	39
BstX I	CCAN,NNNN`NTGG	2	BstY I	R`GATC,Y	13
Bsu36 I	CC`TNA,GG	-	Cac8 I	GCN NGC	52
Cfr10 I	R`CCGG,Y	10	Cla I	AT`CG,AT	1
Csp6 I	G`TA,C	14	Dde I	C`TNA,G	9
Dpn I	GA TC	34	DpnII	`GATC,	34
Dra I	TTT AAA	9	Dra III	CAC,NNN`GTG	1
Drd I	GACNN,NN`NNGTC	5	Dsa I	C`CRYG,G	10
Eae I	Y`GGCC,R	12	Eag I	C`GGCC,G	1
Ear I	CTCTTC 7/10	4	Eco47 III	AGC GCT	3
Eco57 I	CTGAAG 21/19	10	Eco72 I	CAC GTG	-
EcoN I	CCTNN`N,NNAGG	-	Eco1019 I	RG`GNC,CY	5
EcoR I	G`AATT,C	2	EcoR II	`CCWGG,	20
EcoR V	GAT ATC	-	Ehe I	GGC GCC	7
Esp I	GC`TNA,GC	-	Fnu4H I	GC`N,GC	69
Fok I	GGATG 14/18	20	Fse I	GG,CCGG`CC	-
Fsp I	TGC GCA	3	Gdi II	`YGGC,CG	9
Gsu I	CTGGAG 21/19	3	Hae I	WGG CCW	11
Hae II	R,CGCC`Y	15	Hae III	GG CC	44
Hga I	GACGC 9/14	7	HgiA I	G,WGCW`C	8
HgiE II	ACNNNNNNGGT	-1/132	Hha I	G,CG`C	64
HinC II	GTY RAC	2	Hind II	GTY RAC	2
Hind III	A`AGCT,T	-	Hinf I	G`ANT,C	22
HinI I	GR`CG,YC	15	HinP I	G`CG,C	64
Hpa I	GTT AAC	1	Hpa II	C`CG,G	39
Hph I	GGTGA 12/11	19	Kas I	G`CGCG,C	7
Kpn I	G,GTAC`C	1	Mae I	C`TA,G	13
Mae II	A`CG,T	33	Mae III	`GTNAC,	41
Mbo I	`GATC,	34	Mbo II	GAAGA 12/11	17
Mlu I	A`CGCG,T	1	Mme I	TCCRAC 25/23	6
Mnl I	CCTC 10/10	68	Msc I	TGG CCA	4
Mse I	T`TA,A	37	Msl I	CAYNN NNRTG	6
Msp I	C`CG,G	39	MspA1 I	CMG CKG	14
Mun I	C`AATT,G	2	Nae I	GCC GGC	3
Nar I	GG`CG,CC	7	Nci I	CC`S,GG	18
Nco I	C`CATG,G	6	Nde I	CA`TA,TG	2
NgoM I	G`CCGG,C	3	Nhe I	G`CTAG,C	-
Nla III	,CATG`	39	Nla IV	GGN NCC	40
Not I	GC`GGCC,GC	-	Nru I	TCG CGA	-
Nsi I	A,TGCA`T	2	Nsp7524 I	R`CATG,Y	9
NspB II	CMG CKG	14	NspH I	R,CATG`Y	9
PacI	TTA`AT,TAA	2	Paer7 I	C`TCGA,G	-
Pal I	GG CC	44	PflM I	CCAN,NNN`NTGG	2
Ple I	GAGTC 9/10	3	Pme I	CTTT AAR	1
Pml I	CAC GTG	-	PpuM I	RG`GWC,CY	2
Psp1406 I	AA`CG,TT	1	PspA I	C`CCGG,G	3
Pst I	C,TGCA`G	5	Pvu I	CG,AT`CG	-

Pvu II	CAG CTG	7	Rsa I	GT AC	14
Rsr II	CG`GWC,CG	1	Sac I	G,AGCT`C	2
Sac II	CC,GC`GG	2	Sal I	G`TCGA,C	-
Sap I	GCTCTTC 8/11	3	Sau3A I	`GATC,	34
Sau96 I	G`GNC,C	32	Sca I	AGT ACT	-
ScrF I	CC`N,GG	38	Sec I	C`CNNG,G	45
SfaN I	GCATC 9/13	16	Sfc I	C`TRYA,G	7
Sfi I	GGCCN,NNN`NGGCC	-	Sma I	CCC GGG	3
SnaB I	TAC GTA	2	Spe I	A`CTAG,T	-
Sph I	G,CATG`C	3	Spl I	C`GTAC,G	-
Srf I	GCCC GGGC	-	Ssp I	AAT ATT	1
Stu I	AGG CCT	2	Sty I	C`CWWG,G	15
SwaI	ATTT AAAT	1	Taq I	T`CG,A	16
Tfi I	G`AWT,C	10	Tsp45 I	`GTSAC,	24
Tth111 I	GACN`N,NGTC	2	Tth111 II	CAARCA 16/14	9
Vsp I	AT`TA,AT	6	Xba I	T`CTAG,A	2
Xca I	GTA TAC	1	Xcm I	CCANNNN,N`NNNTGG3	
Xho I	C`TCGA,G	-	Xho II	R`GATC,Y	13
Xma I	C`CCGG,G	3	Xma III	C`GGCC,G	1