

# pTagFP635-zyxin vector

### Cat# FP389

## Vector description

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

TagFP635 codon usage is optimized for high expression in mammalian cells, i.e. humanized (Haas et al., 1996). Human zyxin is fused to the TagFP635 N-terminus.

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

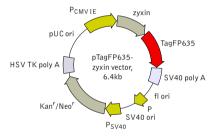
**Note:** The plasmid DNA was isolated from dam<sup>+</sup>-methylated *E. coli*. Therefore some restriction sites are blocked by methylation. If you wish to digest the vector using such sites you will need to transform the vector into a dam<sup>-</sup> host and make fresh DNA

The vector backbone also contains immediate early promoter of cytomegalovirus ( $P_{\text{CMV IE}}$ ) 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 ( $P_{SV40}$ ) provides neomycin resistance gene ( $Neo^r$ ) 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.

## **Vector map**

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



# Expression in mammalian cells

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

### Location of features

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

Enhancer region: 59-465

TATA box: 554-560

Transcription start point: 583

Zyxin: 636-2348

TagFP635: 2370-3083

SV40 early mRNA polyadenylation signal

Polyadenylation signals: 3236-3241 3265-3270

mRNA 3' ends: 3274 3286

f1 single-strand DNA origin: 3333-3788

Bacterial promoter for expression of Kan<sup>r</sup> gene

-35 region: 3850-3855

-10 region: 3873-3878

Transcription start point: 3885 SV40 origin of replication: 4129-4264

SV40 early promoter

Enhancer (72-bp tandem repeats): 3962-4033 4034-4105

21-bp repeats: 4109-4129, 4130-4150 4152-4172

Early promoter element: 4185-4191

Major transcription start points: 4181, 4219, 4225, 4230

Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences:

Start codon (ATG): 4313-4315

Stop codon: 5105-5107

G->A mutation to remove Pst I site: 4495

C->A (Arg to Ser) mutation to remove BssH II site: 4841

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

Polyadenylation signals: 5343-5348 5356-5361

pUC plasmid replication origin: 5692-6335

### 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  $\mu$ g/ml) to *E. coli* hosts. Copy number in *F. coli* is about 500

### References:

Gorman C. High efficiency gene transfer into mammalian cells. In DNA cloning: A Practical Approach, Vol. II. Ed. D. M. Glover. (IRL Press, Oxford, U.K.). 1985; 143-90.

Haas J, Park EC, Seed B. Codon usage limitation in the expression of HIV-1 envelope glycoprotein. Curr Biol. 1996; 6 (3):315-24. / pmid: 8805248

Kozak M. An analysis of 5'-noncoding sequences from 699 vertebrate messenger RNAs. Nucleic Acids Res. 1987; 15 (20):8125-48. / pmid: 3313277

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