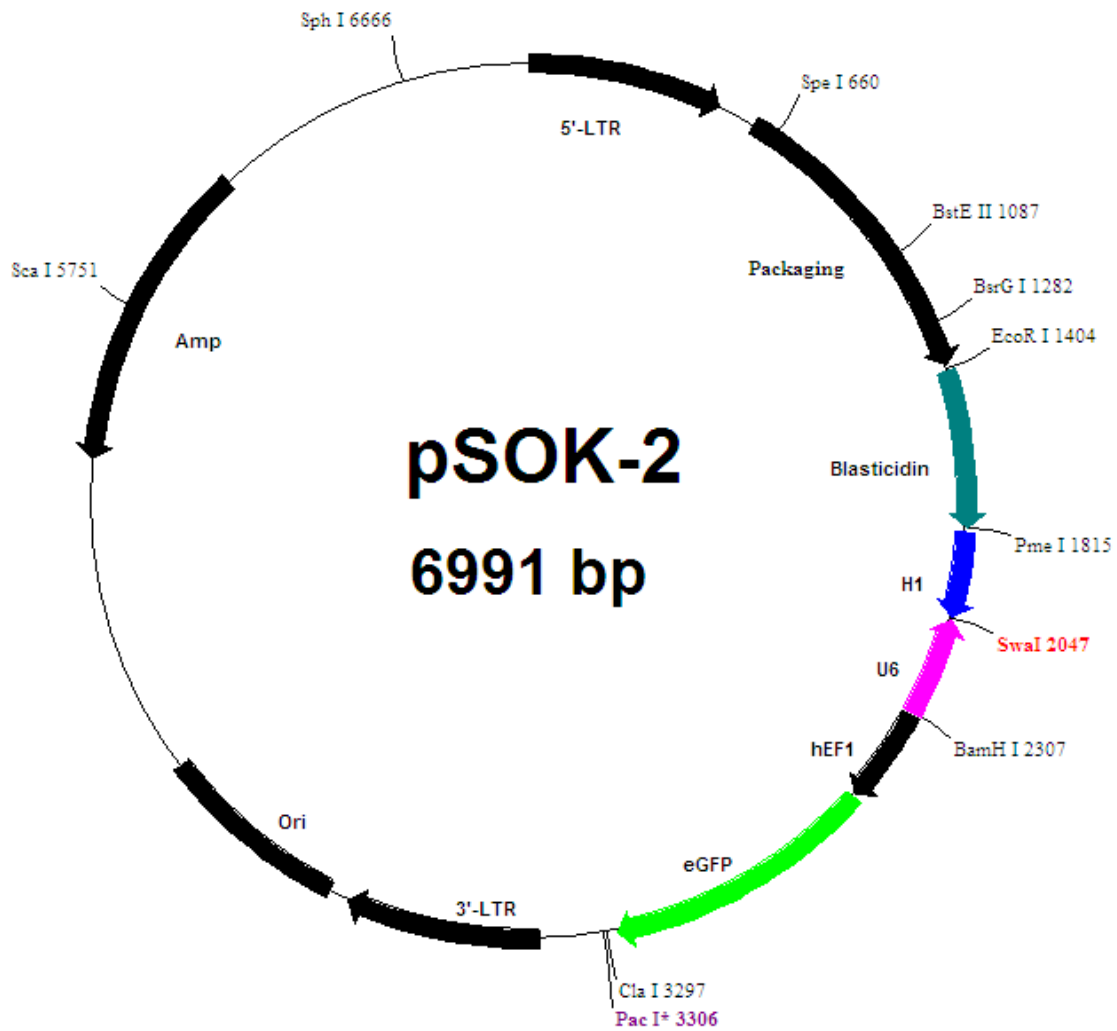


**Vector:** pSOK2

**Antibiotic Selection:** Amp

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

**Date of Construction:** June, 2013



## pSOK2 Full-Length Sequence

TGAAAGACCCACCTGTAGGTTTGGCAAGCTAGCTTAAGTAACGCCATTTTGCAAGGCATGGAAAAACATAAATGAGAATA  
 GAGAAGTTAGATCAAGGTTAGGAACAGAGAGACAGCAGAATATGGGCCAAACAGGATATCTGTGGTAAGCAGTTCCCTCCCCG  
 GCTCAGGGCCAAGAACAGATGGTCCCCAGATGCGGTCCCGCCCTCAGCAGTTTCTAGAGAACCATCAGATGTTTCCAGGGTG  
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 tgtatgagacca **ATTTAAAT** tcctttccacaagatataaaagccaagaaatcgaaatactttcaagttacggta  
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 gtattttgaactaataatctttgtgtttacagtc aaattaattctaattatctcttaacagccttgtaatgata  
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 AAGCCTATAGAGTACGAGCCATAGATAAAAATAAAGATTTTTATTTAGTCTCCAGAAAAGGGGGAA

Unique enzymes in pSOK2:

Spe I	A`CTAG,T	660				3196	3256	3259	3552		
BstE II	G`GTNAC,C	1087				3558	3854	4068	4171		
BsrG I	T`GTAC,A	1282				4227	4237	4261	4304		
EcoR I	G`AATT,C	1404				4311	4332	4423	4451		
Bbv II	GAAGAC 7/11	1497				4578	4597	4718	4828		
Bbs I	GAAGAC 8/12	1498				4963	4972	5334	5425		
Nru I	TCG CGA	1644				5616	5662	5783	5827		
Pme I	CTTT AAAC	1815				5904	6013	6112	6159		
BsiC I	TT`CG,AA	1828				6333	6372	6382	6408		
BstB I	TT`CG,AA	1828				6446	6459	6485	6542		
Bsp120 I	G`GGCC,C	1866				6801					
Apa I	G,GGCC`C	1870				Afl II	(3)	35	1006	3388	
