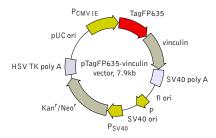


pTagFP635-vinculin vector

The vector sequence has been compiled using the informa-tion from sequence databases, published literature, and other sources, together with partial sequences obtained by Evrogen. This vector has not been completely sequenced.



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

Product	Cat.#	Size	
pTagFP635-vinculin vector	FP388	20 μ g	
The price does not include delivery. The price varies in different countries. Please contact your local distributor for exact prices and delivery information.			

TagFP635 Reporter mammalian Reporter codon usage Promoter for TagFP635 P_{CMV} IE Host cells mammalian

Selection prokaryotic - kanamycin

eukaryotic - neomycin (G418)

mammalian expression vector

Replication prokaryotic - pUC ori

eukaryotic - SV40 ori

Use far-red fluorescent labeling of vinculin

Location of features

P_{CMV IE}: 1-589 Enhancer region: 59-465 TATA box: 554-560 Transcription start point: 583

Kozak consensus translation initiation site: 606-616

TagFP635

Start codon (ATG): 613-615

Last amino acid in TagFP635: 1321-1323

Vinculin: 1393-4593 Stop codon: 4591-4593

SV40 early mRNA polyadenylation signal

Polyadenylation signals: 4746-4751 & 4775-4780 mRNA 3' ends: 4784 & 4796

f1 single-strand DNA origin: 4843-5298 Bacterial promoter for expression of Kan^r gene -35 region: 5360-5365; -10 region: 5383-5388

Transcription start point: 5395 SV40 origin of replication: 5639-5774

SV40 early promoter

Enhancer (72-bp tandem repeats): 5472-5543 & 5544-5615

21-bp repeats: 5619-5639, 5640-5660 & 5662-5682

Early promoter element: 5695-5701 Major transcription start points: 5691, 5729, 5735 &

5740

Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences: Start codon (ATG): 5823-5825: Stop codon: 6615-6617

G->A mutation to remove Pst I site: 6005

C->A (Arg to Ser) mutation to remove BssH II site: 6351 Herpes simplex virus (HSV) thymidine kinase (TK)

polyadenylation signal

Polyadenylation signals: 6853-6858 & 6866-6871 pUC plasmid replication origin: 7202-7845

Vector description

Vector type

pTagFP635-vinculin is a mammalian expression vector encoding TagFP635-vinculin fusion protein. The vector can be used for fluorescent labeling of vinculin in living cells.

TagFP635 codon usage is optimized for high expression in mammalian cells, i.e. humanized [Haas et al. 1996]. Human vinculin is fused to the TagFP635 C-terminus. To increase mRNA translation efficiency. Kozak consensus translation initiation site is generated upstream of TagFP635-vinculin coding sequence [Kozak

pTagFP635-vinculin can be used as a source of TagFP635-vinculin hybrid sequence. The vector backbone contains unique restriction sites that permit its excision and further insertion into expression vector of choice.

Note: The plasmid DNA was isolated from dam*-methylated E.coli. Therefore some restriction sites are blocked 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.

The vector backbone also contains immediate early promoter of cytomegalovirus ($P_{\text{CMV IE}}$) for protein expressions. sion, 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 (PSV40) provides neomycin resistance gene (Neor) expression to select stably transfected eukaryotic cells using G418. Bacterial promoter (P) provides kanamycin resistance gene expression (Kan^r) in E. coli. Kan^r/Neo^r gene is linked with herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signals.

Expression in mammalian cells

pTagFP635-vinculin can be transfected into mammalian cells by any known transfection method. CMV promoter provides strong, constitutive expression of the TagFP635-vinculin fusion in eukaryotic cells. If required, stable transformants can be selected using G418 [Gorman 1985].

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 μ g/ml) to E. coli hosts. Copy number in E. coli is about 500.

References

Gorman (1985). "High efficiency gene transfer into mammalian cells." In: DNA cloning: A Practical Approach, Vol. II. Ed. by Glover. (IRL Press, Oxford, U.K.) Pp. 143-190.

Haas et al. (1996) "Codon usage limitation in the expression of HIV-1 envelope glycoprotein." Curr Biol, 6 (3): 315-324 / pmid:

Kozak (1987) "An analysis of 5'-noncoding sequences from 699 vertebrate messenger RNAs." Nucleic Acids Res, 15 (20): 8125-8148 / pmid: 3313277

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