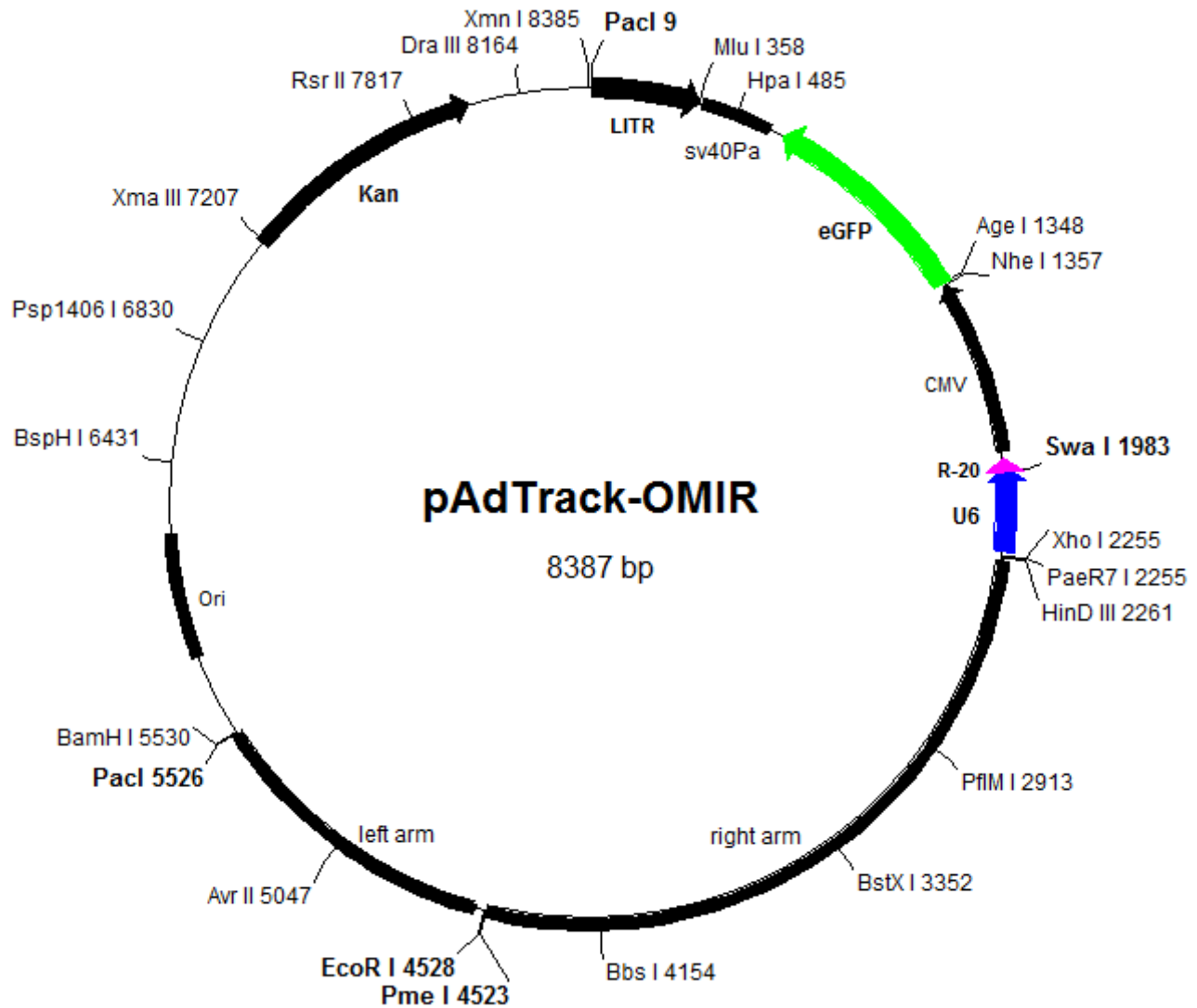


Vector: pAdTrack-OMIR (adeno shuttle vector for one-step mature miRNA expression)

Antibiotic Selection: Kan

Creator(s): Xian Chen, Molecular Oncology Laboratory of The University of Chicago

Date of Construction: June, 2013



Forward (5p) Oligo:

5'-tgtggaagggaaggaATTTxxxxxxx-5p ttttttaaggctgggcaggaagagg-3'

Reverse (3p) Oligo:

5'-atcttgtggaagggaCGAACACCyyyyyyyy-3p ttttttAAATCGCCACATAGGTTTCGT-3'