SwaI	ATTT AAAT	2047				Afl III	(1)	4378			
BsaA I	YAC GTR	2181				Age I	(2)	2402	2567		
BamH I	G`GATC,C	2307				Aha II	(10)	742	1398	1838	1944
BspM II	T`CCGG,A	2310						5808	6190	6494	6677
Mun I	C`AATT,G	2395						6791	6812		
Bsp1286 I	G,DGCH`C	3189				Ahd I	(4)	1020	3801	3847	5271
Cla I	AT`CG,AT	3297				Alu I	(37)	30	34	336	580
PacI	TTA ATTAA	3304						668	676	1515	1620
Afl III	A`CRYG,T	4378						1821	2554	2600	2633
Sca I	AGT ACT	5751						2705	2738	2954	3002
Sph I	G,CATG`C	6666						3113	3287	3383	3387
Number of enzymes = 23								3691	4010	4043	4138
								4202	4320	4546	4636
								4682	4939	5460	5560
								5623	6302	6321	6566
								6922			

The following enzymes do not cut in pSOK2:

Alw I	(17)	453	1051	1357	1727
		2303	2314	2320	3075
		3272	4940	5026	5026
		5123	5124	5588	5903
		5909			
AlwN I	(2)	3512	4794		
Apa I	(1)	1870			
ApaL I	(3)	4692	5938	6435	
Apo I	(3)	872	1404	2048	
Asc I	(2)	368	3723		
Ase I	(5)	2218	4021	4149	4208
		5443			
Asp718	(2)	401	3756		
Ava I	(7)	331	397	498	2369
		2471	3686	3752	
Ava II	(14)	186	199	254	551
		690	1219	1572	3236
		3541	3554	3609	5409
		5631	6886		
BamH I	(1)	2307			
Ban I	(17)	244	401	1142	1397
		1687	1943	2330	2405

pSOK: sites sorted by name:

Aat II	(3)	745	1841	6193	
Acc65 I	(2)	401	3756		
Aci I	(69)	197	203	566	588
		688	727	860	901
		1042	1133	1199	1298
		1317	1355	1532	1614
		1681	1849	1864	1883
		2424	2461	2541	2798
		2839	2906	2945	3083

