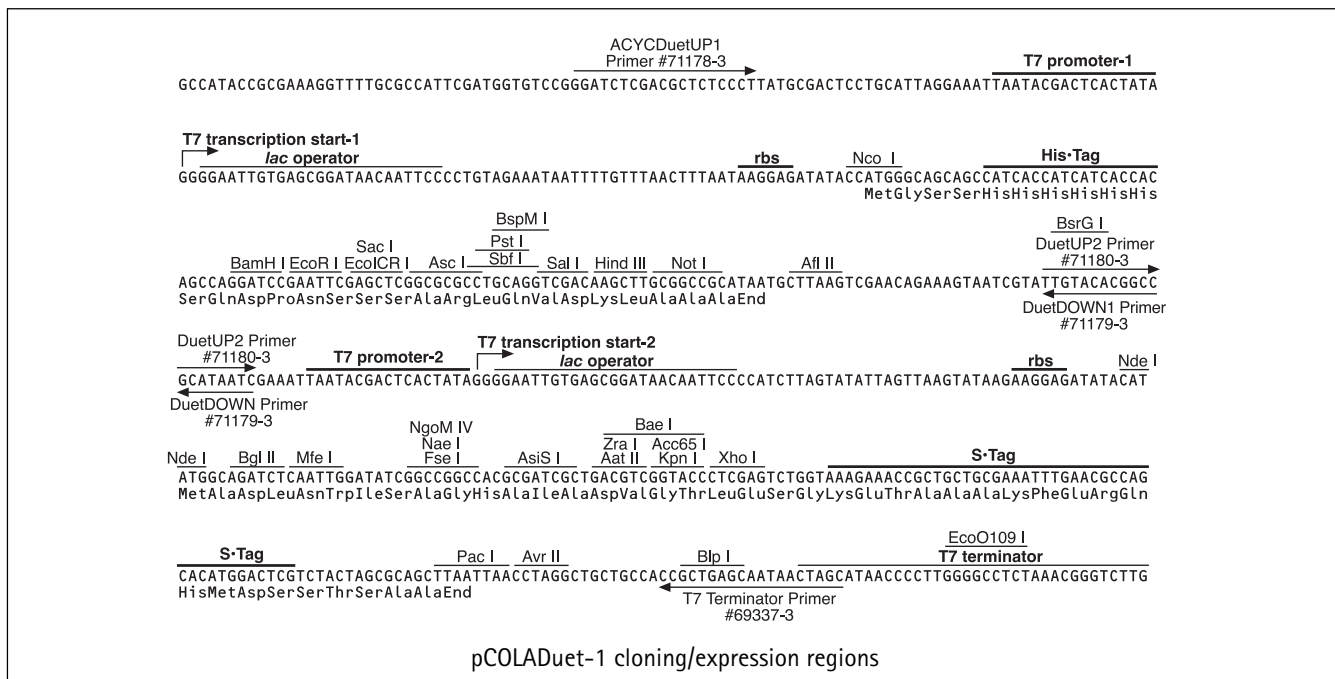
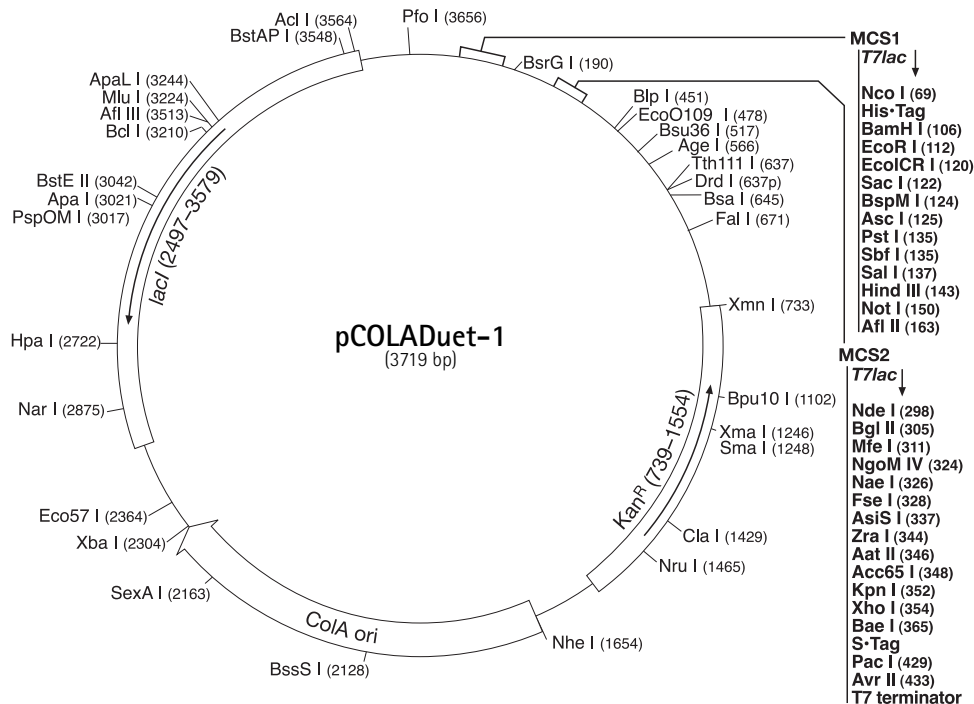