3p Reverse-Complement

5'-ACGAACCTATGTGGCGATTTaaaaaap3-yyyyyyyGGTGTTCGTcctttccacaagat-3'

pAdTrack-OMIR Full-Length Sequence

NNNTTAATTAANNNTCCCTTCCAGCTCTCTGCCCTTTTGGATTGAAGCCAATATGATAATGAGGGGTGGAGTTTG
TGACGTGGCGCGGGGCGTGGGAACGGGGCGGGTGACGTAGTAGTGTGGCGGAAGTGTGATGTTGCAAGTGTGGCGGA
ACACATGTAAGCGACGGATGTGGCAAAAGTGACGTTTTTGGTGTGCGCCGGTGTACACAGGAAGTGACAATTTTCGC
GCGGTTTTAGGCGGATGTTGTAGTAAATTTGGGCGTAACCGAGTAAGATTTGGCCATTTTCGCGGGAAAACCTGAATA
AGAGGAAGTGAATCTGAATAATTTTGTGTTACTCATAGCGCGTAANNNCGCGTTAAGATACATTGATGAGTTTGGGA
CAAACCACAACCTAGAATGCAGTGAATAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCAT
TATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATTCAATTTATGTTTTAGGTTTCAGGTTTCAGGGGGAGGTGTGGGAGG
TTTTTTAAAGCAAGTAAAACCTCTACAAATGTGGTATGGCTGATTATGATCAGTTATCTAGATCCGGTGGATCTGAG
TCCGGACTTGTACAGCTCGTCCATGCCGAGAGTATCCCGGCGGGTACGAACTCCAGCAGGACCATGTGATCGC
GCTTCTCGTTGGGGTCTTTGCTCAGGGCGGACTGGGTGCTCAGGTAGTGGTTGTTCGGGCAGCAGCACGGGGCCGTCCG
CCGATGGGGGTGTTCTGCTGGTAGTGGTCCGGCAGCTGCACGCTGCCGTCTCGATGTTGTGGCGGATCTTGAAGTT
CACCTTGATGCCGTTCTTCTGCTTGTTCGGCCATGATATAGACGTTGTGGCTGTTGTAGTTGTACTCCAGCTTGTGCC
CCAGGATGTTGCCGTCCTCCTTGAAGTCGATGCCCTCAGCTCGATGCGGTTACCAGGGTGTTCGCCCTCGAACTTC
ACCTCGGGCGGGTCTTGTAGTTGCCGTCGTCCTTGAAGAAGATGGTGCCTCCTGGACGTAGCCTTCGGGCATGGC
GGACTTGAAGAAGTCGTGCTGCTTCATGTGGTCCGGGTAGCGGCTGAAGCACTGCACGCCGTAGGTGAGGGTGGTCA
CGAGGGTGGGCCAGGGCACGGGCAGCTTCCGGTGGTGCAGATGAACTTCAGGGTTCAGCTTCCCGTAGGTGGCATCG
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CAATAGGGGGCGTACTTGGCATATGATACACTTGTACTGCAAGTGGGCAGTTTACCAGTAAATACTCCACCCAT
TGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAACATACGTCAATTTGACGTCAATGGGCGGGGGTTCGTT
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GCTTCAGCCGCTGCAGCCACCGCCCGGGATTGTGACTGACTTTGCTTTCTGAGCCCGCTTGCAAGCAGTGCAGC
TTCCCGTTCATCCGCCCGCATGACAAGTTGACGGCTCTTTTGGCACAATTGGATTCTTTGACCCGGGAACCTAATG
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GGGCGCTGGCAAGTGTAGCGGTACGCTGCGCGTAACCACCACACCCGCGCTTAATGCGCCGNNNNNNN

Unique enzymes in pAdTrack-OMIR:

Mlu I	A`CGCG,T	358	Bbv II	GAAGAC 7/11	4155
Hpa I	GTT AAC	485	Bst1107 I	GTA TAC	4508
Age I	A`CCGG,T	1348	Xca I	GTA TAC	4508
Nhe I	G`CTAG,C	1357	Pme I	CTTT AAAC	4523
Swa I	ATTT AAAT	1983	EcoR I	G`AATT,C	4528
PaeR7 I	C`TCGA,G	2255	Avr II	C`CTAG,G	5047
Xho I	C`TCGA,G	2255	BamH I	G`GATC,C	5530
Hind III	A`AGCT,T	2261	BspH I	T`CATG,A	6431
PEI M I	CCAN,NNN`NTGG	2913	Psp1406 I	AA`CG,TT	6830
BstX I	CCAN,NNN`NTGG	3352	Eag I	C`GGCC,G	7207
Bbs I	GAAGAC 8/12	4154	Xma III	C`GGCC,G	7207
			Rsr II	CG`GWC,CG	7817

Dra III CAC,NNN`GTG 8164
 Xmn I GAANN|NNTTC 8385
 Number of enzymes = 25

Ban II (8) 2387 2444 2600 3365
 Bbe I (3) 4264 7056 7304
 Bbs I (1) 4154
 Bbv I (20) 764 1190 2345
 2348 2540 2567 2627
 2719 3654 3670 3730
 4253 5113 6130 6133
 6339 7374 7416 7937