		2615	3599	3756	4122			4225	4227	4425	5006
		5219	6493	6676	6790			5336	5829	6161	6261
		6811						6263	6366		
Ban II	(8)	338	351	1677	1870	BstY I	(9)	2307	2313	3079	5019
		3693	3706	6743	6757			5030	5116	5128	5896
Bbe I	(6)	1401	1947	6497	6680			5913			
		6794	6815			Bsu36 I	(2)	781	1018		
Bbs I	(1)	1498				Cac8 I	(39)	28	32	319	370
Bbv I	(12)	931	939	1524	1629			582	586	631	650
		2831	3115	3122	4797			1199	1508	1622	1866
		4800	5006	5700	6311			1898	1902	2703	2736
Bbv II	(1)	1497						2784	3108	3115	3381
Bcn I	(16)	164	399	400	571			3385	3674	3725	4171
		724	2610	3264	3519			4200	4309	4395	4432
		3754	3755	4759	5455			4992	5383	6396	6544
		5806	6307	6342	6693			6564	6568	6664	6705
Bfa I	(15)	31	219	661	673			6753	6795	6825	
		1394	1522	1879	2410	Cfr10 I	(7)	629	2402	2567	2730
		2559	3384	3574	4873			5351	6814	6823	
		5126	5461	6896		Cla I	(1)	3297			
Bgl I	(2)	5391	6509			Csp6 I	(11)	402	658	1283	2189
Bpm I	(5)	3024	3264	3321	5341			2452	3008	3290	3757
		6959						5750	6426	6937	
Bsa I	(7)	476	530	1149	2031	Dde I	(21)	75	167	208	476
		3809	3830	5332				781	1018	1290	1808
BsaA I	(1)	2181						3183	3201	3428	3522
BsaB I	(2)	1711	2312					3563	3832	3903	4653
BsaH I	(10)	742	1398	1838	1944			5062	5228	5768	6194
		5808	6190	6494	6677			6429			
BsaJ I	(40)	161	240	250	397	Dpn I	(30)	94	459	1046	1178
		461	497	546	547			1363	1647	1708	1722
		694	733	734	746			2309	2315	2322	3081
		747	1167	1236	1245			3229	3267	3448	4946
		1265	1266	1411	1716			5021	5032	5040	5118
		1854	1907	2303	2472			5130	5235	5576	5594
		2578	2608	2748	2911			5640	5898	5915	5951
		2935	2990	3516	3595			6536	6760		
		3605	3752	3815	4117	DpnII	(30)	92	457	1044	1176
		4538	6612	6690	6696			1361	1645	1706	1720
BsaW I	(7)	2310	2327	2402	2567			2307	2313	2320	3079
		4584	4731	5562				3227	3265	3446	4944
BseR I	(5)	464	1293	1332	2610			5019	5030	5038	5116
		3818						5128	5233	5574	5592
Bsg I	(5)	1600	2520	2708	2805			5638	5896	5913	5949
		3129						6534	6758		
BsiC I	(1)	1828				Dra I	(7)	1815	2047	2126	2139
BsiE I	(11)	592	727	1532	1648			5137	5156	5848	
		2323	2572	4294	4718	Drd I	(3)	1500	4486	6355	
		5641	5790	6537		Dsa I	(4)	694	1411	2578	6696
BsiHKA I	(7)	338	3189	3693	4696	Eae I	(12)	582	724	1109	1130
		5857	5942	6439				1414	1529	2651	3040
BsmA I	(22)	107	475	531	609			4217	5659	6693	6825
		734	832	949	1073	Eag I	(2)	724	1529		
		1132	1150	1328	1431	Ear I	(7)	1070	1311	2295	4000
		2032	3338	3461	3808			4262	6066	6554	
		3829	5333	6107	6260	Eco47 III	(3)	923	2563	6732	
		6304	6976			Eco57 I	(7)	1509	2732	2778	2975
BsmB I	(7)	733	831	1072	1131			4925	5939	6943	
		1329	6261	6303		EcoN I	(3)	784	1392	2413	
BsmF I	(12)	172	185	564	729	EcoO109 I	(7)	254	1219	1572	2285
		764	1205	1585	1944			3609	6247	6701	
		2351	3527	3540	6672	EcoR I	(1)	1404			
BsoF I	(43)	727	901	917	920	EcoR II	(23)	239	546	733	746
		925	928	1133	1513			1167	1192	1215	1235
		1532	1606	1609	1612			1265	1715	1907	2623
		1615	1618	1782	2541			2748	2860	2935	2989
		2739	2798	2820	3104			3594	4116	4404	4525
		3111	3162	3165	3259			4538	6611	6888	
		4203	4284	4302	4305	EcoR V	(3)	140	3494	6879	
		4423	4578	4721	4786	Ehe I	(6)	1399	1945	6495	6678
		4789	4995	5323	5662			6792	6813		
		5689	5784	6013	6300	Fnu4H I	(43)	727	901	917	920
		6409	6513	6586				925	928	1133	1513
Bsp120 I	(1)	1866						1532	1606	1609	1612
Bsp1286 I	(1)	3189						1615	1618	1782	2541
BspH I	(4)	5098	6106	6211	6735			2739	2798	2820	3104
BspM II	(1)	2310						3111	3162	3165	3259
Bsr I	(20)	374	1030	1058	1546			4203	4284	4302	4305
		1570	1871	2437	2458			4423	4578	4721	4786
		3192	3729	4179	4785			4789	4995	5323	5662
		4798	4912	5318	5436			5689	5784	6013	6300
		5479	5746	5918	6624			6409	6513	6586	
BsrB I	(5)	1042	1851	4070	4311	Fok I	(13)	423	1140	1300	1464
		6112						1699	2607	2973	3777
BsrD I	(2)	5332	5506					5237	5418	5705	6348
BsrG I	(1)	1282						6592			
BssH II	(5)	317	368	1896	3672	Fsp I	(2)	5493	6516		
		3723				Gdi II	(11)	723	725	1129	1528
BssS I	(5)	1587	2760	4551	5935			1530	2652	3039	4218
		6242						5658	6694	6826	
BstB I	(1)	1828				Gsu I	(5)	3023	3263	3322	5342
BstE II	(1)	1087						6960			
BstN I	(23)	241	548	735	748	Hae I	(6)	584	1111	1416	4393
		1169	1194	1217	1237			4404	4856		
		1267	1717	1909	2625	Hae II	(11)	925	1401	1947	2565
		2750	2862	2937	2991			4256	4626	6497	6680
		3596	4118	4406	4527			6734	6794	6815	
		4540	6613	6890		Hae III	(33)	129	173	584	726
BstU I	(26)	317	319	370	903			762	774	987	1111
		905	1135	1644	1864			1132	1197	1416	1531
		1898	1900	2424	2908			1868	2286	2653	2753
		3226	3672	3674	3725			3042	3154	3483	3528
								4219	4393	4404	4422

