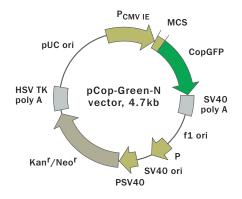


Mammalian expression vector pCop-Green-N



For vector sequence, please visit our Web site at www.evrogen.com/support/vector-info.shtml

Product	Cat.#	Size				
pCop-Green-N	FP502	20 μg				

Please contact your local distributor for exact prices and delivery information.

Vector type mammalian expression vector

Reporter CopGFP
Reporter codon usage mammalian
Promoter for CopGFP PCMV IE
Host cells mammalian

Selection prokaryotic — kanamycin

eukaryotic — neomycin (G418)

Replication prokaryotic — pUC ori eukaryotic — SV40 ori

Multiple cloning site (MCS)

Nhel	BglII	Sacl	HindIII	EcoRI	S	Sall	Kpnl	Apal	BamHI	Agel		CopGFP
GCTA.GCG.CTA.CCG.GAC.	TCA.GAT.CTC	.GAG.CTC.AA	AG.CTT.CC	GA.ATT.CTO	G.CAG.T	CG.ACG	.GTA.CCG.	CGG.GCC.	CGG.GAT.CC	A.CCG.GT	C.GCC.ACC	.ATG.CCC.GCC
Fco47III	Xh	ol .			Pstl*		S	acll Sma	I/Xmal			

^{* -} not unique sites.

Use

- Generation of fusions to the CopGFP N-terminus
- Expression of CopGFP or its fusions in mammalian cells

Vector description

pCop-Green-N vector is an eukaryotic (mammalian) expression vector encoding green fluorescent protein CopGFP. The vector allows to generate fusions to the CopGFP N-terminus and to express CopGFP fusions or CopGFP alone in mammalian cells.

CopGFP codon usage is optimized for high expression in mammalian cells (humanized, Haas et~al., 1996). To increase CopGFP translation, Kozak consensus translation initiation site is generated upstream of CopGFP sequence (Kozak, 1987). Multiple cloning site (MCS) is located between $P_{\text{CMV},\text{IF}}$ and CopGFP coding sequence.

The vector backbone comprises immediate early promoter of cytomegalovirus (P_{CMV IE}) for protein expression, SV40 origin for replication in mammalian cells expressing SV40 T-antigen, pUC origin of replication for propagation in *E. coli* and f1 origin for single-stranded DNA production. SV40 polyadenylation signals (SV40 poly A) direct proper processing of the 3' end of the reporter mRNA.

SV40 early promoter provides neomycin resistance gene expression to select stably transfected eukaryotic cells using G418. Bacterial promoter (P) provides kanamycin resistance gene expression in *E. coli*. Kan^r/Neo^r gene is linked with herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signals.

Generation of fusions

A localization signal or a gene of interest should be cloned into MCS of the vector. It will be expressed as a fusion to the CopGFP N-terminus when inserted in the same reading frame as CopGFP and no in-frame stop codons are present. The inserted sequence should contain an initiating ATG codon. CopGFP-tagged fusions retain fluorescent properties of the native protein allowing fusion localization *in vivo*.

Notes: The plasmid DNA was isolated from dam⁺-methylated *E.coli*. Therefore some restriction sites are bloked by methylation. If you wish to digest the vector using such sites you will need to transform the vector into a dam⁻ host and make fresh DNA.

Expression in mammalian cells

pCop-Green-N vector can be transfected into mammalian cells by any known transfection method. If required, stable transformants can be selected using G418 (Gorman, 1985). Unmodified pCop-Green-N will express CopGFP, when transfected into eukaryotic (mammalian) cells.

Propagation in E. coli

Suitable host strains for propagation in *E. coli* include DH5alpha, HB101, XL1-Blue, and other general purpose strains. Plasmid incompatibility group is pMB1/ColE1. The vector confers resistance to kanamycin (30 μ g/ml) to *E. coli* hosts. Copy number in *E. coli* is about 500.

Location of features

PCMV IE: 1-589

Enhancer region: 59-465 TATA box: 554-560

Transcription start point: 583

MCS: 591-671 CopGFP

Kozak consensus translation initiation site: 672-682 Start codon (ATG): 679-681; Stop codon: 1496-1498

SV40 early mRNA polyadenylation signal

Polyadenylation signals: 1551-1556 & 1580-1585

mRNA 3' ends: 1589 & 1601

f1 single-strand DNA origin: 1648-2103 (Packages the noncoding strand of CopGFP) Bacterial promoter for expression of Kan^r gene

-35 region: 2165-2170; -10 region: 2188-2193 Transcription start point: 2200

SV40 origin of replication: 2444-2579 SV40 early promoter

Enhancer (72-bp tandem repeats): 2277-2348 & 2349-2420 21-bp repeats: 2424-2444, 2445-2465, & 2467-2487

Early promoter element: 2500-2506

Major transcription start points: 2496, 2534, 2540 & 2545

Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences: Start codon (ATG): 2628-2630; stop codon: 3420-3422

G->A mutation to remove PstI site: 2810

C->A (Arg to Ser) mutation to remove BssHII site: 3156

Herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signal

Polyadenylation signals: 3658-3663 & 3671-3676 pUC plasmid replication origin: 4007-4650

Kozak M. (1987) Nucleic Acids Res. 15:8125-8148.

Haas J. et al. (1996) Curr. Biol. 6: 315-324.

Gorman C. (1985) In DNA cloning: A Practical

Approach, Vol. II. Ed. D. M. Glover. (IRL Press, Oxford,

Notice to Purchaser:

References

U.K.), pp. 143-190.

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