The following enzymes do not cut in
 pAdTrack-OMIR:

Acc65 I Afl II Asc I Asp718 Bgl II
 Blp I BsiC I BsiW I BstB I Bsu36 Bbv II
 I Bcl I (1) 4155
 Cla I Eco72 I EcoN I EcoR V Esp I Bcn I (19) 656 7142
 Fse I Kpn I Not I Nru I Pml I (2) 587
 Pvu I Sal I Sca I Sfi I Spe I (19) 656 1310 2420 2684
 2685 2864 2865 3210
 3602 3692 4101 4327
 4939 4940 6092 6788
 6909 7306 7466
 397 598 1358 1949
 2268 5048 6206 6459
 6794 6860 7002 8315
 Bgl I (7) 1672 1743 1865 2744
 4795 6724 8384
 Bpm I (5) 657 897 2428 5056
 6674
 Bsa I (5) 2492 2880 3856 4368
 6665
 BsaA I (7) 1612 2122 3143 5354
 6874 7605 8164
 BsaB I (3) 586 2807 7160
 BsaH I (8) 1506 1692 1775 1828
 3308 4261 7053 7301
 BsaJ I (37) 925 980 1004 1167
 1307 1337 1588 2411
 2418 2566 2682 2862
 3084 3096 3208 3365
 3378 3502 3503 3600
 4028 4044 4115 4196
 4220 4232 4250 4325
 4850 4937 4938 5047
 5871 6906 7110 7464
 7733
 BsaW I (10) 603 618 1348 3247
 4634 4861 5917 6064
 7034 7332
 BseR I (4) 1311 3954 4270 4839
 Bsg I (7) 792 1116 1213 2634
 3244 4299 5224
 BsiE I (7) 1349 2882 4928 5627
 6051 7210 8197
 BsiHKA I (9) 734 2357 2387 3255
 4015 4390 6029 7414
 7604
 Bsm I (4) 406 499 3337 4368
 BsmA I (11) 1515 2493 2517 2879
 2963 3300 3857 4367
 4960 6666 7144
 BsmB I (2) 2964 3299
 BsmF I (8) 1454 1622 1773 2879
 3901 4485 7452
 BsoF I (68) 470 659 753 756
 807 814 1098 1120
 1179 2334 2337 2340
 2343 2346 2529 2538
 2541 2550 2553 2556
 2616 2708 2998 3372
 3375 3643 3646 3659
 3716 3719 3722 3812
 3985 4104 4155 4223
 4242 4264 4345 4553
 5099 5102 5617 5635
 5638 5756 5911 6054
 6119 6122 6328 6656
 7107 7210 7262 7273
 7363 7368 7405 7446
 7533 7536 7539 7775
 7871 7912 7926 8344
 Bsp120 I (2) 3361 3667
 Bsp1286 I (2) 2357 7604
 BspH I (1) 6431
 BspM I (8) 2719 3581 3621 3941
 4615 6959 7188 7569
 BspM II (3) 618 4634 4861
 Bsr I (19) 729 1658 3023 3703
 4818 5123 6118 6131
 6245 6651 6769 6812
 6884 6929 7010 7094
 7244 7445 8078
 BsrB I (4) 5070 5644 7914 8308
 BsrD I (3) 3363 6665 7534

pAdTrack-OMIR: sites sorted by name:

Aat II (4) 1509 1695 1778 1831
 Acc I (2) 3569 4507
 Aci I (107) 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 2340 2343 2346
 2472 2535 2538 2541
 2550 2562 2566 2568
 2600 2631 2635 2765
 2854 2872 3000 3118
 3371 3374 3459 3463
 3480 3576 3656 3795
 3812 3870 3913 3974
 3984 4220 4222 4264
 4340 4345 4354 4682
 4954 5068 5098 5273
 5405 5426 5444 5551
 5577 5594 5637 5644
 5665 5756 5784 5911
 5930 6051 6161 6296
 6305 6667 6758 6888
 7011 7107 7210 7273
 7367 7431 7532 7535
 7775 7815 7820 7870
 7886 7912 8054 8193
 8308 8335 8363
 Afe I (4) 1356 3038 4153 5089
 Afl III (4) 158 358 4674 5711
 Age I (1) 1348
 Aha II (8) 1506 1692 1775 1828
 3308 4261 7053 7301
 Ahd I (2) 2872 6604
 Alu I (46) 25 469 632 806
 917 965 1181 1214
 1286 1319 1386 2263
 2385 2618 2710 2990
 3076 3126 3276 3618
 3645 3715 3721 3924
 4244 4301 4388 4632
 4847 5171 5653 5879
 5969 6015 6272 6793
 6922 6975 7001 7047
 7138 7407 7865 7994
 8212 8251
 Alw I (27) 596 616 646 843
 1370 1950 2723 2815
 3014 3405 3568 3946
 5526 5537 6273 6359
 6359 6456 6457 6857
 6858 7121 7138 7168
 7479 7546 7725
 AlwN I (6) 2552 2710 2795 4516
 6127 6925
 Apa I (2) 3365 3671
 ApaL I (2) 3251 6025
 Apo I (7) 257 430 1984 3262
 4528 7966 7977
 Ase I (5) 5 2159 5003 5522
 6776
 Ava I (4) 2255 2682 2862 4937
 Ava II (8) 680 2866 2890 3915
 4056 4134 6742 7817
 Avr II (1) 5047
 BamH I (1) 5530
 Ban I (11) 1300 1483 3048 3469
 4260 5164 6552 7052
 7300 7335 8201