	Cat No.
pCOLADuet-1 DNA	71406-3
pCOLADuet-1 sequence landmarks	
T7 promoter-1	3703-3719
T7 transcription start-1	1
His•Tag® coding sequence	83-100
Multiple cloning sites-1 (Nco I-Afl II)	68-168
T7 promoter-2	214-230
T7 transcription start-2	231
Multiple cloning sites-2 (Nde I-Avr II)	297-438
S•Tag™ coding sequence	366-410
T7 terminator	462-509
Kan ^R	739-1554
ColA ori	1664-2299
lacI coding sequence	2497-3579

pCOLADuet™-1 is designed for the coexpression of two target genes from a single plasmid. The vector encodes two multiple cloning sites (MCS) each of which is preceded by a T7 promoter, lac operator, and ribosome binding site (rbs). MCS-1 encodes the six-amino acid His•Tag® sequence for the creation of a N-terminal fusion and MCS2 encodes the 15 amino acid S•Tag™ peptide after the last restriction site for the creation of a C-terminal fusion if desired. Genes inserted into MCS-1 can be sequenced using the ACYCDuetUP1 Primer and DuetDOWN1 Primer. Genes inserted into MCS-2 can be sequenced using the DuetUP2 Primer and T7 Terminator Primer. The vector has the COLA replicon from ColA(1) and the kanamycin resistance gene. This vector can be transformed into the same cell with plasmids containing compatible origins of replication and drug resistance genes for coexpression of up to 8 target genes.

Reference

- Zverev, V.V. and Khmel, I.A. (1985) *Plasmid* 14, 192-199.