		4856	5314	5394	5661		1173	1814	1823	2046	
		6248	6546	6695	6703		2125	2138	2218	3302	
		6827					3306	3389	3923	4021	
Hga I	(10)	381	642	895	1491		4149	4208	5084	5136	
		3736	4488	5066	5798		5141	5155	5208	5443	
		6356	6825				5482	5847	6219	6400	
HgiA I	(7)	338	3189	3693	4696		6597	6867			
HgiE II	(2)	4957	5942	6439		Msl I	(4)	1993	3066	5523	5682
Hha I	(46)	319	321	370	372	Msp I	(32)	163	398	569	630
		648	905	924	1400		723	1401	2311	2328	
		1512	1898	1900	1906		2403	2568	2608	2671	
		1946	2352	2424	2564		2731	3262	3518	3753	
		2869	2910	3226	3674		4096	4585	4732	4758	
		3676	3725	3727	4002		4948	5352	5386	5453	
		4162	4227	4255	4288		5563	5805	6306	6340	
		4558	4625	4725	4899		6692	6815	6824	6839	
		5008	5401	5494	5831	MspA1 I	(11)	588	1620	2554	3854
		6163	6263	6366	6496		4202	4720	4965	5906	
		6517	6650	6679	6733		6372	6566	6922		
		6793	6814			Mun I	(1)	2395			
Hinf I	(18)	442	450	791	802	Nae I	(2)	631	6825		
		1437	1769	1859	2017	Nar I	(6)	1398	1944	6494	6677
		2270	3913	3935	3956		6791	6812			
		4213	4278	4353	4749	Nci I	(16)	163	398	399	570
		5266	6904				723	2609	3263	3518	
HinI I	(10)	742	1398	1838	1944		3753	3754	4758	5454	
		5808	6190	6494	6677		5805	6306	6341	6692	
		6791	6812			Nco I	(2)	1411	2578		
HinP I	(46)	317	319	368	370	Nde I	(2)	2110	6442		
		646	903	922	1398	NgoM I	(2)	629	6823		
		1510	1896	1898	1904	Nhe I	(2)	30	3383		
		1944	2350	2422	2562	Nla III	(25)	62	1205	1415	1964
		2867	2908	3224	3672		2277	2582	2816	2846	
		3674	3723	3725	4000		3041	3236	3281	3415	
		4160	4225	4253	4286		3869	4036	4382	5102	
		4556	4623	4723	4897		5593	5603	5681	5717	
		5006	5399	5492	5829		6110	6215	6299	6666	
		6161	6261	6364	6494	Nla IV	(39)	188	201	246	403
		6515	6648	6677	6731		521	552	692	761	
		6791	6812				893	1144	1221	1399	
Hpa II	(32)	163	398	569	630		1573	1689	1868	1945	
		723	1401	2311	2328		2309	2332	2407	2459	
		2403	2568	2608	2671		2617	3155	3543	3556	
		2731	3262	3518	3753		3601	3758	4124	4410	
		4096	4585	4732	4758		4449	5221	5315	5356	
		4948	5352	5386	5453		5567	6157	6495	6678	
		5563	5805	6306	6340		6702	6792	6813		
		6692	6815	6824	6839	Nru I	(1)	1644			
Hph I	(19)	1157	1183	1401	1978	Nsp7524 I	(4)	1960	4378	6295	6662
		2593	2598	2926	2950	NspB II	(11)	588	1620	2554	3854
		3079	5115	5342	5756		4202	4720	4965	5906	