BsrG I	(2)	207	626			EcoR II	(22)	925	979	1054	1166
BssH II	(5)	2848	2850	4106	4237			1291	1666	1859	2906
		8365						3083	3377	3422	3502
BssS I	(5)	1155	4267	4374	5884			3908	4027	4114	4180
		7893						4379	4411	5737	5858
Bst1107 I	(1)	4508				Ehe I	(3)	5871	7686		
BstE II	(2)	3928	4817			Fnu4H I	(68)	4262	7054	7302	
BstN I	(22)	927	981	1056	1168			470	659	753	756
		1293	1668	1861	2908			807	814	1098	1120
		3085	3379	3424	3504			1179	2334	2337	2340
		3910	4029	4116	4182			2343	2346	2529	2538
		4381	4413	5739	5860			2541	2550	2553	2556
		5873	7688					2616	2708	2998	3372
BstU I	(35)	88	231	233	294			3375	3643	3646	3659
		350	360	693	1011			3716	3719	3722	3812
		1892	2401	2568	2637			3985	4104	4155	4223
		2850	2852	2854	3459			4242	4264	4345	4553
		3974	4108	4174	4222			5099	5102	5617	5635
		4239	4309	4342	4579			5638	5756	5911	6054
		5279	5446	5758	6339			6119	6122	6328	6656
		6669	7367	7668	8056			7107	7210	7262	7273
		8348	8365	8367				7363	7368	7405	7446
BstX I	(1)	3352						7533	7536	7539	7775
BstY I	(16)	600	609	836	1363			7871	7912	7926	8344
		1954	2716	5530	6352	Fok I	(19)	184	258	942	1308
		6363	6449	6461	6850			2614	2944	3143	3439
		6862	7114	7472	7718			3481	3715	4323	6570
Cac8 I	(54)	804	811	1135	1183			6751	6923	7106	7168
		1216	1359	1673	1866			7625	7650	8014	
		2405	2472	2566	2600	Fsp I	(2)	6826	7403		
		2604	2608	2635	2726	Gdi II	(9)	876	1263	4179	4391
		2850	2852	3074	3092			4788	7206	7208	7771
		3461	3809	3881	4108			7798			
		4112	4239	4246	4349	Gsu I	(5)	658	898	2429	5057
		4796	4958	5277	5642			6675			
		5728	5765	6325	6716	Hae I	(7)	285	3443	4114	5726
		6953	6977	7028	7045			5737	6189	7383	
		7195	7381	7600	7666	Hae II	(12)	1358	3040	3808	4155
		7672	7700	7704	7745			4264	5091	5589	5959
		7749	7803	8267	8310			7056	7304	8315	8323
		8324	8367			Hae III	(37)	285	765	877	1166
Cfr10 I	(11)	202	1185	1348	3661			1266	1675	1868	2227
		4157	4463	4794	6684			2417	2861	3094	3363
		7620	7801	8265				3377	3443	3501	3669
Csp6 I	(12)	208	627	909	1411			3746	4027	4114	4180
		1568	1619	1652	1707			4230	4351	4394	4789
		1732	2130	3996	7606			5198	5726	5737	5755
Dde I	(11)	613	715	733	2594			6189	6647	6727	7209
		2703	3277	3638	3734			7383	7774	7801	8014
		5986	6395	6561				8156			
Dpn I	(35)	589	602	611	652	Hga I	(6)	1422	2415	3315	4164
		690	838	1365	1956			5821	6399		
		2718	2810	3020	3400	HgiA I	(9)	734	2357	2387	3255
		3433	3563	3614	3952			4015	4390	6029	7414
		5532	6279	6354	6365			7604			
		6373	6451	6463	6568	HgiE II	(2)	3653	6290		
		6852	6864	7116	7133	Hha I	(55)	88	202	233	350
		7144	7163	7474	7552			695	1011	1052	1357
		7633	7642	7720				2403	2724	2850	2852
DpnII	(35)	587	600	609	650			2854	3039	3459	3807
		688	836	1363	1954			4052	4096	4108	4110
		2716	2808	3018	3398			4124	4154	4174	4239
		3431	3561	3612	3950			4241	4263	4309	4344
		5530	6277	6352	6363			4581	5090	5448	5588
		6371	6449	6461	6566			5621	5891	5958	6058
		6850	6862	7114	7131			6232	6341	6734	6827
		7142	7161	7472	7550			6957	7055	7126	7295
		7631	7640	7718				7303	7367	7404	7670
Dra I	(9)	546	1983	2067	2080			7930	8314	8322	8348
		2771	4523	4726	6470			8367	8369	8378	
		6489				HinC II	(2)	485	2649		
Dra III	(1)	8164				Hind II	(2)	485	2649		
Drd I	(4)	4200	5819	7328	8119	HinD III	(1)	2261			
Dsa I	(9)	1337	1588	2411	2566	Hinf I	(20)	615	1539	2211	2672
		3365	4044	4196	4220			2792	2925	3213	3769
		7733						4321	4865	4879	4901
Eae I	(11)	283	875	1264	4112			5611	5686	6082	6599
		4178	4392	4787	7207			7786	7920	8091	8113
		7381	7772	7799		HinI I	(8)	1506	1692	1775	1828
Eag I	(1)	7207						3308	4261	7053	7301
Ear I	(4)	2236	5595	7645	7855	HinP I	(55)	86	200	231	348
Eco47 III	(4)	1356	3038	4153	5089			693	1009	1050	1355
Eco57 I	(9)	946	1143	1189	2530			2401	2722	2848	2850
		2761	4167	6258	7446			2852	3037	3457	3805
		7878						4050	4094	4106	4108
EcoO109 I	(5)	2226	2890	3093	4056			4122	4152	4172	4237
		4350						4239	4261	4307	4342
EcoR I	(1)	4528						4579	5088	5446	5586

