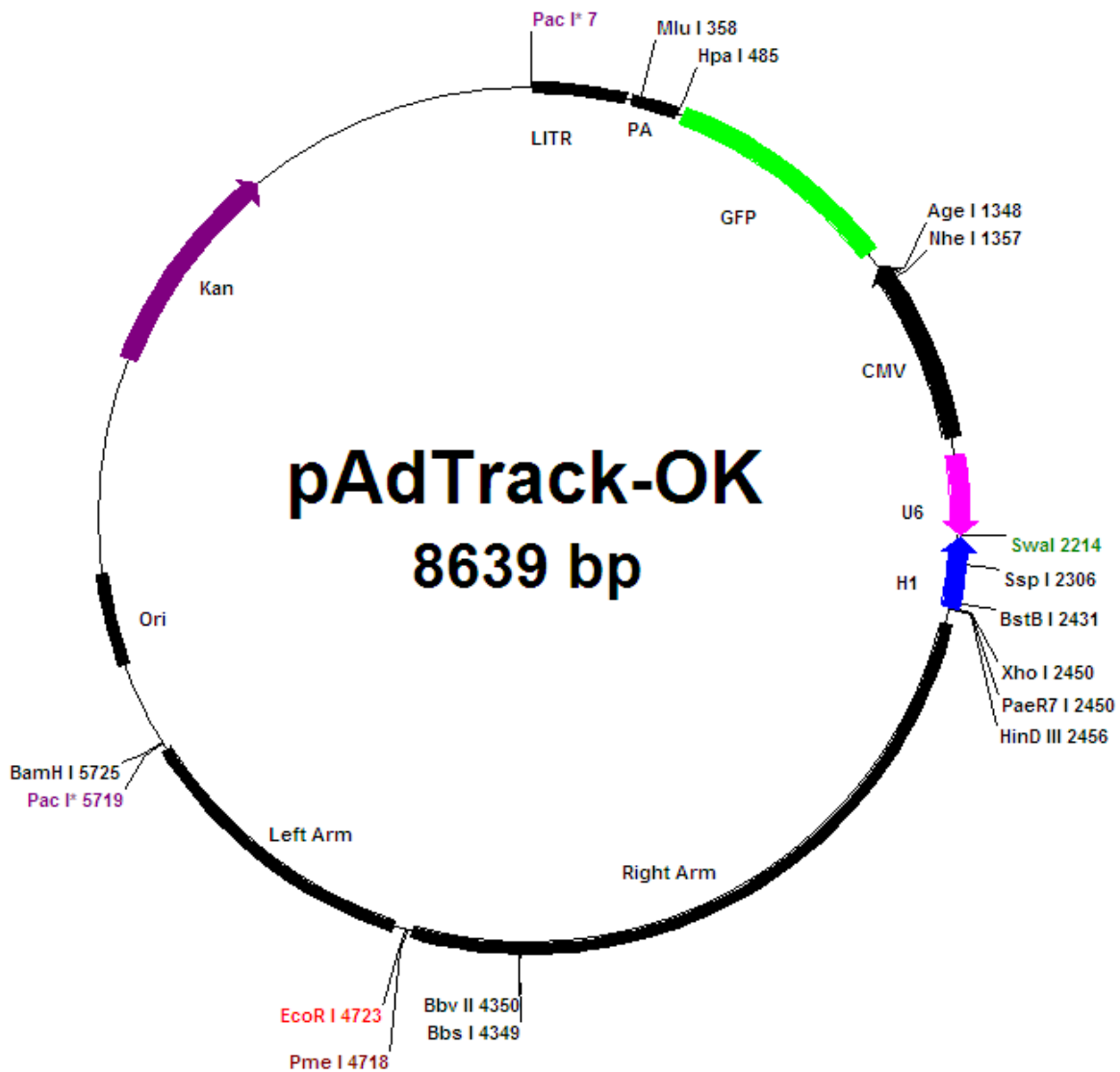


Vector: pAdTrack-OK

Antibiotic Selection: Kan

Creator(s): Fang Deng & Zhengjian Yan, Molecular Oncology Lab of The University of Chicago Medical Center

