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TECHNICAL SUPPORT

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pCpGfree-basic

A mSEAP reporter plasmid without a promoter and devoid of CpG dinucleotides

Catalog # pcpf-bas

For research use only

Version 21F04-MMv02

PRODUCT INFORMATION

Content:

- 20 µg of pCpGfree-basic plasmid provided as lyophilized DNA
- *E. coli* GT115 strain provided lyophilized on a paper disk
- 1 ml of Zeocin™ (100 mg/ml)

Storage and Stability:

- Products are shipped at room temperature.
- Lyophilized DNA is stable 1 year when stored at -20 °C.
- Resuspended DNA is stable 6 months when stored at -20 °C.
- Bacteria should be stored at -20 °C and are stable up to 1 year.
- Store Zeocin™ at 4 °C or at -20 °C. The expiry date is specified on the product label.

Quality control:

Plasmid construct has been confirmed by restriction analysis and sequencing. Plasmid DNA was purified by ion exchange chromatography and lyophilized. Viability of the lyophilized bacteria upon resuspension has been verified.

GENERAL PRODUCT USE

Methylation of CpG dinucleotides within the promoter/enhancer region of genes is often associated with transcriptional silencing. This epigenetic event plays an important role in the regulation of gene activity in normal and cancer cells. Recently, it has been confirmed that the activity of enhancers is correlated with DNA methylation¹.

InvivoGen provides pCpGfree-basic a murine secreted embryonic alkaline phosphatase (mSEAP) reporter plasmid that is completely devoid of CpG dinucleotides and lacks the entire promoter region. It contains a multiple cloning site upstream of the mSEAP reporter gene. Expression of mSEAP in cells transfected with this plasmid depends on the insertion of a functional promoter or enhancer/promoter cassette upstream from the mSEAP gene. Thus, pCpGfree-basic allows to study the effect of CpG methylation on a promoter, alone or combined with enhancer elements.

PLASMID FEATURES

All the elements required for replication and selection of the plasmid in *E. coli* and gene expression in mammalian cells are completely devoid of CpG dinucleotides. Furthermore, all Dam methylation sites (GATC) have been removed to prevent prokaryotic methylation.

Elements for expression in *E. coli*

- Origin of replication: The *E. coli* R6K gamma ori has been modified to remove all CpGs. This origin is activated by the R6K specific initiator protein π , encoded by the *pir* gene².
- Bacterial promoter: EM2K is a CpG-free version of the bacterial EM7 promoter.
- Selectable marker: The Zeocin™ resistance gene is a small gene (<400 bp) that contains numerous CpG dinucleotides. A synthetic new allele was created that contains no CpGs.

Elements for expression in mammalian cells

- The synthetic mSEAPΔCpG gene: a CpG-free allele of the murine SEAP gene constructed by chemical synthesis.
- Polyadenylation signal: The polyadenylation signal is a CpG-free form of the late SV40 polyadenylation signal.

- MAR: Matrix attached regions (MARs) are sequences typically AT-rich that are able to form barriers between independently regulated domains³. pCpGfree plasmids contains two MARs, from the 5' region of the human IFN- β gene or β -globin gene that were chosen because they are naturally CpG-free. The MARs are placed between the bacterial and mammalian transcription units.
- MCS: The multiple cloning site contains several commonly used restriction sites for convenient cloning of a gene of interest.
5' Sda I, Bsp 120I, Avr II, Nsi I, Ppu 10I, Sca I, Bam HI, Spe I, Hind III 3'

Due to the presence of the R6Kγ origin of replication, pCpG plasmids can only be amplified in *E. coli* mutant strain expressing a *pir* mutant gene. They will not replicate in standard *E. coli* strains. Therefore, pCpG plasmids are provided with the *E. coli* GT115 strain, a *pir* mutant also deficient in *Dcm* methylation.

1. Hoivik EA. et al., 2011. DNA Methylation of Intronic Enhancers Directs Tissue-Specific Expression of Steroidogenic Factor 1/Adrenalin 4 Binding Protein (SF-1/Ad4BP). Endocrinology, 152(5):2100-12.
2. Wu F. et al. 1995. A DNA segment conferring stable maintenance on R6K gamma-origin core replicons. J Bacteriol. 177(22):6338-45.
3. Bode J. et al., 1996. Scaffold/matrix-attached regions: topological switches with multiple regulatory functions. Crit Rev Eukaryot Gene Expr. 6(2-3):115-38.

METHODS

Plasmid resuspension

Quickly spin the tube containing the lyophilized plasmid to pellet the DNA. To obtain a plasmid solution at 1 µg/µl, resuspend the DNA in 20 µl of sterile H₂O. Store resuspended plasmid at -20 °C.