		5619	5889	5956	6056	Msp I	(43)	203	604	619	655
		6230	6339	6732	6825			1186	1246	1309	1349
		6955	7053	7124	7293			2418	2683	2863	3208
		7301	7365	7402	7668			3248	3498	3600	3662
		7928	8312	8320	8346			3690	4099	4158	4325
		8365	8367	8376				4464	4635	4795	4862
Hpa I	(1)	485						4938	5918	6065	6091
Hpa II	(43)	203	604	619	655			6281	6685	6719	6786
		1186	1246	1309	1349			6908	7035	7206	7283
		2418	2683	2863	3208			7305	7333	7464	7554
		3248	3498	3600	3662			7621	7802	8266	
		3690	4099	4158	4325	MspAl I	(13)	2552	2568	2710	2872
		4464	4635	4795	4862			3618	3645	3721	4222
		4938	5918	6065	6091			5098	6053	6298	7047
		6281	6685	6719	6786			7407			
		6908	7035	7206	7283	Mun I	(2)	494	2666		
		7305	7333	7464	7554	Nae I	(3)	4796	7803	8267	
		7621	7802	8266		Nar I	(3)	4261	7053	7301	
Hph I	(19)	120	841	970	994	Nci I	(19)	655	1309	2419	2683
		1322	1327	1578	2933			2684	2863	2864	3209
		3923	4094	4394	4490			3601	3691	4100	4326
		4812	5415	6448	6675			4938	4939	6091	6787
		6849	7480	8166				6908	7305	7465	
Kas I	(3)	4260	7052	7300		Nco I	(5)	1337	1588	2411	4196
Mae I	(12)	397	598	1358	1949			7733			
		2268	5048	6206	6459	Nde I	(2)	1716	2051		
		6794	6860	7002	8315	NgoM I	(3)	4794	7801	8265	
Mae II	(31)	81	113	187	889	Nhe I	(1)	1357			
		1060	1273	1506	1611	Nla III	(39)	162	642	687	882
		1692	1775	1816	1828			1077	1107	1341	1592
		2116	2121	2912	3142			1652	2218	2353	2407
		3407	5236	5348	5353			2415	2947	2997	3220
		5419	5451	6414	6830			3273	3337	3568	3763
		6873	7417	7604	8108			3778	3883	4038	4200
		8120	8163	8273				4427	4461	4608	4614
Mae III	(39)	77	109	183	218			4678	4794	4860	5715
		266	337	455	663			6435	6994	7175	7520
		1152	1450	1799	1886			7706	7737	7763	
		2038	2117	2575	2922	Nla IV	(31)	764	1302	1485	2443
		3315	3513	3928	4077			2867	2891	3050	3363
		4479	4799	4817	5179			3471	3492	3635	3669
		5237	5330	5349	5354			3745	3917	4057	4262
		5398	5420	5452	6067			4352	4639	5166	5532
		6130	6246	6529	7421			5743	5782	6554	6648
		7727	8337	8349				6689	7054	7302	7337
Mbo I	(35)	587	600	609	650			8203	8224	8236	
		688	836	1363	1954	Nsi I	(2)	3161	3339		
		2716	2808	3018	3398	Nsp7524 I	(8)	158	2403	3759	3774
		3431	3561	3612	3950			3879	4674	5711	7702
		5530	6277	6352	6363	NspB II	(13)	2552	2568	2710	2872
		6371	6449	6461	6566			3618	3645	3721	4222
		6850	6862	7114	7131			5098	6053	6298	7047
		7142	7161	7472	7550			7407			
		7631	7640	7718		NspH I	(8)	162	2407	3763	3778
Mbo II	(18)	856	1049	1052	1097			3883	4678	5715	7706
		2224	3306	3396	3597	Pac I	(2)	9	5526		
		3634	4155	5583	6372	Paer7 I	(1)	2255			
		6445	6846	7661	7871	Pal I	(37)	285	765	877	1166
		7953	8303					1266	1675	1868	2227
Mlu I	(1)	358						2417	2861	3094	3363
Mne I	(5)	2500	2695	5925	6109			3377	3443	3501	3669
		8140						3746	4027	4114	4180
Mnl I	(55)	56	304	521	530			4230	4351	4394	4789
		569	830	950	1001			5198	5726	5737	5755
		1013	1151	1244	1250			6189	6647	6727	7209
		1331	1411	2238	2263			7383	7774	7801	8014
		2541	2760	2765	2880			8156			
		2964	3214	3329	3459	PflM I	(1)	2913			
		3528	3974	4068	4121	Ple I	(5)	623	1547	4873	6090
		4226	4248	4251	4263			8099			
		4283	4456	4584	4859	Pme I	(1)	4523			
		4982	5049	5073	5331	PpuM I	(2)	2890	4056		
		5478	5609	5819	5892	Psp1406 I	(1)	6830			
		6143	6543	6624	6770	PspA I	(3)	2682	2862	4937	
		7066	7152	7216	7352	Pst I	(4)	2530	2557	2987	3720
		7709	7901	8192		Pvu II	(6)	2710	3618	3645	3721
Msc I	(3)	285	4114	7383				7047	7407		
Mse I	(36)	5	9	363	484	Rsa I	(12)	209	628	910	1412
		545	1943	1982	2066			1569	1620	1653	1708
		2079	2159	2691	2770			1733	2131	3997	7607
		3122	3707	3755	4522	Rsr II	(1)	7817			
		4725	4828	5003	5142	Sac I	(2)	2387	4390		
		5245	5368	5522	5526	Sac II	(2)	2569	4223		
		6417	6469	6474	6488	Sap I	(3)	5595	7645	7855	
		6541	6776	6815	7974	Sau3A I	(35)	587	600	609	650
		7986	8003	8101	8371			688	836	1363	1954
Msl I	(5)	1587	3350	3415	4609			2716	2808	3018	3398
		7738						3431	3561	3612	3950