Date of Construction: October, 2013



pAdTrack-OK Full-Length Sequence

NNNTTAATTAANNNTCCCTTCCAGCTCTCTGCCCTTTTGGATTGAAGCCAATATGATAATGAGGGGGTGGAGTTTGTGACGTGGCGCGGG
CGGTGGGAACGGGGCGGGTACGTAGTAGTGTGGCGGAAGTGTGATGTTGCAAGTGTGGCGGAACACATGTAAGCGACGGATGTGGCAAAA
GTGACGTTTTTGGTGTGCGCCGGTGTACACAGGAAGTGACAATTTTCGCGCGGTTTTAGGCGGATGTTGTAGTAAATTTGGCGTAACCGA
GTAAGATTTGGCCATTTTCGCGGGAAAACGAATAAGAGGAAGTGAATCTGAATAATTTTGTGTTACTCATAGCGCGTAANNNGCGTTA
AGATACATTGATGAGTTTGGACAACCACAACCTAGAATGCAGTGAAAAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTG
TAACCATTATAAGCTGCAATAACAAGTTAAACAACAACATTCGATTCATTTTATGTTTCAGGTTTCAAGTTTCAAGGGGAGGTGTGGGAGGTTTTTTA
AAGCAAGTAAAACTCTACAAATGTGGTATGGCTGATTATGATCAGTTATCTAGATCCGGTGGATCTGAGTCCGGACTTGTACAGCTCGTC
CATGCCGAGAGTGTACCCGGCGGGTACGAACTCCAGCAGGACCATGTGATCGCGCTTCTCGTTGGGGTCTTTGCTCAGGGCGGACTGG
GTGCTCAGGTAGTGGTTGTGCGGCAGCAGCACGGGGCCGTGCCGATGGGGGTGTTCTGCTGGTAGTGGTGGCGAGCTGCACGCTGCCGT
CCTCGATGTTGTGGCGGATCTTGAAGTTCACCTTGTATGCCGTTCTTCTGCTTGTGCGCCATGATATAGACGTTGTGGCTGTTGTAGTTGTA
CTCCAGCTTGTGCCCCAGGATGTTGCCGTCTCTTGAAGTCGATGCCCTTCAGCTCGATCGGTTACCAGGGTGTGCCCTCGAACTTC
ACCTCGGCGCGGGTCTTGTAGTTGCCGTGCTCCTTGAAGAAGTGGTGCCTCCTGGACGTAGCCTTCGGGCATGGCGGACTTGAAGAAGT
CGTGTGCTTATGTTGGTTCGGGTAGCGGCTGAAGCACTGCAGCCGTAGTTCAGGGTGGTTCACGAGGGTGGCCAGGGCACGGGCAGCTT
GCCGGTGGTGCAGATGAACCTCAGGGTCAGCTTCCCGTAGGTGGCATCGCCCTCGCCCTCGCCGGACAGCTGAACCTTGTGGCCGTTTACG
TCGCGCTCCAGCTCGACAGGATGGGCACCACCCCGGTGAACAGCTCCTCGCCCTTGCCTACCATGGTGGCAGCCGGTAGCGTAGCGGAT
CTGACGGTTCATAACCAGCTCTGCTTATATAGACCTCCACCCTACACGGCTACCGCCATTTGCGTCAATGGGGCGGAGTTGTTACGA
CATTTTGGAAAGTCCCGTTGATTTTGGTGCCAAAACAACCTCCATTGACGTCAATGGGGTGGAGACTTGGAAAATCCCCGTGAGTCAAACC
GCTATCCACGCCATTGATGTAAGTCCAAAACCGCATCACCATGGTAATAGCGATGACTAATACGTAGATGTAAGTCCAAAGTAGGAAAGT
CCATAAGGTCATGTAAGTGGGCATAATGCCAGCGGGCCATTACCGTCAATGACGTCAATAGGGGGCGTACTTGGCATATGATACACTTGA
TGTACTGCCAAGTGGGCGTTTACCGTAAATACTCCACCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGAAACATACGTC
TTATTGACGTCAATGGGGCGGGTCTGTGGGGCGTACCGGCGGCTTTTACCGTAAAGTATGTAACGGCAACTCCATATATAGGCT
ATGAACATAGTACCCGTAATGATTACTATTANNNTAGCAaag**tcggcaggaagaggcctatttcccatctctctcaattcttct**
catatacgtatacaggctgttagagagataatagaatlaattgactgtaaacacaaagataattagtaaaaaaacgtgacgtagaaagt
aataatttcttggtagtttgcagttttaaataatgttttaaataaggactatacatgcttaccgtaacttgaagatttctgatttctt
ggctttatatacttgtgaaaggaATTTAAATtggtctcatacagaactataagattcccaaatccaaagacatttccagtttatgggtg
atttccagaacacatagcagacatgcaaatattgaggggcgcaactccctgtccctcacagccatcttcttccagggcgcaagcgct
gggtgttccgccttagtgactgagcctgggcccgcgattccttggaggggtgtgagcgtgagcgttogaattaagctgtttATCCTCGAGAAG
CTTTCTAGAGNNTAAGGGTGGGAAAGAATATATAAGTGGGGCTTATGATGTTTGTATCTGTTTTCGACGAGCCCGCCGCCATGA
GCACCAACTCGTTTGTAGTGAAGCATTTGTGAGCTCATATTTGACAACCGCGCATGCCCCCATGGCCCGGGTGGCTGAGAATGTGATGGGCTC
CAGCATTGATGGTGCAGCCCGTCTGCCCGCAAACCTACTACCTTACCTACGAGACCGTGTCTGGAACGCCGTTGGAGACTGCAGCCTCC
GCCGCGCTTACAGCGCTGCAGCACCGCCCGCGGGATTGTGACTGACTTTGCTTTCCTGAGCCCGCTTGAAGCAGTGCAGCTTCCCGTT
CATCCGCCCGGATGACAAGTGCAGGCTCTTTTGGCACAATTTGATTCTTTGACCCGGGAACCTAATGTGCTTCTCAGCAGCTGTTGGA
TCTGCGCCAGCAGGTTTCTGCCCTGAAGGCTTCTCCCTCCCAATGCGGTTTAAAACATAAATAAAAAACCAGACTCTGTTTGGATTG
