

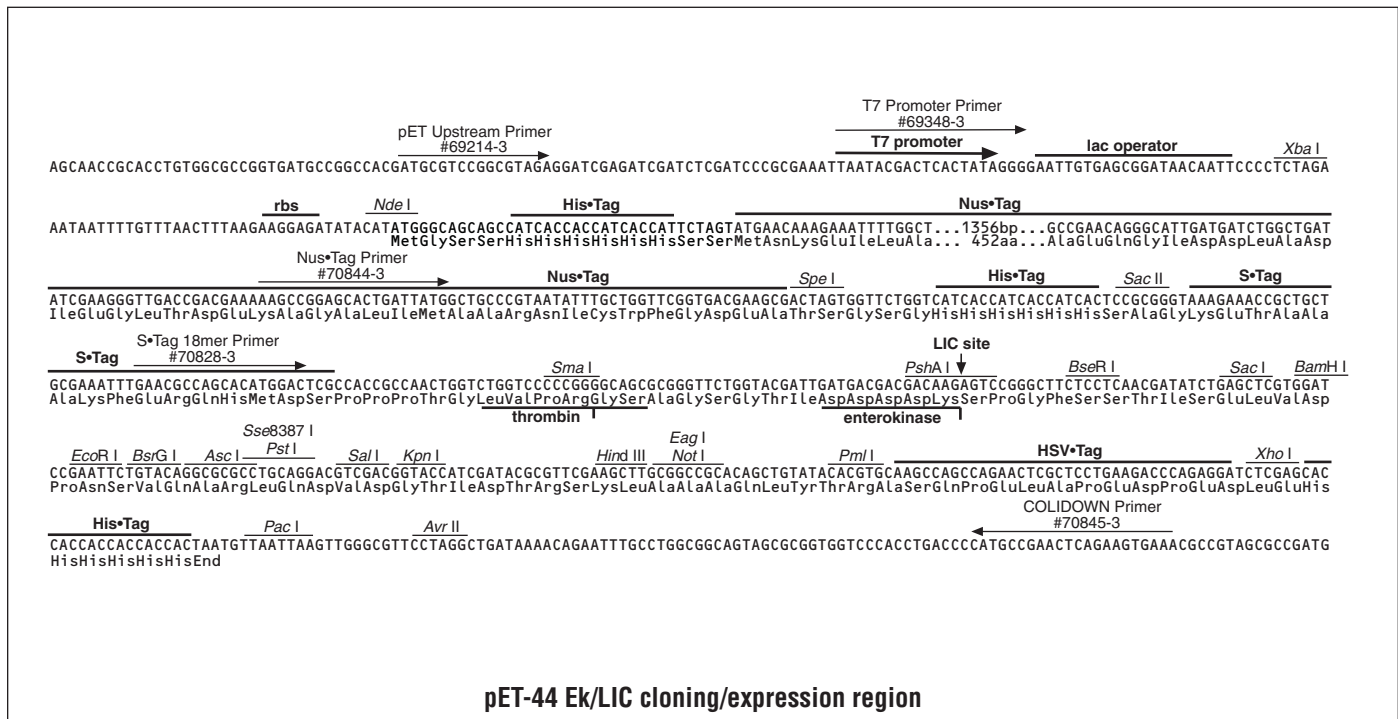
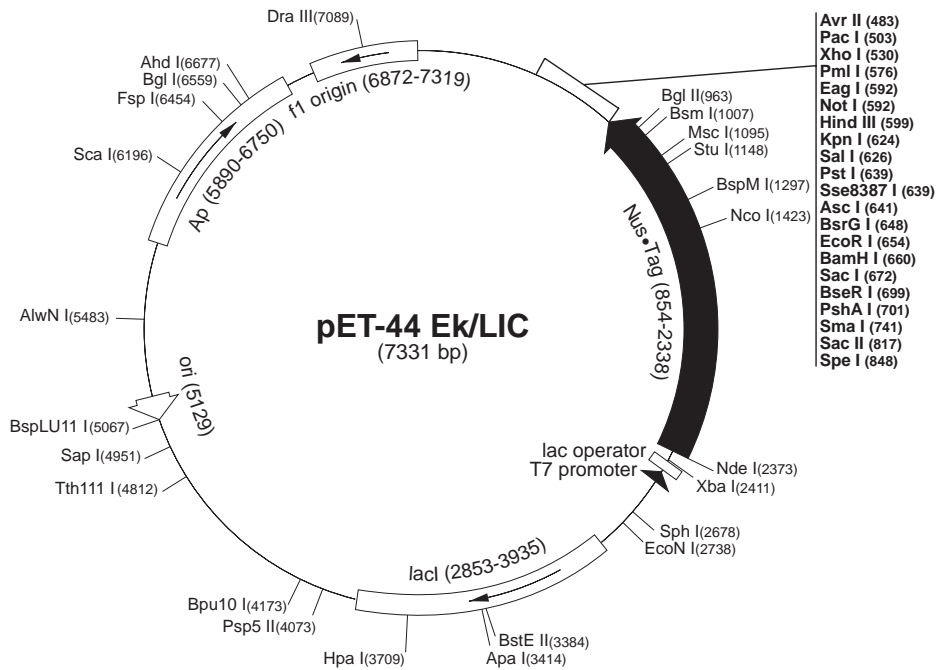
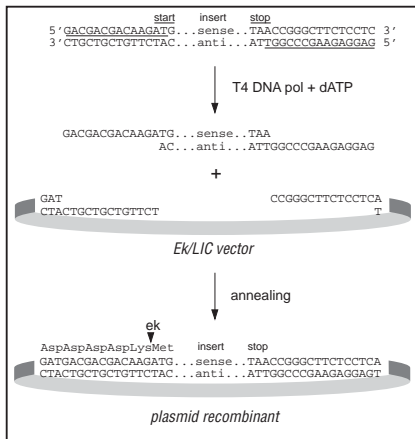
pET-44 Ek/LIC Vector