		5530	6277	6352	6363	AlwN I	CAG,NNN`CTG	6	Apa I	G,GGCC`C	2
		6371	6449	6461	6566	ApaL I	G`TGCA,C	2	Apo I	R`AATT,Y	7
		6850	6862	7114	7131	Asc I	GG`CGCG,CC	-	Ase I	AT`TA,AT	5
		7142	7161	7472	7550	Asp718	G`GTAC,C	-	Ava I	C`YCGR,G	4
		7631	7640	7718		Ava II	G`GWC,C	8	Avr II	C`CTAG,G	1
Sau96 I	(29)	680	763	1164	1673	BamH I	G`GATC,C	1	Ban I	G`GYRC,C	
		1866	2226	2415	2860	Ban II	G,RCGY`C	8	Bbe I	G,GCGC`C	3
		2866	2890	3093	3361	Bbs I	GAAGAC 8/12	1	Bbv I	GCAGC 13/17	
		3362	3500	3667	3668	Bbv II	GAAGAC 7/11	1	Bcl I	T`GATC,A	2
		3744	3915	4025	4056	Bcn I	CC,S`GG	19	Bfa I	C`TA,G	
		4134	4229	4350	5196	Bgl I	GCCN,NNN`NGGC	-	Bgl II	A`GATC,T	-
		6646	6725	6742	7817	Blp I	GC`TNA,GC	7	Bpm I	CTGGAG 22/20	5
ScrF I	(41)	8155				Bsa I	GGTCTC 7/11	5	BsaA I	YAC GTR	7
		655	927	981	1056	BsaB I	GATNN NNATC	3	BsaH I	GR`CG,YC	8
		1168	1293	1309	1668	BsaJ I	C`CNNG,G	37	BsaW I	W`CCGG,W	
		1861	2419	2683	2684	BseR I	GAGGAG 16/14	4	Bsg I	GTGCAG 22/20	7
		2863	2864	2908	3085	BsiC I	TT`CG,AA	-	BsiE I	CG,RY`CG	7
		3209	3379	3424	3504	BsiHKA I	G,WGCW`C	9	BsiW I	C`CTAG,G	-
		3601	3691	3910	4029	Bsm I	GAATG,C 7	4	BsmA I	GTCTC`/9	
		4100	4116	4182	4326	BsmB I	CGTCTC 7/11	2	BsmF I	GGGAC 15/19	8
		4381	4413	4938	4939	BsoF I	GC`N,GC	68	Bsp120 I	G`GGCC,C	2
		5739	5860	5873	6091	Bsp1286 I	G, DGCH`C	2	BspH I	T`CATG,A	1
		6787	6908	7305	7465	BspM I	ACCTGC 10/14	8	BspM II	T`CCGG,A	3
Sec I	(37)	7688				Bsr I	ACT,GG`	19	BsrB I	GAG CGG	4
		925	980	1004	1167	BsrD I	GCAATG, 8	3	BsrG I	T`GTAC,A	2
		1307	1337	1588	2411	BssH II	G`CGCG,C	5	Bsss I	C`TCGT,G	5
		2418	2566	2682	2862	Bst1107 I	GTA TAC	1	BstB I	TT`CG,AA	-
		3084	3096	3208	3365	BstE II	G`GTNAC,C	2	BstN I	CC`W,GG	-
		3378	3502	3503	3600	BstU I	CG CG	35	BstX I	CCAN,NNNN`NTGG	1
		4028	4044	4115	4196	BstY I	R`GATC,Y	16	Bsu36 I	CC`TNA,GG	-
		4220	4232	4250	4325	Cac8 I	GCN NGC	54	Cfr10 I	R`CCGG,Y	
		4850	4937	4938	5047	Cla I	AT`CG,AT	-	Csp6 I	G`TA,C	
SfaN I	(20)	5871	6906	7110	7464	Dde I	C`TNA,G	11	Dpn I	GA TC	
		7733				DpnII	`GATC,	35	Dra I	TTT AAA	9
		429	846	945	960	Dra III	CAC,NNN`GTG	1	Drd I	GACNN,NN`NNGTC	4
		1236	1590	3147	3167	Dsa I	C`CRYG,G	9	Eae I	Y`GGCC,R	
Sfc I	(6)	3952	4344	4425	5549	Eag I	C`GGCC,G	1	Ear I	CTCTTC 7/10	4
		5587	5807	7260	7515	Eco47 III	AGC GCT	4	Eco57 I	CTGAAG 21/19	9
		7599	7663	7731	7938	Eco72 I	CAC GTG	-	EcoN I	CCTNN`N,NNAGG	-