ATCAAGCAAGTGTCTGCTGCTTTATTTAGGGTTTTGGCGCGCGGTAGGCCCGGGACCAGCGGTCTCGGTGTTGAGGGTCTCTGTGA
TTTTTTCCAGGACGTGGTAAAGGTGACTCTGGATGTTTCAGATACATGGGCATAAGCCCGCTCTGCGGGTGGAGTAGCACCCTGCAGAG
TTTATGCTGCGGGTGGTGTGTAGATGATCCAGTCTGAGCAGGAGCGCTGGCGTGGTGCCTAAAAATGCTTTCAGTACCAAGCTGATT
GCCAGGGGCAGGCCCTTGGTGAAGTGTTTACAAAGCGGTTAAGCTGGGATGGGTGCATACGTGGGGATAGAGATGCATCTTGGACTGTA
TTTTTAGGTTGGCTATGTTCCAGCCATATCCCTCCGGGATTCATGTTGTGCAGAACCACCAGCACAGTGTATCCGGTGCACCTTGGGAAA
TTTGTATGTAGCTTAGAAGGAAATGCGTGAAGAAGCTTGAGACGCCCTTGTGACCTCCAGATTTTCCATGCATTTCGTCCATAATGATG
GCAATGGGCCACGGGGCGGGCCTGGGCGAAGATATTTCTGGGATCACTAACGTCAATAGTTGTGTTCCAGGATGAGATCGTCATAGGCCA
TTTTTACAAAGCGGGCGGAGGGTGCCAGACTGCGGTATAATGGTTCATCCGGCCAGGGGCGTAGTTACCCTCACAGATTTGCATTTCA
CCAGCTTTGAGTTGCAGTGCAGGGGATCATGTCTACCTGCGGGCGATGAGAAAACGGTTTCCGGGGTAGGGGAGATCAGCTGGGAAGAA
AGCAGGTTCCTGAGCAGCTGCAGCTTACCAGCAGCGGTGGGCCGTAATCACACCTATTACCGGGTGCAACTGGTAGTTAAGAGAGCTGC
AGCTGCCGTATCCCTGAGCAGGGGGGCCACTTCGTTAAGCATGTCCCTGACTCGCATGTTTTCCCTGACCAATCCGCCAGAAGGGCTC
GCCGCCAGCGATAGCAGTCTTGAAGGAAGCAAAGTTTTTCAACGGTTTGAAGCCGTCGCGGTAGGCATGCTTTTGTAGCGTTTGACCA
AGCAGTTCAGCGGGTCCACAGCTCGGTCACTGCTTACGGCATCTCGATCCAGCATACTCTCTGTTTCGCGGGTGGGGCGGCTTTC
GCTGTACGGCAGTAGTCCGTGCTCGTCCAGACGGGCCAGGGTCATGCTTTCCACGGGGCAGGGTCTCGTCAGCGTAGTCTGGGTACG
GTGAAGGGGTGCGCTCCGGCTGCGCGTGGCCAGGTCGCTTGAAGCTGGTCTGCTGGTGTGAAGCGCTGCCGGTCTCGCCCTCGC
CGTCGGCCAGTAGCATTGACCATGTTGTATGATGCTACAGCCCTCCCGCGGTGGCCCTGGCGCGCAGCTGTCCCTTGGAGGAGGCGCC
GCACGAGGGGCAGTGCAGACTTTTGAAGGGCTAGAGCTTGGGCGGAGAAATACCGATTCCGGGGAGTAGGCATCCGCGCCGAGGCCCG
CAGACGGTCTCGCATTCCAGAGCCAGGTGAGCTTGGCCGTTCCGGGTCAAACCCAGGTTTCCCCATGCTTTTTGATGCGTCTTCTTAC
CTCTGGTTTTCCATGAGCCGGTTCACGCTCGGTGACGAAAAGGCTGTCCGTGTCCTCGTATACAGACTNNGTTTAAACGAATTCNNNTA
TAAATGCAAGTGTGCTCAAAAAATCAGGCAAAAGCCTCGCGCAAAAAAGAAAACACATCGTAGTCATGCTCATGCAGATAAAGGCAGGT
AAGCTCCGGAACCACAGAAAAAGACACCATTTTTCTCAAAATGTCTGCGGGTTTTCTGCATAAACACAAAATAAATAACAAAA
ACATTTAAACATTAGAAGCTGTCTTACAACAGGAAAAACCCCTTATAAGCATAAGACGGACTACGCCATCCGCGGTGACCGTAAAA
AAACTGGTCCCGTGATTAATAAAGCACCACCAGCAGCTCCTCGGTGATGTCGGGAGTCATAATGTAAGACTCGGTAAACACATCAGTTGA
TTCATCGGTGAGTAAAGAGCAGCCGAAATAGCCCGGGGAATACATACCCGAGGCGTAGAGACAACATTACAGCCCCATAGGAGGT
ATAACAAAATTAATAGGAGAGAAAAACACATAAACACCTGAAAAACCCCTCTGCTTAGGCAAAATAGCACCTCCCGCTCCAGAACAACAT
ACAGCGCTTACAGCGGCAGCTTAACAGTCAAGCTTACCAGTAAAAAAGAAAACCTATTAATAAACCACCTCGACACGGCACCAGCTCA
ATCAGTCAAGTGTAAAAAAGGGCAAGTGCAGAGCGAGTATATATAGGACTAAAAAATGACGTAACGGTTAAAGTCCACAAAAAACACC
AGAAAACCGCACGGAACCTACGCCAGAAACGAAAGCCAAAAACCCCAACTTCTCAAATCGTCACTCCGTTTTCCACGTTTCCAGTTACGTA
ACTTCCATTTTAAAGAAAATGACATTTCCACACATACAGTACTCCGCCCTTAAACCTACGTCACCCGCCCCGTTCCACGCCCGCGG
CCACGTCAAAAACCTCCACCCCTCATTATCATATTGGCTTCAATCCAAAATAAGGTATATATTGATGATNNTTAATTAAGGATCCNNNC
GGTGTAAATACCGCACAGATGCGTAAGGAGAAAAATACCGCATCAGGCGCTTCCGCTTCTCGCTCACTGACTCGCTGCGCTCGGTGCT
TCGGCTGCGGCGAGCGGTATCAGCTCACTCAAAGCGGTAAACGGTTATCCACAGAATCAGGGGATAACGAGGAAAGAACATGTGAGCA
AAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCTTGTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATC
GACGCTCAAGTCAAGGTTGGCGAAAACCCGACAGGACTATAAAGATACCAGGCGTTTCCCTTGAAGCTCCCTCGTGCCTCTCTGTTCC

