

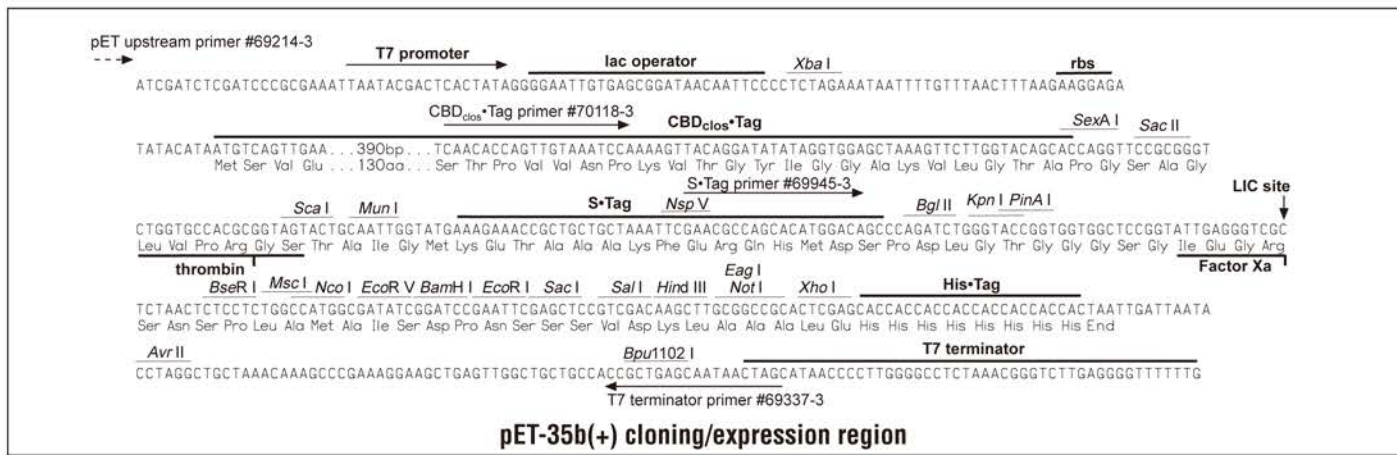
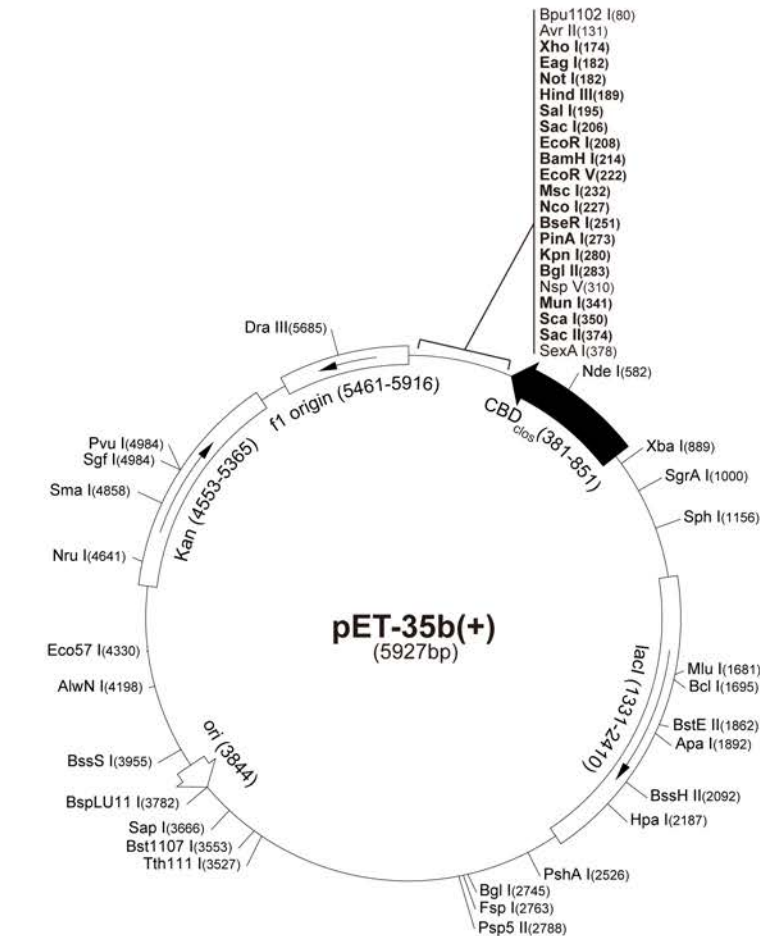
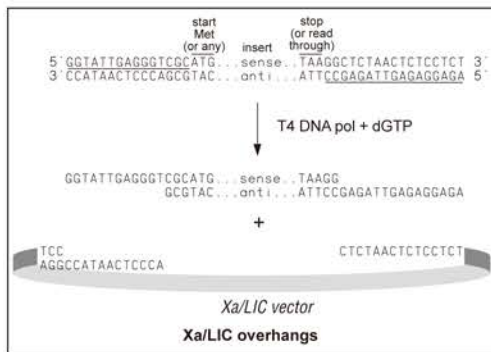
pET-35b(+) Vector