		5964	5997	6281	6290		6372	6566	6922		
		6783	6828	6909		NspH I	(4)	1964	4382	6299	6666
Kas I	(6)	1397	1943	6493	6676	PacI	(1)	3304			
		6790	6811			Pal I	(33)	129	173	584	726
Kpn I	(2)	405	3760				762	774	987	1111	
Mae I	(15)	31	219	661	673		1132	1197	1416	1531	
		1394	1522	1879	2410		1868	2286	2653	2753	
		2559	3384	3574	4873		3042	3154	3483	3528	
		5126	5461	6896			4219	4393	4404	4422	
Mae II	(16)	742	1081	1122	1838		4856	5314	5394	5661	
		1994	2175	2180	2517		6248	6546	6695	6703	
		2644	2857	3028	5081		6827				
		5497	5870	6190	6632	Fle I	(3)	799	3921	4757	
Mae III	(23)	39	994	1022	1087	Fme I	(1)	1815			
		1231	1874	2097	2176	PpuM I	(4)	254	1219	1572	3609
		2762	3251	3392	3870	Psp1406 I	(3)	2517	5497	5870	
		4734	4797	4913	5196	PspA I	(2)	397	3752		
		5527	5585	5738	5926	Pst I	(3)	921	929	1105	
		6314	6605	6625		Pvu I	(4)	1648	2323	5641	6537
Mbo I	(30)	92	457	1044	1176	Pvu II	(5)	1620	2554	4202	6566
		1361	1645	1706	1720		6922				
		2307	2313	2320	3079	Rsa I	(11)	403	659	1284	2190
		3227	3265	3446	4944		2453	3009	3291	3758	
		5019	5030	5038	5116		5751	6427	6938		
		5128	5233	5574	5592	Sac I	(2)	338	3693		
		5638	5896	5913	5949	Sap I	(2)	4000	4262		
		6534	6758			Sau3A I	(30)	92	457	1044	1176
Mbo II	(20)	1086	1299	1445	1502		1361	1645	1706	1720	
		1530	1927	2283	2823		2307	2313	2320	3079	
		2868	2871	3064	4016		3227	3265	3446	4944	
		4250	5039	5112	5867		5019	5030	5038	5116	
		5945	6054	6542	6754		5128	5233	5574	5592	
Mme I	(6)	829	1106	3817	3869		5638	5896	5913	5949	
		4592	4776				6534	6758			
Mnl I	(58)	167	216	368	387	Sau96 I	(32)	127	171	186	199
		432	460	484	506		254	551	690	760	
		510	566	776	872		773	985	1196	1219	
		1168	1271	1304	1310		1572	1866	1867	2285	
		1313	1319	1349	1352		2752	3153	3236	3481	
		1367	1374	1390	1454		3526	3541	3554	3609	
		1795	1796	1921	2297		5313	5392	5409	5631	
		2378	2467	2588	2669		6247	6544	6701	6886	
		2675	2768	2906	2918	Sca I	(1)	5751			
		2969	3089	3571	3723	ScrF I	(39)	163	241	398	399
		3742	3787	3814	3838		548	570	723	735	
		3898	4227	4276	4486		748	1169	1194	1217	
		4559	4810	5210	5291		1237	1267	1717	1909	
		5437	5643	6238	6296		2609	2625	2750	2862	
		6556	6840				2937	2991	3263	3518	
Msc I	(3)	584	1111	1416			3596	3753	3754	4118	
Mse I	(30)	36	1007	1119	1149		4406	4527	4540	4758	