	Cat. No.
pET-44 Ek/LIC DNA	71143-3

pET-44 Ek/LIC sequence landmarks

T7 promoter	2446-2462
T7 transcription start	2445
His•Tag coding sequence	2345-2362
Nus•Tag coding sequence	854-2338
His•Tag coding sequence	821-838
S•Tag coding sequence	767-811
Multiple cloning sites (<i>BseR I-Xho I</i>)	699-530
HSV•Tag coding sequence	536-571
His•Tag coding sequence	512-529
T7 terminator	26-73
<i>lacI</i> coding sequence	2853-3932
pBR322 origin	5129
Ap (<i>bla</i>) coding sequence	5890-6750
f1 origin	6872-7319

The pET-44 Ek/LIC Vector is prepared for rapid, directional cloning of PCR-amplified DNA for high-level expression of polypeptides fused with N-terminal His•Tag[®], Nus•Tag[™] and S•Tag[™] sequences. Using specifically designed primers for amplification and the pET-44 Ek/LIC Vector Kit (Cat. No. 71144-3), inserts can be efficiently cloned without the need for restriction digestion or ligation. Unique sites are shown on the circle map. Note that the sequence is numbered by the pBR322 convention, so the T7 expression region is reversed on the circle map. The cloning/expression region of the coding strand transcribed by T7 RNA polymerase is shown below. The f1 origin is oriented so that infection with helper phage will produce virions containing single stranded DNA that corresponds to the coding strand. Therefore, single stranded sequencing should be performed using the COLIDOWN Primer (Cat. No. 70845-3). Vector encoded sequences can be completely removed when cloning into the Ek/LIC site (as shown below left) by cleaving the fusion protein with enterokinase.