The pET-35b(+)⁺ (Cat. No. 70103-3) vector is designed for expression of CBD_{clos} fusion proteins. A variety of cloning sites and strategies are available. Rapid directional cloning of PCR-amplified DNA is available with the pET-35 Xa/LIC Vector Kit (Cat. No. 70115-3), which contains linearized pET-35b(+)⁺ ready for annealing with appropriately prepared inserts. When cloned with the Xa/LIC method, resulting CBD_{clos}•TagTM fusion proteins can be cleaved precisely at the vector-encoded junction using Factor Xa. 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 T7 terminator primer (Cat. No. 69337-3).

pET-35b(+) sequence landmarks

T7 promoter	924-940
T7 transcription start	923
CBD _{clos} •Tag coding seq.	381-851
S•Tag coding sequence	291-335
Multiple cloning sites (<i>BseR</i> I - <i>Xho</i> I)	174-241
His•Tag coding sequence	150-173
T7 terminator	26-72
<i>lacI</i> coding sequence	1331-2410
pBR322 origin	3844
Kan coding sequence	4553-5365
f1 origin	5461-5916

Note: Primer sequence extensions required for Xa/LIC compatibility are underlined in the diagram below.



pET-35b(+) cloning/expression region

pET-35b(+) Restriction Sites

Enzyme	# Sites	Locations
AccI	2	196 3552
AccI	78	
AflIII	2	1681 3782
AluI	25	
AlwI	12	
Alw26I	6	1378 1783 1909 2296 3423
		5000
AlwNI	1	4198
ApaI	1	1892
ApaLI	3	1661 3596 4096
ApoI	10	208 312 573 754 837
		1956 4597 4781 5487 5498
AvaI	2	174 4856
AvaII	5	2233 2609 2697 2788 3067
AvrII	1	131
BamHI	1	214
BanI	10	276 361 1003 1024 1138
		1601 2320 2450 2576 5722
BanII	6	206 1065 1079 1892 4639
		5760
BbsI	4	1827 2166 2540 2900
BbvI	26	
BcgI	4	210 1973 2541 3393
BcgI'	4	176 2007 2507 3359
BclI	1	1695
BfaI	8	70 132 643 890 2796
		4277 4584 5836
BglI	1	2745
BglII	1	283
BpmI	4	1519 2008 2642 3309
Bpu10I	2	2888 5001
Bpu1102I	1	80
BsaAI	2	3534 5685
BsaBI	3	954 964 2979
BsaHI	5	1004 1025 1139 1638 2321
BsaJI	12	
BsaWI	9	2 261 273 2000 2503
		2971 3988 4135 5119
BseRI	1	251
BsgI	3	1532 1732 2942
BsiEI	5	185 2466 3698 4122 4984
BsiHKAI	7	175 206 1181 1665 2776
		3600 4100
BsII	26	
BsmI	2	4868 4945
BsmBI	3	2296 3423 5000
BsmFI	4	1142 2683 3053 5900
Bsp1286I	12	
BspEI	2	2 2971
BspLU11I	1	3782
BsrI	23	
BsrBI	4	910 3715 5383 5829
BsrDI	2	1728 2094
BsrFI	8	273 991 1000 1367 2579
		2739 4938 5786
BssHII	1	2092
BssSI	1	3955
Bst1107I	1	3553
BstEII	1	1862
BstXI	3	1483 1612 1735
BstYI	8	214 283 1245 2457 2974
		4423 4434 5233
Cac8I	42	
ClaI	2	958 4675
CviJI	89	
DdeI	10	80 101 2253 2888 3050
		3590 4057 4466 5001 5365
DpnI	22	
DraI	2	753 759
DraIII	1	5685
DrdI	3	3475 3890 5640
DsaI	4	227 371 1118 2754

