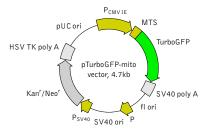


pTurboGFP-mito vector

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

Product	Cat.#	Size	
pTurboGFP-mito vector	FP517	20 μ g	
Vector type	mammalian expression vector		
Reporter	TurboGFP		
Reporter codon usage	mammalian		
Promoter for TurboGFP	P _{CMV IE}		
Host cells	mammalian		
Selection	prokaryotic - kanamycin eukaryotic - neomycin (G418)		
Replication	prokaryotic - pUC ori		
	eukaryotic - SV40) ori	
Use	green fluorescent labeling of mitochondria		

Location of features

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

TurboGFP-mito fusion Start codon (ATG): 597-599

Mitochondrial targeting sequence (MTS): 597-683 Start of TurboGFP coding sequence (ATG): 705-707

Stop codon: 1401-1403

SV40 early mRNA polyadenylation signal Polyadenylation signals: 1557-1562 & 1586-1591

Polyadenylation signals: 1557-1562 & 1586-1592 mRNA 3' ends: 1595 & 1607

f1 single-strand DNA origin: 1654-2109 Eukaryotic promoter for expression of Kan^r gene -35 region: 2171-2176; -10 region: 2194-2199

Transcription start point: 2206 SV40 origin of replication: 2450-2585

SV40 origin of replication: 2

Enhancer (72-bp tandem repeats): 2283-2354 & 2355

21-bp repeats: 2430-2450, 2451-2471 & 2473-2493 Early promoter element: 2506-2512

Major transcription start points: 2502, 2540, 2546 & 2551

Kanamycin/neomycin resistance gene Neomycin phosphotransferase coding sequences:

Start codon (ATG): 2634-2636; Stop codon: 3426-3428 G->A mutation to remove Pst I site: 2816

C->A (Arg to Ser) mutation to remove BssH II site: 3162 Herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signal

Polyadenylation signals: 3664-3669 & 3677-3682 pUC plasmid replication origin: 4013-4656

Vector description

pTurboGFP-mito is a mammalian expression vector intended for green fluorescent labeling of mitochondria in living cells. The vector encodes green fluorescent protein TurboGFP 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 TurboGFP N-terminus.

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

pTurboGFP-mito vector can be used as a source of TurboGFP-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 ($P_{\text{CMV IE}}$) 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.

Expression in mammalian cells

pTurboGFP-mito vector can be transfected into mammalian cells by any known transfection method. CMV promoter provides strong, constitutive expression of the TurboGFP-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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