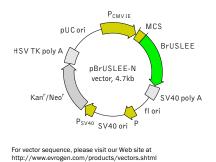


# pBrUSLEE-N vector

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.



Product	Cat.#	Size				
pBrUSLEE-N vector	FP214	20 $\mu$ g				
Vector type	mammalian expre	mammalian expression vector				
Reporter	BrUSLEE					
Reporter codon usage	mammalian					
Promoter for BrUSLEE	P <sub>CMV IE</sub>					
Host cells	mammalian					
Selection	prokaryotic - kanamycin					
	eukaryotic - neom	iycin (G418)				
Replication	prokaryotic - pUC	ori				
	eukaryotic - SV40	ori				
Use	BrUSLEE expression in mammalian cells; generation of					
	fusions to the BrL	fusions to the BrUSLEE N-terminus				

### Multiple cloning site (MCS)

Afe I	Xho I	Hind III	Pst I*	Kpn I Apa I	BamH I	Nco I*
Nhe I	Bgl II Sac I	E	EcoR I Sal I	Sac II*   Sma I/	Xma I Age I	BrUSLEE
						<u> </u>
G.CTA.GCG.CTA.CCG.GAC.T	CA.GAT.CTC.GAG.CTC	. AAG.CTT.CGA.	ATT.CTG.CAG.TCG.A	CG.GTA.CCG.CGG.GCC.CG	G.GAT.CCA.CCG.GTC.(	GCC.ACC. <i>ATG.G</i>
LALPDS	S D L E L	KLR	ILQS	TVPRAR	DPPV	A T M
* – not unique sites.						

#### - not unique sites.

# Location of features

P<sub>CMV IE</sub>: 1-589 Enhancer region: 59-465 TATA box: 554-560 Transcription start point: 583 MCS: 592-678 BrUSLEE Kozak consensus translation initiation site: 672-682 Start codon (ATG): 679-681; Stop codon: 1396-1398 SV40 early mRNA polyadenylation signal Polyadenylation signals: 1551-1556 & 1580-1585 mRNA 3<sup>°</sup> ends: 1588 & 1601

f1 single-strand DNA origin: 1648-2103

Eukaryotic promoter for expression of Kan<sup>r</sup> gene -35 region: 2165-2170; -10 region: 2188-2193

Transcription start point: 2200 SV40 origin of replication: 2444-2579

SV40 early promoter

Enhancer (72-bp tandem repeats): 2277-2348 & 2349-2420

21-bp repeats: 2424-2444, 2445-2465 & 2467-2487 Early promoter element: 2500-2506

Major transcription start points: 2496, 2534, 2540 & 2545

Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences: Start codon (ATG): 2628-2630; stop codon: 3420-3422 G->A mutation to remove Pst I site: 2810 C->A (Arg to Ser) mutation to remove BssH II site: 3156

Herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signal Polyadenylation signals: 3658-3663 & 3671-3676

pUC plasmid replication origin: 4007-4650

#### 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

# Vector description

pBrUSLEE-N is a mammalian expression vector encoding green fluorescent protein BrUSLEE. The vector allows generation of fusions to the BrUSLEE N-terminus and expression of BrUSLEE fusions or BrUSLEE alone in eukaryotic (mammalian) cells.

BrUSLEE 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 BrUSLEE coding sequence [Kozak 1987]. Multiple cloning site (MCS) is located between  $P_{CMV IE}$  and BrUSLEE coding sequence.

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<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 BrUSLEE 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 BrUSLEE N-terminus when inserted in the same reading frame as BrUSLEE and no in-frame stop codons are present. The inserted sequence should contain an initiating ATG codon. BrUSLEE-tagged fusions retain fluorescent properties of the native protein allowing fusion localization *in vivo*. Unmodified vector will express BrUSLEE 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

pBrUSLEE-N vector can be transfected into mammalian cells by any known transfection method. CMV promoter provides strong, constitutive expression of BrUSLEE 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  $\mu$ g/ml) to *E. coli* hosts. Copy number in *E. coli* is about 500.

### Notice to Purchaser:

BrUSLEE-related materials (also referred to as "Products") are intended for research use only. MSDS information is available at http://www.evrogen.com/MSDS.shtml