Sma I	(3)	5976	6167	4939		EcoO109 I	RG`GNC,CY	5	EcoR I	G`AATT,C	1
SnaB I	(2)	2684	2864			EcoR II	`CCWGG,	22	EcoR V	GAT ATC	-
Sph I	(3)	2684	2864	4939		Ehe I	GGC GCC	3	Esp I	GC`TNA,GC	-
Sph I	(3)	2407	3883	7706		Fnu4H I	GC`N,GC	68	Fok I	GGATG 14/18	
Sty I	(10)	1337	1588	2411	3096	Fse I	GG,CCGG`CC	-	Fsp I	TGC GCA	2
		4196	4232	4250	5047	Gdi II	`YGGC,CG	9	Gsu I	CTGGAG 21/19	5
		7110	7733			Hae I	WGG CCW	7	Hae II	R,GCGC`Y	
Swa I	(1)	1983	823	952	967	Hae III	GG CC	37	Hga I	GACGC 9/14	6
Taq I	(16)	1288	2024	2256	3949	HgiA I	G,WGCW`C	9	HgiE II	ACCNNNNNNGGT	-
		5157	5811	7414	7570	Hha I	G,CG`C	55	Hinc II	GTY RAC	2
		7594	7630	7792	8197	Hind II	GTY RAC	2	Hind III	A`AGCT,T	1
Tfi I	(8)	2211	2672	3213	4321	Hinf I	G`ANT,C	20	HinI I	GR`CG,YC	8
		4901	5686	7786	7920	HinP I	G`CG,C	55	Hpa I	GTT AAC	1
Tsp45 I	(22)	77	109	183	218	Hpa II	C`CG,G	43	Hph I	GGTGA 12/11	
		663	1152	2117	2575	Kas I	G`GCGC,C	3	Kpn I	G,GTAC`C	-
		2922	3315	3928	4077	Mae I	C`TA,G	12	Mae II	A`CG,T	
		4479	4799	4817	5179	Mae III	`GTNAC,	39	Mbo I	`GATC,	
		5330	5420	5452	7421	Mbo II	GAAGA 12/11	18	Mlu I	A`CGCG,T	1
		7727	8337			Mme I	TCCRAC 25/23	5	Mnl I	CCTC 10/10	
Tth111 I	(2)	2503	7419			Msc I	TGG CCA	3	Mse I	T`TA,A	
Tth111 II	(10)	859	2622	2788	2827	Msl I	CAYNN NNRTG	5	Msp I	C`CG,G	
		3915	4686	6300	6309	MspAl I	CMG CKG	13	Mun I	C`AATT,G	2
		6339	7737			Nae I	GCC GGC	3	Nar I	GG`CG,CC	3
Vsp I	(5)	5	2159	5003	5522	Nci I	CC`S,GG	19	Nco I	C`CATG,G	5
		6776				Nde I	CA`TA,TG	2	NgoM I	G`CCGG,C	3
Xba I	(2)	597	2267			Nhe I	G`CTAG,C	1	Nla III	,CATG`	
Xca I	(1)	4508				Nla IV	GGN NCC	31	Not I	GC`GGCC,GC	-
Xcm I	(3)	2796	4387	7117		Nru I	TCG CGA	-	Nsi I	A,TGCA`T	2
Xho I	(1)	2255				Nsp7524 I	R`CATG,Y	8	NspB II	CMG CKG	
Xho II	(16)	600	609	836	1363	NspH I	R,CATG`Y	8	Pac I	TTA,AT`TAA	2
		1954	2716	5530	6352	PaeR7 I	C`TCGA,G	1	Pal I	GG CC	
		6363	6449	6461	6850	Pf1M I	CCAN,NNN`NTGG	1	Ple I	GAGTC 9/10	5
		6862	7114	7472	7718	Pme I	CTTT AAAC	1	Pml I	CAC GTG	-
Xma I	(3)	2682	2862	4937		PpuM I	RG`GWC,CY	2	Psp1406 I	AA`CG,TT	1
Xma III	(1)	7207				PspA I	C`CCGG,G	3	Pst I	C,TGCA`G	4
Xmn I	(1)	8385				Pvu I	CG,AT`CG	-	Pvu II	CAG CTG	6
						Rsa I	GT AC	12	Rsr I	CG`GWC,CG	1
						Sac I	G,AGCT`C	2	Sac II	CC,GC`GG	2
						Sal I	G`TCGA,C	-	Sap I	GCTCTC 8/11	3
						Sau3A I	`GATC,	35	Sau96 I	G`GNC,C	