GACCTGCCGCTTACCGGATACTGTCCGCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCATAGCTCAGCTGTAGGTATCTCAGTTCCG
 GTGTAGGTCGCTCGCTCCAAGCTGGGCTGTGTGCACGAACCCCCGTTACGCCGACCGCTGCGCCTTATCCGGTAACTATCGTCTTGAGT
 CCAAGCTGGTAAAGACACGACTTATCGCCACTGGCAGGACCCACTGTTAAACAGGATAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTT
 CTTGAAGTGGTGGCCAACTACGGCTACACTAGAAGCAGTATTTGGTATCTGCGCTGTGAAGCCAGTTACCTTCGGAAAAAGAGTT
 GGTAGCTCTTGATCCGGCAAAACAAACCACCGTGGTAGCGGTGGTTTTTTTGTGCAAGCAGCAGATTACCGCGCAAAAAAAGGATCTC
 AAGAAGATCCTTTGATCTTTTCTACGGGTCTGACGCTCAGTGAACGAAAACTCACGTTAAGGGATTTTGGTCATGAGATTATCAAAAAG
 GATCTTACCTAGATCCTTTTAAATTAATAAAGTAAAAATCAATCTAAAGTATATATGAGTAACTTTGGTCTGACAGTTACCAATGC
 TTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCTGACTCCCCGTCGTGATAGATAACTACGATACGGG
 AGGGCTTACCATCTGGCCCCAGTGTGCAATGATAACCGGAGACCCACGCTCACCGGCTCCAGATTTATCAGCAATAAACAGCCAGCCGG
 AAGGCGGAGCGCAGAAGTGGTCCTGCAACTTTATCCGCTCCATCCAGTCTATTAATTTGTGCGGGGAAGCTAGAGTAAGTAGTTCCGCCA
 GTTAATAGTTTGGCGCAAGTGTGTTGNNNNNNAAAGGACTCTCACCTAGATCCTTTTTCAGTGAAGAACCGTCCGCGAAGCCGCTGCTG
 ACCCGGATGAATGTGAGCTACTGGCTATCTGGACAAGGGAACCGCAAGCGCAAAAGAGAAGCAGGTAGCTTGCAGTGGGCTTACATGG
 CGATAGCTAGACTGGGCGGTTTTATGGACAGCAAGCGAACCGGAATTGCCAGCTGGGGCGCCCTCTGGTAAGGTTGGGAAGCCCTGCAAAG
 TAAACTGGATGGCTTTCTCGCGCCAAGGATCTGATGGCGCAGGGGATCAAGCTCTGATCAAGAGACAGGATGAGGATCGTTTCGCATGAT
 TGAACAAGATGGATTGCACGAGGTTCTCCGCGCGCTTGGTGGAGAGGCTATTTCGGCTATGACTGGGCACAACAGACAATCGGCTGCTCT
 GATGCCCGCGTGTCCGGCTGTCAGCGCAGGGGCGCCGGTTCTTTTTGTCAAGACCGACCTGTCCGGTGCCTGAATGAAGTGAAGACG
 AGGCAGCGCGGCTATCGTGGCTGGCCACGACGGGCGTTCCTTGCGCAGCTGTGCTCGACGTTGTCACTGAAGCGGGAAGGACTGGCTGCT
 ATTTGGGCGAAGTGGCGGGGACGATCTCCTGTCACTCTGCTCCTGCGGAAAGTATCCATCATGGTATGCAATGCGCGGCTGCTGCTG
 CATACGCTTGATCCGGTACTGCCCATTTCGACCAACCAAGCGAAACATCGCATCGAAGGAGCACGTAAGTCCGATGGAAGCCGGTCTTGTGCG
 ATCAGGATGATCTGGACGAAGAGCATCAGGGGCTCGCGCCAGCCGAACGTTTCGCCAGGCTCAAGGCGAGCATGCCCGACGGCGAGGATCT
 CGTCGTGACCCATGGCGATGCTTGTGCGGAATATCATGTTGAAAAATGGCCGCTTTTTCTGGATTCATCGACTGTGGCCGCTGGGTGTG
 GCGGACCGCTATCAGGACATAGCGTTGGCTACCCGTGATATGCTGAAGAGCTTGGCGCGCAATGGGCTGACCGCTTCTCGTCTTTACG
 GTATCGCCGCTCCCGATTCGCGAGCGCATCGCCTTCTATCGCCTTCTTTCGACGAGTTCTTCTGAATTTTGTAAAAATTTTTGTAAAAATCAGCT
 CATTTTTTAAACCATAGGCCGAAATCGGCAACATCCCTTATAAATCAAAGAATAGACCCGATAGGGTTGAGTGTGTTCAGTTTGGAA
 CAAGAGTCCACTATAAAGAAGCTGGACTCCAACGCTCAAAGGGCGAAAAACCGTCTATCAGGGCGATGGCCCACTGAAACCATCACCC
 AAATCAAGTTTTTTTGGCGTGGAGGTCGCGTAAAGCTCTAAATCGGAACCCATAAGGGAGCCCCGATTTAGAGCTTACGCGGGAAAGCCGG
 CGAACGTGGCGAGAAAGGAAGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAGTGTAGCGGTACGCTGCGCGTAACCACCAC
 ACCCGCGCGCTTAATGCGCCGNN