pET-44 Ek/LIC cloning/expression region

pET-44 Ek/LIC Restriction Sites

Enzyme	# Sites	Locations	Enzyme	# Sites	Locations	Enzyme	# Sites	Locations			
AatII	2	633 2208	DpnI	42		Sau96I	20				
AccI	4	580 627 1549 4837	DraI	3	6099 6791 6810	Scal	1	6196			
AcI	9		DraIII	1	7089	ScrFI	33				
AflIII	4	575 608 3203 5067	DrdI	3	4760 5175 7044	SfaNI	23				
AhdI	1	6677	Dsal	4	814 984 1423 2640	Sfcl	7	635 2314 2445 5332 5523			
AluI	27		EaeI	8	592 991 1093 1380 2511			6431 7308			
Alw26I	8	371 2900 3305 3431 3818			2643 3877 6284	SgrAI	2	1919 2522			
		4708 5834 6611	EagI	1	592	SmaI	1	741			
AlwI	20		EarI	3	2821 4951 5878	SpeI	1	848			
AlwNI	1	5483	Ecl136II	1	670	SphI	1	2678			
Apal	1	3414	Eco47III	2	2608 4321	Sse8387I	1	639			
ApaLI	4	3183 4881 5381 6005	Eco57I	4	532 1846 5615 6011	SspI	2	877 6881			
ApoI	7	466 654 788 2323 3478	EcoNI	1	2738	StuI	1	1148			
		6891 6902	EcoO109I	3	53 2636 4073	StyI	4	57 483 1423 2119			
Ascl	1	641	EcoRI	1	654	Tail	25				
Aval	2	530 739	EcoRII	13		TaqI	28				
Avall	7	440 745 3755 4073 4352	EcoRV	3	677 931 1522	TfiI	4	3882 4117 4621 5042			
		6313 6535	EheI	4	2527 2548 2662 3844	ThaI	52				
AvrII	1	483	FauI	20		TseI	30				
BamHI	1	660	Fnu4HI	54		Tsp45I	10	859 1745 2132 3384 4506			
BanI	10	620 1470 2525 2546 2660	FokI	14				4719 4814 6204 6415 7262			
		3123 3842 3972 6724 7126	FspI	1	6454	Tsp509I	23				
BanII	5	672 2587 2601 3414 7164	HaeII	15		TspRI	16				
BbsI	5	538 1228 3349 3688 4185	HaeIII	31		Tth111I	1	4812			
BbvI	30		Hgal	15		VspI	4	2460 3888 3947 6502			
Bcgl	3	3529 4644 6139	Hhal	59		XbaI	1	2411			
BclI	2	1953 3217	HincII	4	628 920 1315 3709	XcmI	3	3059 3575 3593			
Bfal	11		HindIII	1	599	XhoI	1	530			
BglI	1	6559	Hinfl	16		XmnI	3	2271 4625 6077			
BglII	1	963	HpaI	1	3709						
Bpml	4	3041 3530 4594 6608	HphI	31		Enzymes that do not cut pET-44 Ek/LIC:					
Bpu10I	1	4173	KpnI	1	624	AflII	Bsu36I	FseI	MunI	NheI	NsiI
Bpu1102I	2	80 1298	MaellI	22		PinAI	PmeI	RsrII	SanDI	SexAI	SfiI
BsaAI	3	576 4819 7089	MbolI	25		Sgfl	SnaBI	SrfI	SunI	Swal	
BsaBI	3	2476 2486 4264	Mlul	2	608 3203						
BsaHI	9	200 630 2205 2526 2547	MnlI	29							
		2661 3160 3843 6137	MscI	1	1095						
Bsal	2	371 6611	MseI	33							
BsaJI	17		MslI	11							
BsaWI	8	2 1596 3522 4025 4256	MspA1I	13							
		5273 5420 6381	MspI	42							
BseRI	1	699	Mwol	47							
BsgI	4	1976 3054 3254 4227	NarI	4	2526 2547 2661 3843						
BsiEI	10	595 1135 1273 1809 2190	NciI	20							
		3988 4983 5407 6159 6308	NcoI	1	1423						
BsiHKAI	11		NdeI	1	2373						
BsII	26		NgoAIV	2	2513 7190						
BsmBI	2	3818 4708	NlaIII	28							
BsmFI	5	453 758 2664 4338 7304	NlaIV	25							
BsmI	1	1007	NotI	1	592						
Bsp1286I	15		Nrul	3	1230 1655 2279						
BspEI	2	2 4256	NspI	4	2678 4412 4704 5071						
BspLU11I	1	5067	NspV	2	604 2265						
BspMI	1	1297	Pacl	1	503						
BsrBI	6	249 1343 2432 5000 5834	PfIMI	2	778 2785						
		7233	PleI	12							
BsrDI	6	1093 1474 3250 3616 6443	PmlI	1	576						
		6617	PshAI	1	701						
BsrFI	9	1047 1403 1467 1919 2513	Psp1406I	6	237 2865 4392 6075 6448						
		2522 2889 6592 7190			6874						
BsrGI	1	648	Psp5II	1	4073						
BsrI	29		PstI	1	639						
BssHII	5	641 1567 1759 2248 3614	PvuI	3	1273 2190 6308						
BssSI	4	665 1639 5240 6008	PvuII	4	586 3803 3896 4658						
Bst1107I	2	581 4838	RcaI	4	2601 5787 5836 5868						
BstEII	1	3384	RsaI	8	622 650 721 976 2198						
BstXI	3	3005 3134 3257			3350 4873 6196						
BstYI	13		SacI	1	672						
Cac8I	52		SacII	1	817						
Clal	5	615 1524 1602 2190 2480	Sall	1	626						
CviJI	114		Sapl	1	4951						
Ddel	15		Sau3AI	42							