

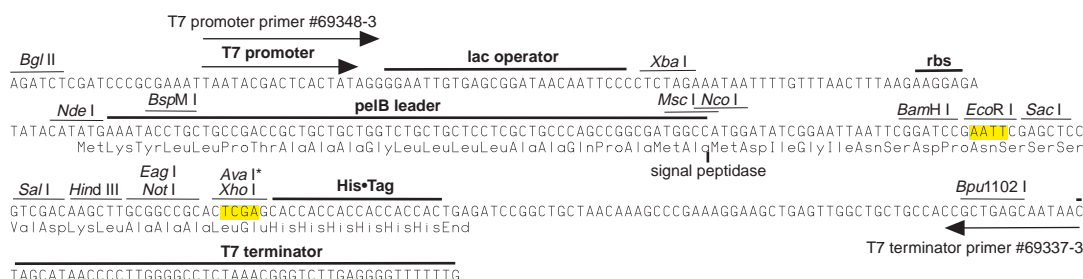
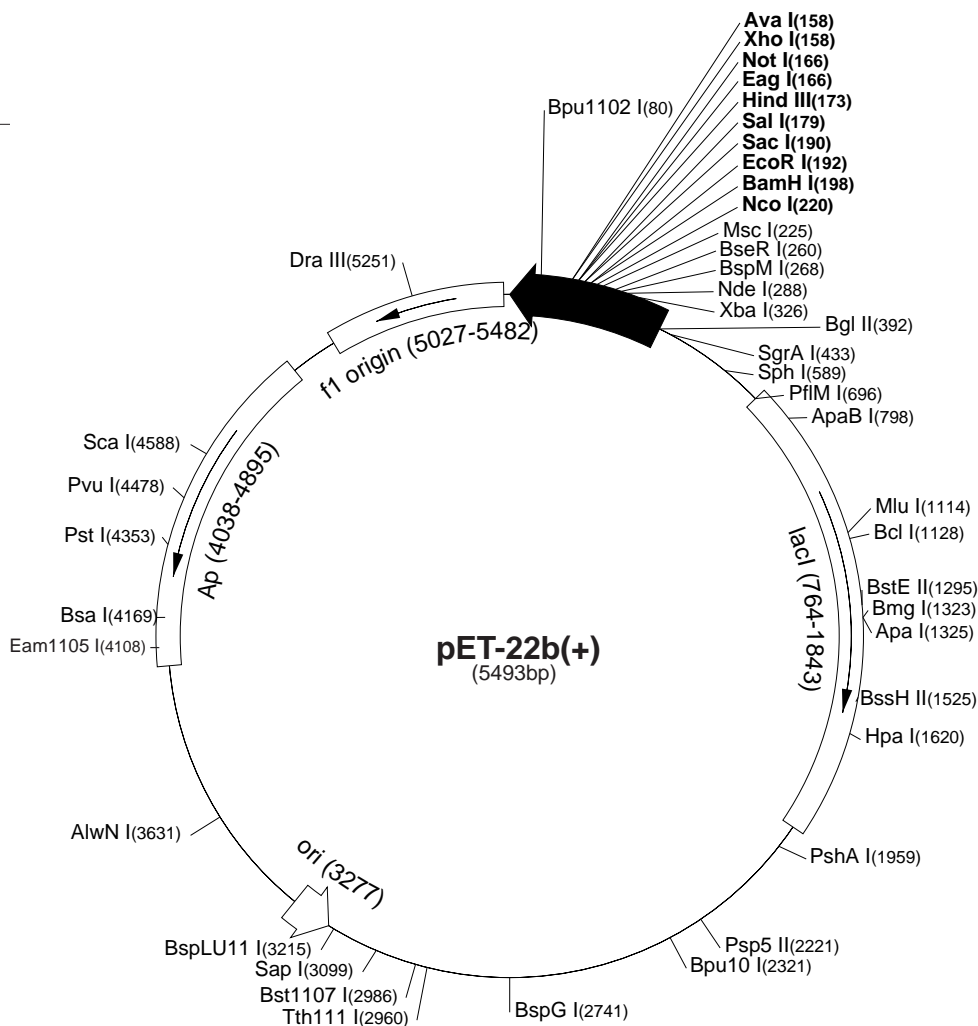
pET-22b(+) Vector

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The pET-22b(+) vector (Cat. No. 69744-3) carries an N-terminal *pelB* signal sequence for potential periplasmic localization, plus optional C-terminal His•Tag® sequence. 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 circular 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 T7 terminator primer (Cat. No. 69337-3).

pET-22b(+) sequence landmarks

T7 promoter	361-377
T7 transcription start	360
<i>pelB</i> coding sequence	224-289
Multiple cloning sites	
(<i>Nco</i> I - <i>Xho</i> I)	158-225
His•Tag coding sequence	140-157
T7 terminator	26-72
<i>lacI</i> coding sequence	764-1843
pBR322 origin	3277
<i>bla</i> coding sequence	4038-4895
f1 origin	5027-5482