pCOLADuet™-1 Restriction Sites

TB408 0304

Enzyme	# Sites	Locations	Enzyme	# Sites	Locations			
AatII	1	346	EarI	4	1306 1562 2332 3607			
Acc65I	1	348	Ecil	2	1730 3439			
AccI	3	138 411 1922	Ecl136II	1	120			
AccI	1	3564	Eco57I	1	2364			
AflII	1	163	Eco57MI	3	2364 2903 3392			
AflIII	1	3224	EcoCRI	1	120			
AgeI	1	566	EcoNI	3	1209 2107 3692			
ApaI	1	3021	EcoO109I	1	478			
ApaLI	1	3244	EcoRI	1	112			
AscI	1	125	EcoRV	2	319 2080			
Asel	6	213 732 921 2482 2541 3702	Eal	1	671			
AsiSI	1	337	FseI	1	328			
AvaI	2	354 1246	HaeII	3	2589 2832 3613			
AvrII	1	433	HincII	2	139 2722			
BaeI	1	365	HindIII	1	143			
BamHI	1	106	HpaI	1	2722			
BanI	4	348 2455 2585 3304	KasI	1	2585			
BanII	3	122 1471 3021	KpnI	1	352			
BbeI	1	2589	MfeI	1	311			
BbsI	2	2739 3078	MluI	1	3224			
BceAI	5	211 801 1708 2740 3367	MslI	3	2858 2888 3176			
BcgI	2	162 2904	NaeI	1	326			
BciVI	4	728 1604 1681 2772	NarI	1	2586			
BclI	1	3210	NcoI	1	69			
BfrBI	2	1008 1274	NdeI	1	298			
BglII	1	305	NgoMIV	1	324			
BipI	1	451	NheI	1	1654			
Bme1580I	2	3021 3248	NotI	1	150			
BmgBI	1	1908	NruI	1	1465			
BmrI	3	2426 3066 3303	NsiI	2	1010 1276			
BmtI	1	1658	NspI	1	1654			
BpmI	2	2903 3392	Pacl	1	429			
Bpu10I	1	1102	PfiMI	3	401 862 3649			
BpuEI	4	515 2024 2252 2416	PfoI	1	3656			
BsaHI	4	343 1870 2586 3269	PinAI	1	566			
Bsal	1	645	PspOMI	1	3017			
BsaWI	6	551 566 983 1988 2402 2905	PstI	1	135			
BsaXI	2	655 2556	PvuI	2	337 2255			
BseYI	3	1939 2690 2825	PvuII	2	2535 2628			
BsgI	2	3179 3379	SacI	1	122			
BsiEI	8	153 199 325 337 636 1124 2255 2445	Sall	1	137			
BsiHKAI	2	122 3248	SbfI	1	135			
BsmAI	8	645 1102 1604 1695 2609 2996 3122 3527	SexAI	1	2163			
BsmBI	2	1102 2609	SfcI	4	29 131 226 3715			
BsmFI	1	1894	SfoI	1	2587			
BsmI	2	1163 1240	SmaI	1	1248			
Bsp1286I	4	122 1471 3021 3248	SmlI	6	163 354 494 2003 2267 2431			
BspCNI	5	443 530 1094 2051 2645	SphI	1	1654			
BspHI	2	725 1602	Sse8387I	1	135			
BspMI	1	124	SspI	2	1197 1571			
BsrBI	4	13 243 723 1608	StyI	3	69 433 473			
BsrDI	2	2817 3183	TaqII	2	870 2431			
BsrFI	4	324 566 1164 3538	TatI	1	190			
BsrGI	1	190	TspGWI	5	1303 1315 1863 1906 2298			
BssHII	2	125 2813	Tth111I	1	637			
BstAPI	1	3548	XbaI	1	2304			
BstEII	1	3042	XcmI	3	2839 2857 3373			
BstXI	3	3178 3301 3430	XhoI	1	354			
BstYI	7	106 305 869 2263 2274 2448 3660	XmaI	1	1246			
Bsu36I	1	517	XmnI	1	733			
BtgI	1	69	ZraI	1	344			
BtrI	1	1908	Enzymes that do not cut pCOLADuet-1:					
BtsI	5	543 1176 1263 2497 2865	AarI	AfeI	AhdI	AleI	Alol	AlwNI
Clal	1	1429	BbvCI	BglI	BpII	BsaAI	BsaBI	BseRI
DrdI	1	637	BsiWI	BspEI	BspLU11I	BssSI	Bst1107I	BstBI
EaeI	5	150 196 322 326 2550	BstZ17I	DraI	DraIII	FspAI	FspI	MscI
EagI	3	150 196 322	NspV	PciI	PmeI	PmlI	Ppil	PpuMI
			PshAI	Psil	Psrl	RsrII	SacII	SanDI
			SapI	Scal	Sfil	SgrAI	SnaBI	SpeI
			SrfI	StuI	Swal			