



			NheI			NotI			HindIII			SalI
	BamHI			MluI			EagI	ClaI			EcoRV	
601	GGGATCCTCT	AGTCAGCTGA	CGCGTGCTAG	CGCGGCCGCA	TCGATAAGCT	TGTCGACGAT						
	CCCTAGGAGA	TCAGTCGACT	GCGCAGCATC	GCGCCGGCGT	AGCTATTCTGA	ACAGCTGCTA						
	EcoRV											
661	ATCTCCAGAG											
	TAGAGGTCTC											

pBI-CMV5 Vector Map and Multiple Cloning Site.

Description

pBI-CMV5 is a mammalian bidirectional expression vector designed to constitutively express a protein of interest and *Metridia* luciferase (MetLuc), a secreted reporter protein that can be easily detected in the medium surrounding the cells.

Metridia luciferase is a naturally secreted luciferase from the marine copepod *Metridia longa*. The human codon-optimized *Metridia* luciferase gene encodes a 219 amino acid (24 kDa) polypeptide that includes a 17 amino acid N-terminal signal peptide necessary for secretion (1). *Metridia* luciferase is expressed and secreted into the surrounding medium, where it is easily detected by the addition of a chemiluminescent substrate.

Protein expression is driven by one of two constitutively active, minimal human cytomegalovirus promoters: $P_{\min\text{CMV1}}$ (located upstream of the multiple cloning site [MCS]), drives the expression of the protein of interest, and $P_{\min\text{CMV2}}$ drives the expression of MetLuc. To allow propagation and selection in *E. coli*, the vector contains a CoIE1 origin of replication and an ampicillin resistance gene (Amp^r).

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Use

The pBI-CMV5 vector, available as part of the Bidirectional Secreted Luciferase System (Cat. No. 631757), is designed to constitutively express a protein of interest and *Metridia* luciferase. The gene of interest must contain an initiation codon and a stop codon.

pBI-CMV5 can be transfected into mammalian cells using any standard transfection method. Cells expressing the secreted *Metridia* luciferase can be detected by adding chemiluminescent substrate (provided) to a sample of the cell medium.

Location of features

- Enhancer: 64–473
- P_{minCMV1} (minimal human cytomegalovirus promoter 1): 474–599
- MCS (multiple cloning site): 602–663
- SV40 polyA signals: 675–862
- ColE1 origin of replication: 1038–1637
- Amp^r (ampicillin resistance gene): 1799–2659 (complementary)
- SV40 polyA signals: 2795–2982 (complementary)
- MetLuc (*Metridia* luciferase): 3017–3676
- P_{minCMV2} (minimal human cytomegalovirus promoter 2): 3691–3759

Propagation in *E. coli*

- Recommended host strain: DH5 α TM and other general purpose strains.
- Selectable marker: plasmid confers resistance to ampicillin (100 $\mu\text{g/ml}$) in *E. coli* hosts.
- *E. coli* replication origin: ColE1
- Copy number: low
- Plasmid incompatibility group: pMB1/ColE1

References

1. Markova, S.V. *et al.* (2004) *J. Bio. Chem.* **279**(5):3212-3217.

Note: The vector sequence was compiled from information in the sequence databases, published literature, and other sources, together with partial sequences obtained by Clontech. This vector has not been completely sequenced.

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Metridia luciferase

Markova, S.V., Golz, S., Frank, L.A., Kalthof, B. & Vysotski, E.S. (2004) Cloning and expression of cDNA for a luciferase from the marine copepod *Metridia longa*. A novel secreted bioluminescent reporter enzyme. *J. Biol. Chem.* **279**(5):3212–3117.

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