pET-22b(+) cloning/expression region

pET-22b(+) Restriction Sites

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Enzyme	# Sites	Locations	Enzyme	# Sites	Locations	Enzyme	# Sites	Locations
AccI	2	180 2985	BsrFI	8	231 424 433 800 2012	NotI	1	166
AccIII	8	881 1609 1940 2724 2865			2172 4188 5352	NspI	4	589 2560 2852 3219
		3167 4407 5091	BssHII	1	1525	Pfi1108I	2	2001 4126
Acil	81		Bst1107I	1	2986	PfiMI	1	696
AflIII	2	1114 3215	BstEII	1	1295	PleI	9	375 663 750 1546 3109
AluI	25		BstXI	3	916 1045 1168			3594 4097 5186 5194
AlwI	16		BstYI	12		PshAI	1	1959
Alw21I	9	159 190 614 1098 2209	Cac8I	39		Psp5II	1	2221
		3033 3533 4694 4779	CjeI	24		Psp1406I	6	776 2144 2540 4334 4707
Alw44I	4	1094 3029 3529 4775	CjePI	18				5036
AlwNI	1	3631	CviJI	87		PstI	1	4353
Apal	1	1325	CviRI	24		PvuI	1	4478
ApaBI	1	798	DdeI	11		PvuII	3	1714 1807 2806
ApoI	4	192 1389 5053 5064	DpnI	27		RcaI	3	512 3935 4943
AvaI	1	158	DraI	3	3974 3993 4685	RsaI	3	1261 3021 4588
Avall	7	1666 2042 2130 2221 2500	DraIII	1	5251	SacI	1	190
		4246 4468	DrdI	3	2908 3323 5206	Sall	1	179
BamHI	1	198	DrdII	2	837 5256	SapI	1	3099
BanI	9	436 457 571 1034 1753	DsaI	3	220 551 2187	Sau96I	18	
		1883 2009 4056 5288	EaeI	6	166 223 422 554 1788	Sau3AI	27	
BanII	5	190 498 512 1325 5326			4496	Scal	1	4588
BbsI	4	1260 1599 1973 2333	EagI	1	166	ScrFI	19	
BbvI	31		Eam1105I	1	4108	SfaNI	20	
BccI	14		EarI	3	732 3099 4903	Sfcl	5	360 3480 3671 4349 5470
Bce83I	7	21 1928 2098 3306 3604	EcII	4	891 3289 3435 4263	SgrAI	1	433
		3845 4713	Eco47III	3	519 2020 2469	SphI	1	589
Bcefl	5	633 974 1601 3717 5277	Eco57I	2	3763 4775	Sspl	2	4912 5043
Bcgl	10	160 194 1406 1440 1940	EcoNI	1	649	Styl	2	57 220
		1974 2792 2826 4613 4647	EcoO109I	3	53 547 2221	TaqI	13	
BclI	1	1128	EcoRI	1	192	TaqII	9	1022 1240 1913 3117 4456
Bfal	7	70 327 2229 3710 3963	EcoRII	7	837 1152 1692 1749 3241			4641 4794 4811 5155
		4298 5402			3362 3375	TfiI	5	1793 2095 2265 2769 3190
BglI	3	237 2178 4228	EcoRV	2	217 1564	Thal	34	
BglII	1	392	FauI	17		TseI	31	
BmgI	1	1323	FokI	10	1160 1169 2434 2496 2574	Tsp45I	8	1295 2123 2654 2867 2962
BpmI	5	952 1441 2075 2742 4178			2760 2901 4074 4255 4542			4364 4575 5424
Bpu10I	1	2321	Fspl	2	2196 4330	Tsp509I	18	
Bpu1102I	1	80	GdIII	5	166 422 554 1788 4496	Tth111I	1	2960
BsaI	1	4169	HaeI	6	225 842 2163 3230 3241	Tth111II	6	953 1646 2676 3805 3812
BsaAI	2	2967 5251			3693			3844
BsaBI	3	391 397 2412	HaeII	14		UbaII	20	
BsaHI	6	437 458 572 1071 1754	HaeIII	25		VspI	5	207 375 1799 1858 4280
		4645	HgaI	12		XbaI	1	326
BsaJI	7	57 220 551 557 1749	HgiEII	2	712 3801	XcmI	3	970 1486 1504
		2187 3375	HhaI	45		XhoI	1	158
BsaWI	7	2 1433 1936 2404 3421	Hin4I	3	1013 4107 4181	XmnI	3	208 2773 4707
		3568 4399	HincII	2	181 1620			
BsaXI	2	1773 5199	HindIII	1	173			
Bsbl	2	2931 5158	Hinfl	14				
BscGI	13		HpaI	1	1620			
BseRI	1	260	HphI	16				
BsgI	3	965 1165 2375	Maell	15				
Bsil	2	3388 4772	MaellI	18				
BsiEI	7	169 271 1899 3131 3555	MbolI	14				
		4478 4627	MluI	1	1114			
BsII	20		MmeI	3	3430 3614 5228			
BsmAI	7	811 1216 1342 1729 2856	MnII	27				
		4169 4945	MscI	1	225			
BsmBI	2	1729 2856	MseI	29				
BsmFI	4	575 2116 2486 5466	MslI	9	1166 1454 1484 2202 2397			
BsoFI	51				2788 4360 4519 4878			
Bsp24I	10	404 436 955 987 1257	MspI	32				
		1289 3708 3740 3886 3918	MspAII	10	84 267 1144 1714 1807			
Bsp1286I	13				2806 2925 3557 3802 4743			
BspEI	2	2 2404	Mwol	39				
BspGI	1	2741	NarI	4	437 458 572 1754			
BspLU11I	1	3215	NciI	12				
BspMI	1	268	NcoI	1	220			
BsrI	25		NdeI	1	288			
BsrBI	4	347 3148 4949 5395	NgoAIV	5	231 424 2012 2172 5352			
BsrDI	4	1161 1527 4169 4343	NlaIII	25				
			NlaIV	25				

Enzymes that do not cut pET-22b(+):

AatII	AflII	AgeI	AscI	AvrII
BaeI	BsmI	BsrGI	Bsu36I	Clal
FseI	KpnI	MunI	NheI	NruI
NsiI	NspV	PacI	PmeI	PmlI
RleAI	RsrII	SacII	SexAI	SfiI
SgfI	Smal	SnaBI	SpeI	SrfI
Sse8387I	StuI	SunI	Swal	