Unique enzymes in pAdTrace-OK

Mlu I	A`CGCG,T	358				4415	4417	4459	4535
Hpa I	GTT AAC	485				4540	4549	4877	5149
Age I	A`CGG,T	1348				5263	5293	5468	5600
Nhe I	G`CTAG,C	1357				5621	5639	5746	5772
Swa I	ATTT AAAT	2214				5789	5832	5839	5860
Ssp I	AAT ATT	2306				5951	5979	6106	6125
BsiC I	TT`CG,AA	2431				6246	6356	6491	6500
BstB I	TT`CG,AA	2431				6862	6953	7083	7206
PaeR7 I	C`TCGA,G	2450				7302	7405	7468	7562
Xho I	C`TCGA,G	2450				7626	7727	7730	7970
Hind III	A`AGCT,T	2456				8010	8015	8065	8081
PflM I	CCAN,NNN`NTGG	3108	Afl III	(4)	158	358		4869	5906
BstX I	CCAN,NNN`NTGG	3547	Age I	(1)	1348				
Bbs I	GAAGAC 8/12	4349	Aha II	(10)	1506	1692	1775	1828	
Bbv II	GAAGAC 7/11	4350				2315	2421	3503	4456
Bst1107 I	GTA TAC	4703				7248	7496		
Xca I	GTA TAC	4703	Ahd I	(2)	3067	6799			
Pme I	CTTT AAAC	4718	Alu I	(47)	25	469	632	806	
EcoR I	G`AATTC,C	4723				917	965	1181	1214
Avr II	C`CTAG,G	5242				1286	1319	1386	2440
BamH I	G`GATC,C	5725				2458	2580	2813	2905
BspH I	T`CATG,A	6626				3185	3271	3321	3471
Psp1406 I	AA`CG,TT	7025				3813	3840	3910	3916
Eag I	C`GGCC,G	7402				4119	4439	4496	4583
Xma III	C`GGCC,G	7402				4827	5042	5366	5848
Rsr II	CG`GWC,CG	8012				6074	6164	6210	6467
Dra III	CAC,NNN`GTC	8359				6988	7117	7170	7196
Xmn I	GAANN NNTTC	8580				7242	7333	7602	8060
Number of enzymes = 28			Alw I	(26)	596	616	646	843	

The following enzymes do not cut in pAdTrack-OK:

Acc65 I	Afl II	Asc I	Asp718	Bgl II						
Blp I	BsiW I	Bsu36 I	Cla I	Eco72 I						
EcoN I	EcoR V	Esp I	Fse I	Kpn I						
Not I	Nru I	Pml I	Pvu I	Sal I						
Sca I	Sfi I	Spe I	Spl I	Srf I						
					AlwN I	(6)	2747	2905	2990	4711
							6322	7120		
					Apa I	(3)	2395	3560	3866	
					ApaL I	(2)	3446	6220		
					Apo I	(7)	257	430	2209	3457
							4723	8161	8172	
					Ase I	(5)	5	2041	5198	5717
							6971			
					Ava I	(4)	2450	2877	3057	5132
					Ava II	(8)	680	3061	3085	4110
							4251	4329	6937	8012
					Avr II	(1)	5242			
					BamH I	(1)	5725			
					Ban I	(12)	1300	1483	2314	3243
							3664	4455	5359	6747
							7247	7495	7530	8396
					Ban II	(9)	2395	2582	2639	2795
							3560	3866	4585	7861
							8434			
					Bbe I	(4)	2318	4459	7251	7499
					Bbs I	(1)	4349			
					Bbv I	(20)	764	767	1190	2540
							2543	2735	2762	2822

pAdTrack-OK: sites sorted by name:

Aat II	(5)	1509	1695	1778	1831
		2424			
Acc I	(2)	3764	4702		
Aci I	(110)	88	106	126	151
		233	243	294	658
		661	721	834	972
		1011	1078	1119	1361
		1422	1443	1547	1580
		1671	1838	1852	1864
		1892	2376	2395	2410
		2535	2538	2541	2667
		2730	2733	2736	2745
		2757	2761	2763	2795
		2826	2830	2960	3049
		3067	3195	3313	3566
		3569	3654	3658	3675
		3771	3851	3990	4007
		4065	4108	4169	4179

