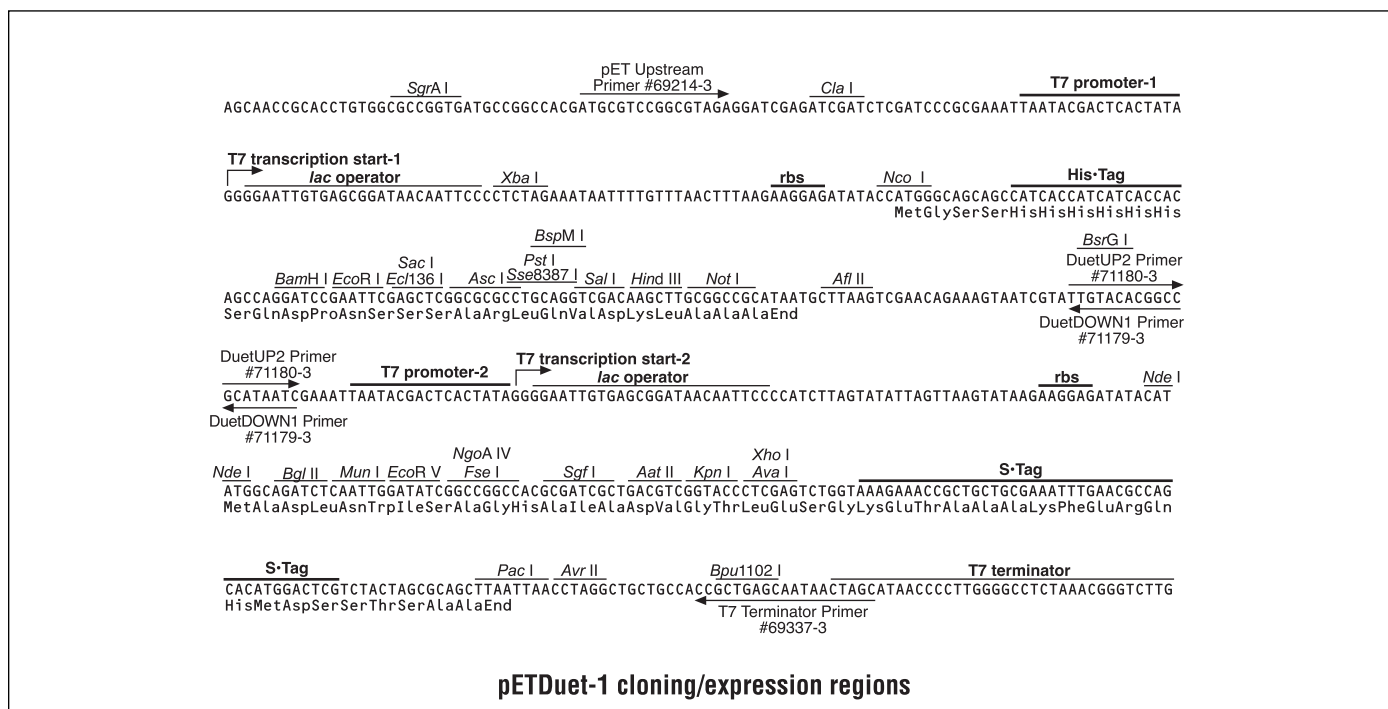
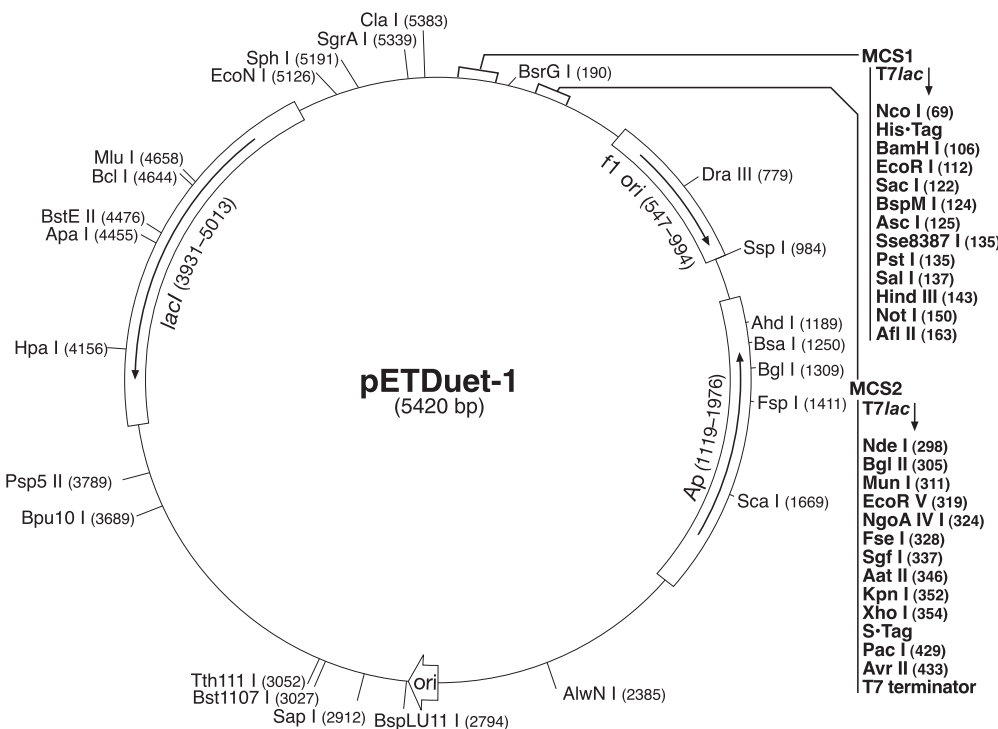