Site usage in pAdTrack-OMIR:

Aat II	G,ACGT`C	4	Acc I	GT`MK,AC	2
Acc65 I	G`GTAC,C	-	Aci I	C`CG,C	
Afe I	AGC GCT	4	Afl II	C`TTAA,G	-
Afl III	A`CRYG,T	4	Age I	A`CCGG,T	1
Aha II	GR`CG,YC	8	Ahd I	GACNN,N`NNGTC	2
Alu I	AG CT	46	Alw I	GGATC 8/9	
27					

Sca I	AGT ACT	-	ScrF I	CC`N,GG	Sty I	C`CWG,G	10	Swa I	ATTT AAAT	1	
41					Taq I	T`CG,A	16	Tfi I	G`AWT,C	8	
Sec I	C`CNGG,G	37	SfaN I	GCATC 9/13	Tsp45 I	`GTSAC,	22	Tth111 I	GACN`N,NGTC	2	
20					Tth111 II	CAARCA 16/14	10	Vsp I	AT`TA,AT	5	
Sfc I	C`TRYA,G	6	Sfi I	GGCCN,NNN`NGGCC	-	Xba I	T`CTAG,A	2	Xca I	GTA TAC	1
Sma I	CCC GGG	3	SnaB I	TAC GTA	2	Xcm I	CCANNNN,N`NNNTGG3		Xho I	C`TCGA,G	1
Spe I	A`CTAG,T	-	Sph I	G,CATG`C	3	Xho II	R`GATC,Y	16	Xma I	C`CCGG,G	3
Spl I	C`GTAC,G	-	Srf I	GCCC GGGC	-	Xma III	C`GGCC,G	1	Xmn I	GAANN NNTTC	1
Ssp I	AAT ATT	-	Stu I	AGG CCT	-						