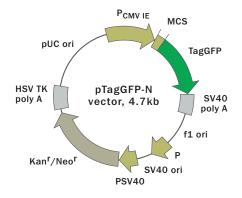


# Mammalian expression vector pTagGFP-N



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

Product	Cat.#	Size
pTagGFP-N	FP122	20 μg

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

Vector type mammalian expression vector

Reporter TagGFP
Reporter codon usage mammalian
Promoter for TagGFP P<sub>CMV IE</sub>
Host cells mammalian

Selection prokaryotic — kanamycin

eukaryotic — neomycin (G418)

Replication prokaryotic — pUC ori eukaryotic — SV40 ori

## Multiple cloning site (MCS)

															TagGI	FP	
Nhel	Bglll	Sacl	HindIII	EcoRI		Sall	Крі	nl .		Apal	BamHI	Agel				$\longrightarrow$	-
G.CTA.GCG.CTA.CCG.GAC.T	CA.GAT.CTC.	GAG.CTC.	AAG.CTT.	CGA.ATT.	CTG.CAG	a.TCG.	ACG. GT	A.CCG.	CGG.	GCC. CGG	G.GAT.CC	A. CCG.	TC.GCC	. ACC.	ATG. A	GC.GGG	1
Afel	Xh	ol			Pstl	_		S	acll*	Smal/X	mal						

<sup>\* -</sup> not unique sites.

#### Use

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

# **Vector description**

pTagGFP-N is an eukaryotic (mammalian) expression vector encoding green fluorescent protein TagGFP. The vector is designed to generate TagGFP-tagged fusions or to express TagGFP in mammalian cells.

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

The vector backbone comprises immediate early promoter of cytomegalovirus ( $P_{CMV \mid E}$ ) 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 polyA) 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<sup>r</sup>/Neo<sup>r</sup> 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 TagGFP N-terminus when inserted in the same reading frame as TagGFP and no intervening stop codons are present. The inserted sequence should contain an initiating ATG codon.

TagGFP fusions retain fluorescent properties of the native protein allowing fusion protein localization *in vivo*.

**Note:** The plasmid DNA was isolated from dam<sup>+</sup>-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<sup>-</sup> host and make fresh DNA.

#### Expression in mammalian cells

pTagGFP-N can be transfected into mammalian cells by any known transfection method. If required, stable transformants can be selected using G418 (Gorman, 1985).

Unmodified pTagGFP-N will express TagGFP, 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/CoIE1. The vector confers resistance to kanamycin (30 µg/ml) to E. coli hosts. Copy number in E. coli is about 500.

#### **Location of features**

P<sub>CMV IE</sub>: 1-589

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

Transcription start point: 583

MCS: 592-678 **TagGFP** 

Kozak consensus translation initiation site: 672-682 Start codon (ATG): 679-681; stop codon: 1393-1395

SV40 early mRNA polyadenylation signal

Polyadenylation signals: 1549-1554 & 1578-1583

mRNA 3' ends: 1587 & 1599

f1 single-strand DNA origin: 1646-2101 Bacterial promoter for expression of Kan<sup>r</sup> gene

-35 region: 2163-2168; -10 region: 2186-2191

Transcription start point: 2198 SV40 origin of replication: 2442-2577

SV40 early promoter

Enhancer (72-bp tandem repeats): 2275-2346 & 2347-2418 21-bp repeats: 2422-2442, 2443-2463 & 2465-2485

Early promoter element: 2498-2504

Major transcription start points: 2494, 2532, 2538 & 2543

Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences: Start codon (ATG): 2626-2628; stop codon: 3418-3420

G->A mutation to remove Pst I site: 2808

C->A (Arg to Ser) mutation to remove BssH II site: 3154

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

Polyadenylation signals: 3656-3661 & 3669-3674 pUC plasmid replication origin: 4005-4648

# Notice to Purchaser:

References

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

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The CMV promoter is covered under U.S. Patents 5,168,062 and 5,385,839 and its use is permitted for research purposes only. Any other use of the CMV promoter requires a license from the University of Iowa Research Foundation, 214 Technology Innovation Center, Iowa City, IA 52242.

#### MATERIAL SAFETY DATA SHEET INFORMATION

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

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

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

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

To the best of our knowledge, these products do not require a Material Safety Data Sheet. However, all the properties of these products (and, if applicable, each of their components) have not been thoroughly investigated. Therefore, we recommend that you use gloves and eye protection, and wear a laboratory coat when working with these products.

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