*pSOK Vector*

		5454	5805	6306	6341			1854	2303	2578	3605
		6613	6692	6890				3815			
Sec I	(40)	161	240	250	397	SwaI	(1)	2047			
		461	497	546	547	Taq I	(16)	794	1038	1349	1360
		694	733	734	746			1705	1828	2083	2629
		747	1167	1236	1245			2923	2950	2965	3094
		1265	1266	1411	1716			3297	3916	4478	5922
		1854	1907	2303	2472	Tfi I	(10)	442	802	1437	1769
		2578	2608	2748	2911			1859	2017	2270	3956
		2935	2990	3516	3595			4213	4353		
		3605	3752	3815	4117	Tsp45 I	(10)	1022	1231	1874	2176
		4538	6612	6690	6696			2762	3251	5527	5738
SfaN I	(24)	185	444	641	938			6314	6625		
		1485	1542	1674	1720	Tth111 I	(4)	390	811	1217	3745
		2679	2955	2970	3069	Tth111 II	(9)	146	422	613	632
		3540	3798	4474	5526			3062	3500	4967	4976
		5719	5966	6327	6419			5006			
		6457	6495	6811	6823	Vsp I	(5)	2218	4021	4149	4208
Sfc I	(10)	15	917	925	1101			5443			
		1496	3368	4643	4834	Xba I	(2)	218	3573		
		5512	6930			Xho II	(9)	2307	2313	3079	5019
Sma I	(2)	399	3754					5030	5116	5128	5896
Spe I	(1)	660						5913			
Sph I	(1)	6666				Xma I	(2)	397	3752		
Ssp I	(3)	1955	3921	6075		Xma III	(2)	724	1529		
Sty I	(9)	250	461	1245	1411	Xmn I	(3)	3984	4022	5870	