



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

pBI-CMV4 Vector Map and Multiple Cloning Site.



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Description

pBI-CMV4 is a mammalian bidirectional expression vector designed to constitutively express a protein of interest and DsRed2, a human codon-optimized variant of the *Discosoma sp.* red fluorescent protein, DsRed (1), that has been engineered for faster maturation and lower non-specific aggregation. The vector allows straightforward detection of transfected mammalian cells by flow cytometry or fluorescence microscopy, as cells expressing the protein of interest can be quickly identified by screening for DsRed2 fluorescence.

Protein expression is driven by one of two constitutively active, minimal human cytomegalovirus promoters: $P_{minCMV1}$ (located upstream of the multiple cloning site [MCS]), drives the expression of the protein of interest, and $P_{minCMV2}$ drives the expression of DsRed2. To allow propagation and selection in *E. coli*, the vector contains a ColE1 origin of replication and an ampicillin resistance gene (Amp^r).

(PR093642; published 3 September 2010)

Use

pBI-CMV4 is designed to constitutively express a protein of interest and the red fluorescent protein DsRed2. The gene of interest must contain an initiation codon and a stop codon.

pBI-CMV4 can be transfected into mammalian cells using any standard transfection method. Cells expressing DsRed2 (excitation and emission maxima: 558 and 583, respectively) can be detected by flow cytometry or fluorescence microscopy 24 hr after transfection. However, in some cases, up to 48 hr may be required for detection of red-emitting cells.

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)
- DsRed2 (human codon optimized): 3017–3693
- P_{minCMV2} (minimal human cytomegalovirus promoter 2): 3711–3779

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
- Plasmid incompatibility group: pMB1/ColE1

References

1. Matz, M. V. *et al.* (1999) *Nat. Biotechnol.* **17**(10):969-973.

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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DH5 α^{TM} is a trademark of Invitrogen Corporation.

The RCFPs (including DsRed-Express and DsRed-Express2) are covered by one or more of the following U.S. Patent Nos. 7,166,444; 7,157,565; 7,217,789; 7,338,784; 7,338,783; 7,537,915 6,969,597, 7,150,979 and 7,442,522.

Living Colors Fluorescent Protein Products:

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