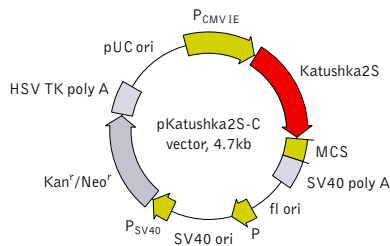


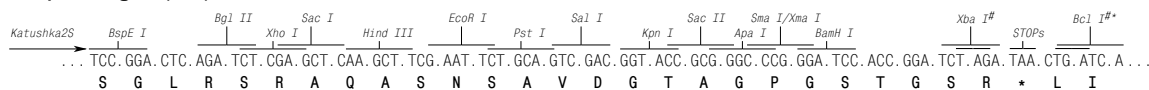
## pKatushka2S-C vector

The vector sequence has been compiled using the information 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/products/vectors.shtml>

### Multiple cloning site (MCS)



\* — not unique sites.

# — sites are blocked by *dam* methylation. If you wish to digest the vector with these enzymes, you will need to transform the vector into a *dam<sup>-</sup>* host and make fresh DNA.

### Location of features

P<sub>CMV IE</sub>: 1-589  
 Enhancer region: 59-465  
 TATA box: 554-560  
 Transcription start point: 583  
 Kozak consensus translation initiation site: 606-616  
 Katushka2S  
 Start codon (ATG): 613-615; Stop codon: 1396-1398  
 Last amino acid in Katushka2S: 1315-1317  
 MCS: 1318-1398  
 SV40 early mRNA polyadenylation signal  
 Polyadenylation signals: 1538-1543 & 1567-1572  
 mRNA 3' ends: 1576 & 1588  
 f1 single-strand DNA origin: 1635-2090  
 Bacterial promoter for expression of Kan<sup>r</sup> gene  
 -35 region: 2152-2157; -10 region: 2175-2180  
 Transcription start point: 2187  
 SV40 origin of replication: 2431-2566  
 SV40 early promoter  
 Enhancer (72-bp tandem repeats): 2264-2335 & 2336-2407  
 21-bp repeats: 2411-2431, 2432-2452 & 2454-2474  
 Early promoter element: 2487-2493  
 Major transcription start points: 2483, 2521, 2527 & 2532  
 Kanamycin/neomycin resistance gene  
 Neomycin phosphotransferase coding sequences:  
 Start codon (ATG): 2615-2617; Stop codon: 3407-3409  
 G->A mutation to remove Pst I site: 2797  
 C->A (Arg to Ser) mutation to remove BssH II site: 3143  
 Herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signal  
 Polyadenylation signals: 3645-3650 & 3658-3663  
 pUC plasmid replication origin: 3994-4637

### References

Gorman, C. (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, J. et al. (1996) "Codon usage limitation in the expression of HIV-1 envelope glycoprotein." *Curr Biol*, 6 (3): 315-324 / pmid: 8805248

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

### Notice to Purchaser:

Katushka2S-related materials (also referred to as "Products") are intended for research use only. The Products are covered by U.S. Pat. 8,138,320; European Pat. 1,994,149; and other Evrogen Patents and/or Patent applications pending. By use of these Products, you accept the terms and conditions of the applicable Limited Use Label License #001: <http://www.evrogen.com/products/Evrogen-FP-license.shtml>. 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.

**MSDS information** is available at <http://www.evrogen.com/MSDS.shtml>

Product	Cat.#	Size
pKatushka2S-C vector	<b>FP761</b>	20 µg
Vector type	mammalian expression vector	
Reporter	Katushka2S	
Reporter codon usage	mammalian	
Promoter for Katushka2S	P <sub>CMV IE</sub>	
Host cells	mammalian	
Selection	prokaryotic - kanamycin eukaryotic - neomycin (G418)	
Replication	prokaryotic - pUC ori eukaryotic - SV40 ori	
Use	Katushka2S expression in mammalian cells; generation of fusions to the Katushka2S C-terminus	

### Vector description

pKatushka2S-C is a mammalian expression vector encoding far-red fluorescent protein Katushka2S. The vector allows generation of fusions to the Katushka2S C-terminus and expression of Katushka2S fusions or Katushka2S alone in eukaryotic (mammalian) cells.

Katushka2S codon usage is optimized for high expression in mammalian cells (humanized) [Haas et al. 1996]. To increase mRNA translation efficiency, Kozak consensus translation initiation site is generated upstream of the Katushka2S coding sequence [Kozak 1987]. Multiple cloning site (MCS) is located between Katushka2S coding sequence and SV40 polyadenylation signal (SV40 poly A).

The vector backbone contains immediate early promoter of cytomegalovirus (P<sub>CMV IE</sub>) 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<sub>SV40</sub>) provides neomycin resistance gene (Neo<sup>r</sup>) expression to select stably transfected eukaryotic cells using G418. Bacterial promoter (P) provides kanamycin resistance gene expression (Kan<sup>r</sup>) 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 Katushka2S fusion proteins

A localization signal or a gene of interest can be cloned into MCS of the vector. It will be expressed as a fusion to the Katushka2S C-terminus when inserted in the same reading frame as Katushka2S and no in-frame stop codons are present. Katushka2S-tagged fusions retain fluorescent properties of the native protein allowing fusion localization *in vivo*. Unmodified vector will express Katushka2S when transfected into eukaryotic (mammalian) cells.

**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.

### Expression in mammalian cells

pKatushka2S-C vector can be transfected into mammalian cells by any known transfection method. CMV promoter provides strong, constitutive expression of Katushka2S or its fusions 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/ColE1. The vector confers resistance to kanamycin (30 µg/ml) to *E. coli* hosts. Copy number in *E. coli* is about 500.