pHSE-SEAP⁺ Vector Information

GenBank Accession No. Submission in progress.

Sold as part of Cat. No. 631910



Restriction Map of pHSE-SEAP. Unique restriction sites are in bold.

Description:

pHSE-SEAP is designed to monitor the activation of heat shock factor (HSF) and heat shockmediated signal transduction pathways. The pHSE-SEAP contains the secreted alkaline phosphatase (SEAP) reporter gene (2–4). This vector also contains three tandem copies of the HSE consensus sequence fused to aTATA-like promoter (P_{TAL}) region from the Herpes simplex virus thymidine kinase (HSV-TK) promoter. After endogenous HSF proteins bind the heat-shock element (HSE; 1), transcription is induced and the reporter gene is activated.

The SEAP coding sequence is followed by the SV40 late polyadenylation signal to ensure proper, efficient processing of the SEAP transcript in eukaryotic cells. Located upstream of HSE is a synthetic transcription blocker (TB), which is composed of adjacent polyadenylation and transcription pause sites for reducing background transcription (5). The vector backbone also contains an f1 origin for single-stranded DNA production, a pUC origin of replication, and an ampicillin resistance gene for propagation and selection in *E. coli*.





United States/Canada 800.662.2566 Asia Pacific +1.650.919.7300 Europe +33.(0)1.3904.6880 Japan +81.(0)77.543.6116

Clontech Laboratories, Inc. ATakara Bio Company 1290 Terra Bella Ave. Mountain View, CA 94043 Technical Support (US) E-mail: tech@clontech.com www.clontech.com

Use:

pHSE-SEAP is designed to measure the binding of transcription factors to HSE, providing a direct measurement of activation for this pathway. For example, exposure of cells to elevated temperatures such as 42°C activates HSF binding to HSE, which initiates transcription of SEAP. Alternatively, you can cotransfect this vector with an expression vector containing your gene of interest to monitor pathway activation. The secreted SEAP enzyme can be assayed directly from the culture medium using one of Clontech's Great EscAPe Chemiluminescence Detection Kits (Cat. Nos. 631701 & 631704). In addition, the SEAP assay permits time-course studies not possible with assays dependent on cell lysates. The pHSE-SEAP Vectors can be transfected into mammalian cells by any standard method. For selecting stable clones, cotransfect with a vector containing an antibiotic resistance gene, such as neomycin, hygromycin, or puromycin.

(PR6X2115; published 12 October 2006)

Location of features:

- Heat shock response element (HSE; 1): 26–70
- TATA-like promoter (P_{TAI}): 76–224
- Secreted alkaline phosphatase (SEAP) gene:
 - SEAP coding sequences: start codon (ATG): 252–254; stop codon: 1809–1811 signal peptide: 252–302 mature protein: 303–1808 C-terminal extension to SEAP: 1770–1808
- SV40 late mRNA polyadenylation signal: 1922–1927 mRNA 3' end: 1941
- pUC plasmid replication origin: 2320-2963
- Ampicillin resistance gene:

Promoter: -35 region: 4041-4036; -10 region: 4018-4013

Transcription start point: 4006

Ribosome binding site: 3983–3979

β-lactamase coding sequences:

start codon (ATG): 3971-3969; stop codon: 3113-3111

β-lactamase signal peptide: 3971-3903

β-lactamase mature protein: 3902-3114

- f1 single-strand DNA origin (packages the noncoding strand of SEAP): 4103–4558
- Transcription blocker (TB): 4689-4842

Synthetic polyadenylation site (6): 4689-4737

Transcription pause site from human α 2 globin gene (7): 4751–4842

Propagation in *E. coli*:

• Suitable host strains: DH5 α and other general purpose strains. Single-stranded DNA production requires a host containing an F' episome such as JM109.

- Selectable marker: plasmid confers resistance to ampicillin (50 µg/ml) in *E. coli* hosts.
- *E. coli* replication origin: pUC
- Copy number: ~500
- Plasmid incompatibility group: pMB1/Col E1

References:

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