Enzyme	# Sites	Locations
EaeI	5	182 230 989 1121 2355
EagI	1	182
EarI	3	1299 3666 4797
Eco47III	3	1086 2587 3036
Eco57I	1	4330
EcoNI	2	1216 4896
EcoO109I	3	53 1114 2788
EcoRI	1	208
EcoRII	10	378 1404 1719 2259 2316
		3808 3929 3942 4872 5229
EcoRV	1	222
FauI	18	
Fnu4HI	44	
FokI	9	1727 1736 3001 3063 3141
		3327 3468 4622 5228
FspI	1	2763
HaeII	14	
HaeIII	24	
HgaI	11	
HhaI	46	
HincII	2	197 2187
HindIII	1	189
HinfI	18	
HpaI	1	2187
HphI	16	
KpnI	1	280
MaeIII	19	
MbolI	13	
MluI	1	1681
MnII	27	
MscI	1	232
MseI	30	
MslI	7	681 1733 2021 2051 2769
		2964 3355
MspI	30	
MspA1I	10	84 325 373 1711 2281
		2374 3373 3492 4124 4369
MunI	1	341
MwoI	39	
NarI	4	1004 1025 1139 2321
NciI	12	
NcoI	1	227
NdeI	1	582
NgoAIV	4	991 2579 2739 5786
NlaIII	27	
NlaIV	25	
NotI	1	182
NruI	1	4641
NsiI	2	4834 5100
NspI	4	1156 3127 3419 3786
NspV	1	310
PIIMI	3	302 1263 5247
PinAI	1	273
PleI	9	938 1230 1317 2113 3676
		4161 5216 5620 5628
PshAI	1	2526
Psp1406I	4	1343 2711 3107 5470
Psp5II	1	2788
PvuI	1	4984
PvuII	3	2281 2374 3373
RcaI	3	1079 4502 5377
RsaI	7	278 350 389 713 1828
		3588 4819
SacI	1	206
SacII	1	374
SalI	1	195
SapI	1	3666
Sau3AI	22	
Sau96I	14	
Scal	1	350
ScrFI	22	
SexAI	1	378

Enzyme	# Sites	Locations
SfaNI	24	
SfcI	4	923 4047 4238 5904
SgfI	1	4984
SgrAI	1	1000
SmaI	1	4858
SphI	1	1156
SspI	2	4909 5477
StyI	3	57 131 227
Swal	2	753 759
TalI	15	
TaqI	17	
TfiI	9	2360 2662 2832 3336 3757
		4895 4951 5123 5214
Thal	37	
TseI	26	
Tsp45I	9	633 686 1862 2690 3221
		3434 3529 5131 5858
Tsp509I	29	
TspRI	13	
Tth111I	1	3527
VspI	6	139 938 2366 2425 5183
		5372
XbaI	1	889
XcmI	3	1537 2053 2071
XhoI	1	174
XmnI	2	3340 5373

Enzymes that do not cut pET-35b(+):

AatII	AflIII	AhdI	AscI	BsaI
BspMI	BsrGI	Bsu36I	FseI	NheI
Pacl	PmeI	PmlI	PstI	RsrII
SanDI	SfiI	SnaBI	SpeI	SrfI
Sse8387I	StuI	SunI	UbaEI	