pAdTrack-OK Vector

		2914	3849	3865	3925		1892	2361	2363	2397
		4448	5308	6325	6328		2596	2763	2832	3045
		6534	7569	7611	8132		3047	3049	3654	4169
Bbv II	(1)	4350					4303	4369	4417	4434
Bcl I	(2)	587	7337				4504	4537	4774	5474
Bcn I	(19)	656	1310	2615	2879		5641	5953	6534	6864
		2880	3059	3060	3405		7562	7863	8251	8543
		3797	3887	4296	4522		8560	8562		
		5134	5135	6287	6983	BstX I	(1)	3547		
		7104	7501	7661		BstY I	(15)	600	609	836
Bfa I	(13)	397	598	1358	1949		2911	5725	6547	6558
		2380	2463	5243	6401		6644	6656	7045	7057
		6654	6989	7055	7197		7309	7667	7913	
		8510				Cac8 I	(57)	804	811	1135
Bgl I	(7)	1672	1743	1865	2939		1216	1359	1673	1866
		4990	6919	8579			2359	2363	2395	2600
Bpm I	(5)	657	897	2623	5251		2667	2761	2795	2799
		6869					2803	2830	2921	3045
Bsa I	(6)	2226	2687	3075	4051		3047	3269	3287	3656
		4563	6860				4004	4076	4303	4307
BsaA I	(7)	1612	2080	3338	5549		4434	4441	4544	4991
		7069	7800	8359			5153	5472	5837	5923
BsaB I	(3)	586	3002	7355			5960	6520	6911	7148
BsaH I	(10)	1506	1692	1775	1828		7172	7223	7240	7390
		2315	2421	3503	4456		7576	7795	7861	7867
		7248	7496				7895	7899	7940	7944
BsaJ I	(39)	925	980	1004	1167		7998	8462	8505	8519
		1307	1337	1588	2350		8562			
		2403	2606	2613	2761	Cfr10 I	(11)	202	1185	1348
		2877	3057	3279	3291		4352	4658	4989	6879
		3403	3560	3573	3697		7815	7996	8460	
		3698	3795	4223	4239	Csp6 I	(12)	208	627	909
		4310	4391	4415	4427		1568	1619	1652	1411
		4445	4520	5045	5132		1732	2070	4191	7801
		5133	5242	6066	7101	Dde I	(11)	613	715	733
		7305	7659	7928			2898	3472	3833	3929
BsaW I	(10)	603	618	1348	3442		6181	6590	6756	
		4829	5056	6112	6259	Dpn I	(34)	589	602	611
		7229	7527				690	838	1365	2913
BseR I	(4)	1311	4149	4465	5034		3005	3215	3595	3628
Bsg I	(7)	792	1116	1213	2829		3758	3809	4147	5727
		3439	4494	5419			6474	6549	6560	6568
BsiC I	(1)	2431					6646	6658	6763	7047
BsiE I	(7)	1349	3077	5123	5822		7059	7311	7328	7339
		6246	7405	8392			7358	7669	7747	7828
BsiHKA I	(9)	734	2552	2582	3450		7837	7915		
		4210	4585	6224	7609	DpnII	(34)	587	600	609
		7799					688	836	1363	2911
Bsm I	(4)	406	499	3532	4563		3003	3213	3593	3626
BsmA I	(12)	1515	2225	2688	2712		3756	3807	4145	5725
		3074	3158	3495	4052		6472	6547	6558	6566
		4562	5155	6861	7339		6644	6656	6761	7045
BsmB I	(2)	3159	3494				7057	7309	7326	7337
BsmF I	(9)	1454	1622	1773	2313		7356	7667	7745	7826
		3074	3943	4096	4680		7835	7913		
		7647				Dra I	(9)	546	2122	2135
BsoF I	(68)	470	659	753	756		2966	4718	4921	6665
		807	814	1098	1120		6684			
		1179	2529	2532	2535	Dra III	(1)	8359		
		2538	2541	2724	2733	Drd I	(4)	4395	6014	7523
		2736	2745	2748	2751	Dsa I	(9)	1337	1588	2606
		2811	2903	3193	3567		3560	4239	4391	4415
		3570	3838	3841	3854		7928			
		3911	3914	3917	4007	Eae I	(11)	283	875	1264
		4180	4299	4350	4418		4373	4587	4982	7402
		4437	4459	4540	4748		7576	7967	7994	
		5294	5297	5812	5830	Eag I	(1)	7402		
		5833	5951	6106	6249	Ear I	(4)	1963	5790	7840
		6314	6317	6523	6851	Eco47 III	(4)	1356	3233	4348
		7302	7405	7457	7468	Eco57 I	(9)	946	1143	1189
		7558	7563	7600	7641		2956	4362	6453	7641
		7728	7731	7734	7970		8073			
		8066	8107	8121	8539	EcoO109 I	(5)	1973	3085	3288
Bsp120 I	(3)	2391	3556	3862			4545			4251
Bsp1286 I	(2)	2552	7799			EcoR I	(1)	4723		
BspH I	(1)	6626				EcoR II	(23)	925	979	1054
BspM I	(8)	2914	3776	3816	4136		1291	1666	1859	2349
		4810	7154	7383	7764		3101	3278	3572	3617
BspM II	(3)	618	4829	5056			3697	4103	4222	4309
Bsr I	(20)	729	1658	2392	3218		4375	4574	4606	5932
		3898	5013	5318	6313		6053	6066	7881	
		6326	6440	6846	6964	Ehe I	(4)	2316	4457	7249
		7007	7079	7124	7205	Fnu4H I	(68)	470	659	753
		7289	7439	7640	8273		807	814	1098	1120
BsrB I	(5)	2410	5265	5839	8109		1179	2529	2532	2535
		8503					2538	2541	2724	2733
BsrD I	(3)	3558	6860	7729			2736	2745	2748	2751
BsrG I	(2)	207	626				2811	2903	3193	3567
BssH II	(6)	2361	3043	3045	4301		3570	3838	3841	3854
		4432	8560				3911	3914	3917	4007
BssS I	(5)	1155	4462	4569	6079		4180	4299	4350	4418
		8088					4437	4459	4540	4748
Bst1107 I	(1)	4703					5294	5297	5812	5830
BstB I	(1)	2431					5833	5951	6106	6249
BstE II	(2)	4123	5012				6314	6317	6523	6851
BstN I	(23)	927	981	1056	1168		7302	7405	7457	7468
		1293	1668	1861	2351		7558	7563	7600	7641
		3103	3280	3574	3619		7728	7731	7734	7970
		3699	4105	4224	4311		8066	8107	8121	8539
		4377	4576	4608	5934	Fok I	(19)	184	258	942
		6055	6068	7883			2809	3139	3338	3634
BstU I	(38)	88	231	233	294		3676	3910	4518	6765
		350	360	693	1011		6946	7118	7301	7363