pETDuet-1 Vector

	Cat. No.
pETDuet-1 DNA	71146-3
pETDuet-1 sequence landmarks	
T7 promoter-1	5404–5420
T7 transcription start-1	1
His•Tag [®] coding sequence	83–100
Multiple cloning sites-1 (<i>Nco</i> I– <i>Afl</i> II)	69–168
T7 promoter-2	214–230
T7 transcription start-2	231
Multiple cloning sites-2 (<i>Nde</i> I– <i>Avr</i> II)	297–438
S•Tag [™] coding sequence	366–410
T7 terminator	462–509
<i>lacI</i> coding sequence	3931–5013
pBR322 origin	2737
<i>bla</i> (Ap) coding sequence	1119–1976
f1 origin	547–994

pETDuet[™]-1 is designed for the coexpression of two target genes. The vector contains two multiple cloning sites (MCS), each of which is preceded by a T7 promoter/*lac* operator and a ribosome binding site (rbs). The vector also carries the pBR322-derived ColE1 replicon, *lacI* gene and ampicillin resistance gene. This vector can be used in combination with pACYCDuet[™]-1 (Cat. No. 71147-3) in an appropriate host strain for the coexpression of up to 4 target genes. Genes inserted into MCS1 can be sequenced using the pET Upstream Primer (Cat. No. 69214-3) and DuetDOWN1 Primer (Cat. No. 71179-3). Genes inserted into MCS2 can be sequenced using the DuetUP2 Primer (Cat. No. 71180-3) and T7 Terminator Primer (Cat. No. 69337-3).



