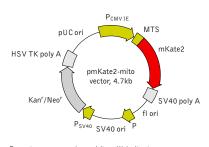


pmKate2-mito vector

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

Location of features

P_{CMV IE}: 1-589 Enhancer region: 59-465 TATA box: 554-560 Transcription start point: 583 mKate2-mito fusion Start codon (ATG): 597-599 Mitochondrial targeting sequence (MTS): 597-683 Start of mKate2 coding sequence (ATG): 705-707 Stop codon: 1401-1403 SV40 early mRNA polyadenylation signal Polyadenylation signals: 1556-1561 & 1585-1590 mRNA 3' ends: 1594 & 1606 f1 single-strand DNA origin: 1653-2108 Bacterial promoter for expression of Kan^r gene -35 region: 2170-2175; -10 region: 2193-2198 Transcription start point: 2205 SV40 origin of replication: 2449-2584 SV40 early promoter Enhancer (72-bp tandem repeats): 2282-2353 & 2354 2425 21-bp repeats: 2429-2449, 2450-2470 & 2472-2492 Early promoter element: 2505-2511 Major transcription start points: 2501, 2539, 2545 & 2550

Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences: Start codon (ATG): 2633-2635; Stop codon: 3425-3427 G->A mutation to remove Pst I site: 2815

Polyadenylation signals: 3663-3668 & 3676-3681 pUC plasmid replication origin: 4012-4655

Product	Cat.#	Size	
pmKate2-mito vector	FP187	20 μ g	
Vector type	mammalian expression vector		
Reporter	mKate2		
Reporter codon usage	mammalian		
Promoter for mKate2	P _{CMV IE}		
Host cells	mammalian		
Selection	prokaryotic - kanamycin		
	eukaryotic - neomycin (G418)		
Replication	prokaryotic - pUC ori		
	eukaryotic - SV4	0 ori	
Use	far-red fluorescent labeling of mitochondria		

Vector description

pmKate2-mito is a mammalian expression vector intended for far-red fluorescent labeling of mitochondria in living cells. The vector encodes far-red fluorescent protein mKate2 fused to mitochondrial targeting sequence (MTS) derived from the subunit VIII of human cytochrome C oxidase [Rizzuto et al. 1989; Rizzuto et al. 1995]. MTS is fused to the mKate2 N-terminus.

mKate2 codon usage is optimized for high expression in mammalian cells (humanized) [Haas et al. 1996].

pmKate2-mito vector can be used as a source of mKate2-MTS 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 contains 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 (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.

Expression in mammalian cells

pmKate2-mito vector can be transfected into mammalian cells by any known transfection method. CMV promoter provides strong, constitutive expression of the mKate2-MTS 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/ColE1. The vector confers resistance to kanamycin (30 μ g/ml) to *E. coli* hosts. Copy number in *E. coli* is about 500.

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

Rizzuto, R. et al. (1989) "A gene specifying subunit VIII of human cytochrome c oxidase is localized to chromosome 11 and is expressed in both muscle and non-muscle tissues." J Biol Chem, 264 (18): 10595–10600 / pmid: 2543673

Rizzuto, R. et al. (1995) "Chimeric green fluorescent protein as a tool for visualizing subcellular organelles in living cells." Curr Biol, 5 (6): 635–642 / pmid: 7552174

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