pAdTrack-OK Vector

NspB II	(13)	2747	2763	2905	3067
		3813	3840	3916	4417
		5293	6248	6493	7242
		7602			
NspH I	(9)	162	2301	2602	3958
		3973	4078	4873	5910
		7901			
PacI	(2)	7	5719		
PaeR7 I	(1)	2450			
Pal I	(38)	285	765	877	1166
		1266	1675	1868	1975
		2393	2612	3056	3289
		3558	3572	3638	3696
		3864	3941	4222	4309
		4375	4425	4546	4589
		4984	5393	5921	5932
		5950	6384	6842	6922
		7404	7578	7969	7996
		8209	8351		
PflM I	(1)	3108			
Ple I	(5)	623	1547	5068	6285
		8294			
Pme I	(1)	4718			
PpuM I	(2)	3085	4251		
Psp1406 I	(1)	7025			
PspA I	(3)	2877	3057	5132	
Pst I	(4)	2725	2752	3182	3915
Pvu II	(6)	2905	3813	3840	3916
		7242	7602		
Rsa I	(12)	209	628	910	1412
		1569	1620	1653	1708
		1733	2071	4192	7802
Rsr II	(1)	8012			
Sac I	(2)	2582	4585		
Sac II	(2)	2764	4418		
Sap I	(3)	5790	7840	8050	
Sau3A I	(34)	587	600	609	650
		688	836	1363	2911
		3003	3213	3593	3626
		3756	3807	4145	5725
		6472	6547	6558	6566
		6644	6656	6761	7045
		7057	7309	7326	7337
		7356	7667	7745	7826
		7835	7913		
Sau96 I	(31)	680	763	1164	1673
		1866	1973	2391	2392
		2610	3055	3061	3085
		3288	3556	3557	3695
		3862	3863	3939	4110
		4220	4251	4329	4424
		4545	5391	6841	6920
		6937	8012	8350	
ScrF I	(42)	655	927	981	1056
		1168	1293	1309	1668
		1861	2351	2614	2878
		2879	3058	3059	3103
		3280	3404	3574	3619
		3699	3796	3886	4105
		4224	4295	4311	4377
		4521	4576	4608	5133
		5134	5934	6055	6068
		6286	6982	7103	7500
		7660	7883		
Sec I	(39)	925	980	1004	1167
		1307	1337	1588	2350
		2403	2606	2613	2761
		2877	3057	3279	3291
		3403	3560	3573	3697
		3698	3795	4223	4239
		4310	4391	4415	4427
		4445	4520	5045	5132
		5133	5242	6066	7101
		7305	7659	7928	
SfaN I	(20)	429	846	945	960
		1236	1590	3342	3362
		4147	4539	4620	5744
		5782	6002	7455	7710
		7794	7858	7926	8133
Sfc I	(6)	2721	2748	3178	3911
		6171	6362		
Sma I	(3)	2879	3059	5134	
SnaB I	(2)	1612	5549		
Sph I	(3)	2602	4078	7901	
Ssp I	(1)	2306			
Sty I	(11)	1337	1588	2403	2606
		3291	4391	4427	4445
		5242	7305	7928	
SwaI	(1)	2214			
Taq I	(17)	823	952	967	994
		1288	2176	2431	2451
		4144	5352	6006	7609
		7765	7789	7825	7987
		8392			
Tfi I	(10)	1988	2241	2399	2867
		3408	4516	5096	5881
		7981	8115		
Tsp45 I	(23)	77	109	183	218
		663	1152	2080	2382
		2770	3117	3510	4123
		4272	4674	4994	5012
		5374	5525	5615	5647

Tth111 I	(2)	2698	7614		
Tth111 II	(10)	859	2817	2983	3022
		4110	4881	6495	6504
		6534	7932		
Vsp I	(5)	5	2041	5198	5717
		6971			
Xba I	(2)	597	2462		
Xca I	(1)	4703			
Xcm I	(3)	2991	4582	7312	
Xho I	(1)	2450			
Xho II	(15)	600	609	836	1363
		2911	5725	6547	6558
		6644	6656	7045	7057
		7309	7667	7913	
Xma I	(3)	2877	3057	5132	
Xma III	(1)	7402			
Xmn I	(1)	8580			

Site usage in pAdTrack-OK:

Aat II	G,ACGT`C	5	Acc I	GT`MK,AC	2
Acc65 I	G`GTAC,C	-	Aci I	C`CG,C	110
Afl II	C`TTAA,G	-	Afl III	A`CRYG,T	4
Age I	A`CCGG,T	1	Aha II	GR`CG,YC	10
Ahd I	GACNN,N`NNGTC	2	Alu I	AG CT	47
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	5	Asp718	G`GTAC,C	-
Ava I	C`YCGR,G	4	Ava II	G`GWC,C	8
Avr II	C`CTAG,G	1	BamH I	G`GATC,C	1
Ban I	G`GYRC,C	12	Ban II	G,RCGY`C	9
Bbe I	G,GGCC`C	4	Bbs I	GAAGAC 8/12	1
Bbv I	GCAGC 13/17	20	Bbv II	GAAGAC 7/11	1
Bcl I	T`GATC,A	2	Bcn I	CC,S`GG	19
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	5	Bsa I	GGTCTC 7/11	6
BsaA I	YAC GTR	7	BsaB I	GATNN NNATC	3
BsaH I	GR`CG,YC	10	BsaJ I	C`CNNG,G	39
Bsaw I	W`CCGG,W	10	BseI I	GAGGAG 16/14	4
Bsg I	GTGCAG 22/20	7	BsiC I	TT`CG,AA	1
BsiE I	CG,RY`CG	7	BsiHK I	G,WGCW`C	9
BsiW I	C`GTAC,G	-	Bsm I	GAATG,C 7	4
BsmA I	GTCCT`/9	12	BsmB I	CGTCTC 7/11	2
BsmF I	GGGAC 15/19	9	BsoF I	GC`N,GC	68
Bsp120 I	G`GGCC,C	3	Bsp1286 I	G,DCGH`C	2
BspH I	T`CATG,A	1	BspM I	ACCTGC 10/14	8
BspM II	T`CCGG,A	3	Bsr I	ACT,GG`	20
BsrB I	GAG CGG	5	BsrD I	GCAATG, 8	3
BsrG I	T`GTAC,A	2	BssH II	G`CGCG,C	6
BssS I	C`TCGT,G	5	Bst1107 I	GTA TAC	1
BstB I	TT`CG,AA	1	BstE II	G`GTNAC,C	2
BstN I	CC`W,GG	23	BstU I	CG CG	38
BstX I	CCAN,NNNN`NTGG	1	BstY I	R`GATC,Y	15
Bsu36 I	CC`TNA,GG	-	Cac8 I	GCN NGC	57
Cfr10 I	R`CCGG,Y	11	Cla I	AT`CG,AT	-
Csp6 I	G`TA,C	12	Dde I	C`TNA,G	11
Dpn I	GA TC	34	DpnII	`GATC,	34
Dra I	TTT AAA	9	Dra III	CAC,NNN`GTG	1
Drd I	GACNN,NN`NNGTC	4	Dsa I	C`CRYG,G	9
Eae I	Y`GGCC,R	11	Eag I	C`GGCC,G	1
Ear I	CTCTTC 7/10	4	Eco47 III	AGC GCT	4
Eco57 I	CTGAAG 21/19	9	Eco72 I	CAC GTG	-
EcoN I	CCTNN`N,NNAGG	-	Eco1019 I	RG`GNC,CY	5
EcoR I	G`AATT,C	1	EcoR II	`CCWGG,	23
EcoR V	GAT ATC	-	Ehe I	GGC GCC	4
Esp I	GC`TNA,GC	-	Fnu4H I	GC`N,GC	68
Fok I	GGATG 14/18	19	Fse I	GG,CCGG`CC	-
Fsp I	TGC GCA	2	Gdi II	`YGGC,CG	9
Gsu I	CTGGAG 21/19	5	Hae I	WGG CCW	7
Hae II	R,GGCC`Y	13	Hae III	GG CC	38
Hga I	GACGC 9/14	6	HgiA I	G,WGCW`C	9
HgiE II	ACCNNNNNNGGT	-1/132	Hha I	G,CG`C	59
Hinc II	GTY RAC	2	Hind II	GTY RAC	2
Hind III	A`AGCT,T	1	Hinf I	G`ANT,C	22
HinI I	GR`CG,YC	10	HinP I	G`CG,C	59
Hpa I	GTT AAC	1	Hpa II	C`CG,G	43
Hph I	GGTGA 12/11	20	Kas I	G`GGCG,C	4
Kpn I	G,GTAC`C	-	Mae I	C`TA,G	13
Mae II	A`CG,T	33	Mae III	`GTNAC,	40
Mbo I	`GATC,	34	Mbo II	GAAGA 12/11	19
Mlu I	A`CGCG,T	1	Mme I	TCCRAC 25/23	5
Mnl I	CCTC 10/10	56	Msc I	TGG CCA	3
Mse I	T`TA,A	37	Msl I	CAYNN NNRTG	5
Msp I	C`CG,G	43	MspAl I	CMG CKG	13
Mun I	C`AATT,G	2	Nae I	GCC GGC	3
Nar I	GG`CG,CC	4	Nci I	CC`S,GG	19
Nco I	C`CATG,G	5	Nde I	CA`TA,TG	2
NgoM I	G`CCGG,C	3	Nhe I	G`CTAG,C	1
Nla III	,CATG`	40	Nla IV	GGN NCC	33
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	13	NspH I	R,CATG`Y	9
PacI	TTA`AT,TAA	2	PaeR7 I	C`TCGA,G	1
Pal I	GG CC	38	Pf1M I	CCAN,NNN`NTGG	1
Ple I	GAGTC 9/10	5	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,G	3
Pst I	C,TGCA`G	4	Pvu I	CG,AT`CG	-
Pvu II	CAG CTG	6	Rsa I	GT AC	12
Rsr II	CG`GWC,CG	1	Sac I	G,AGCT`C	2

pAdTrack-OK Vector

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	31	Sca I	AGT ACT	-
ScrF I	CC`N,GG	42	Sec I	C`CNNG,G	39
SfaN I	GCATC 9/13	20	Sfc I	C`TRYA,G	6
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	-	Sty I	C`CWWG,G	11
SwaI	ATTT AAAT	1	Taq I	T`CG,A	17
Tfi I	G`AWT,C	10	Tsp45 I	`GTSAC,	23
Tth111 I	GACN`N,NGTC	2	Tth111 II	CAARCA 16/14	10
Vsp I	AT`TA,AT	5	Xba I	T`CTAG,A	2
Xca I	GTA TAC	1	Xcm I	CCANNNN,N`NNNTGG3	
Xho I	C`TCGA,G	1	Xho II	R`GATC,Y	15
Xma I	C`CCGG,G	3	Xma III	C`GGCC,G	1