pETDuet-1 cloning/expression regions

pETDuet-1 Restriction Sites

Enzyme	# Sites	Locations	Enzyme	# Sites	Locations	Enzyme	# Sites	Locations
AatII	1	346	DraI	3	1055 1074 1766	SfaNI	20	
AccI	3	138 411 3026	DraIII	1	779	Sfcl	7	131 226 553 1430 2338
Acil	78		DrdI	3	823 2692 3107			2529 5416
AfIII	1	163	DsaI	2	69 5221	Sgfl	1	337
AfIII	2	2794 4658	EaeI	8	150 196 322 326 1577	SgrAI	1	5339
AhdI	1	1189	EagI	3	3984 5218 5350	SphI	1	5191
AluI	25		EarI	3	150 196 322	Sse8387I	1	135
Alw26I	7	1250 2027 3153 4043 4430	Ecl136II	1	120	Sspl	1	984
		4556 4961	Eco47III	2	3544 5257	Styl	3	69 433 473
AlwI	15		Eco57I	2	1856 2252	TaiI	15	
AlwNI	1	2385	EcoNI	1	5126	TaqI	16	
ApaI	1	4455	EcoO109I	3	478 3789 5226	TfiI	4	2820 3241 3745 3980
ApaLI	4	1856 2480 2980 4678	EcoRI	1	112	ThaI	34	
ApoI	5	112 384 959 970 4383	EcoRII	8	102 2633 2646 2767 4022	TseI	28	
AscI	1	125			4079 4619 4934	Tsp45I	7	598 1445 1656 3046 3141
AvaI	1	354	EcoRV	1	319			3354 4476
Avall	5	1327 1549 3510 3789 4107	EheI	4	4021 5203 5317 5338	Tsp509I	22	
AvrII	1	433	FauI	16		TspRI	13	
BamHI	1	106	Fnu4HI	48		Tth111I	1	3052
BanI	9	348 735 1137 3889 4019	FokI	10	1155 1336 1623 3108 3249	VspI	5	213 1361 3916 3975 5403
		4738 5201 5315 5336			3435 3513 3575 4603 4612	XbaI	1	30
BanII	5	122 705 4455 5268 5282	FseI	1	328	XcmI	3	4273 4291 4807
BbsI	3	3676 4173 4512	FspI	1	1411	XhoI	1	354
BbvI	28		HaellI	13		XmnI	2	1788 3240
BcgI	4	162 1728 3223 4338	HaeIII	24				
BclI	1	4644	Hgal	12				
Bfal	10	31 415 434 462 623	HhaI	44				
		1044 1379 2301 3782 3817	HincII	2	139 4156			
BglI	1	1309	HindIII	1	143			
BglII	1	305	HinfI	16				
BpmI	4	1259 3273 4337 4826	HpaI	1	4156			
Bpu10I	1	3689	HphI	18				
Bpu1102I	1	451	KpnI	1	352			
BsaAI	2	776 3046	MaeIII	17				
BsaBI	3	3601 5379 5389	MbolI	13				
BsaHI	7	343 1726 4020 4703 5202	MluI	1	4658			
		5316 5337	MnlI	25				
BsaI	1	1250	MseI	33				
BsaJI	7	69 433 473 2634 4023	MslI	9	1441 1600 1959 3225 3616			
		5215 5221			3811 4292 4322 4610			
BsaWI	7	528 1480 2441 2588 3605	MspA1I	10	375 450 1824 2211 2456			
		3836 4339			3088 3207 3969 4062 4632			
BsgI	3	3640 4613 4813	MspI	29				
BsiEI	9	153 199 325 337 1559	MunI	1	311			
		1708 2460 2884 3879	MwoI	34				
BsiHKAI	8	122 1775 1860 2484 2984	NarI	4	4020 5202 5316 5337			
		3808 4682 5166	NciI	12				
BsII	22		NcoI	1	69			
BsmBI	2	3153 4043	NdeI	1	298			
BsmFI	3	557 3523 5197	NgoAIV	3	324 671 5348			
Bsp1286I	12		NlaIII	26				
BspEI	2	528 3605	NlaIV	22				
BspLU11I	1	2794	NotI	1	150			
BspMI	1	124	NspI	4	2798 3165 3457 5191			
BsrBI	5	13 243 632 2031 2865	Pacl	1	429			
BsrDI	4	1250 1424 4251 4617	PfiMI	2	401 5083			
BsrFI	6	324 671 1269 4972 5339	PleI	12				
		5348	Psp1406I	5	989 1415 1788 3471 4998			
BsrGI	1	190	Psp5II	1	3789			
Bsrl	24		PstI	1	135			
BssHII	2	125 4247	PvuI	2	337 1559			
BssSI	2	1853 2621	PvuII	3	3207 3969 4062			
Bst1107I	1	3027	RcaI	4	1993 2025 2074 5260			
BstEII	1	4476	RsaI	5	192 350 1669 2992 4515			
BstXI	3	4612 4735 4864	SacI	1	122			
BstYI	11		Sall	1	137			
Cac8I	36		SapI	1	2911			
Clal	1	5383	Sau3AI	28				
CviJI	84		Sau96I	15				
Ddel	10	262 451 1146 1686 2111	ScaI	1	1669			
		2520 2987 3527 3689 4087	ScrFI	20				
Dpnl	28							

Enzymes that do not cut pETDuet-1:

BseRI	BsmI	Bsu36I	MscI	NheI	NruI
NsiI	NspV	PinAI	PmeI	PmlI	PshAI
RsrII	SacII	SanDI	SexAI	SfiI	SmaI
SnaBI	SpeI	SrfI	StuI	SunI	Swal