Reconstitution of *E. coli* GT115 strain

Use sterile conditions to do the following:

1. Reconstitute *E. coli* GT115 by adding 1 ml of Luria-Bertani (LB) medium in the tube containing the paper disk. Let sit for 5 minutes.
2. Mix gently by vortexing for 1-2 minutes.
3. Streak bacteria taken from this suspension on a LB agar plate.
4. Place the plate in an incubator at 37°C overnight.
5. Isolate a single colony and grow the bacteria in LB or terrific broth (TB) medium.
6. Prepare competent cells utilizing protocol of choice.

Plasmid amplification and cloning

Plasmid amplification and cloning can be performed in *E. coli* GT115.

Zeocin™ usage

This antibiotic can be used for *E. coli* at 25 µg/ml in liquid or solid media.

TECHNICAL SUPPORT

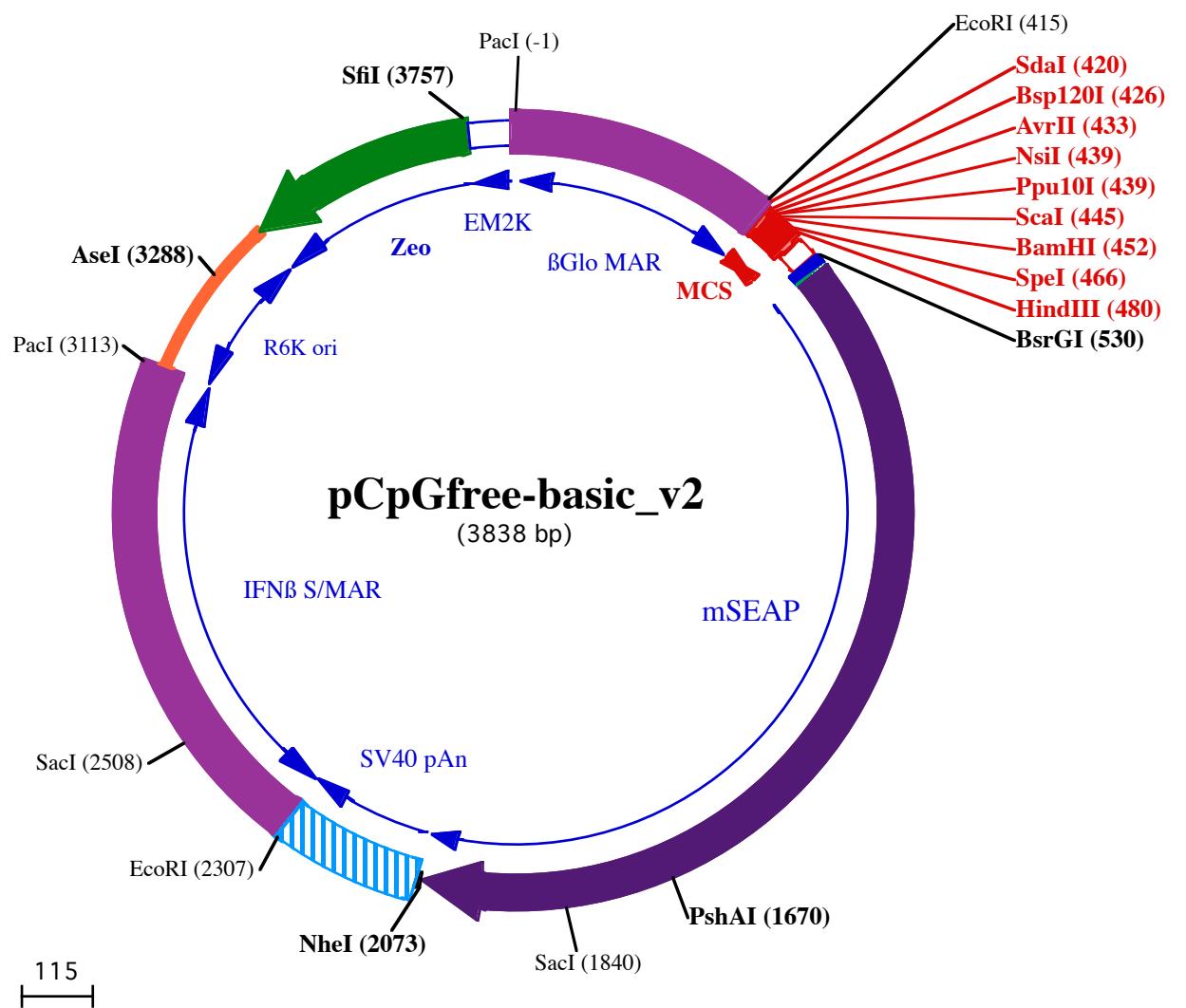
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PacI (-1)

1 TTAATTAAAATTCCTAAGGCATGTGAACGGCTGCTTGGTTTCACTGTACTTCATCTGCTACCTCTGTGACCTGAAACATATTATAATTCCAT

101 TAAGCTGTCATATGATAGATTATCATATGTATTTCTTAAAGGATTTTGTAAGAACTAATTGAATTGATACCTGAAAGCTTATCACACTACCC

201 AATAAATAATAATCTTGTTCAGCTCTGTTCTATAAATATGACCAGTTTATTGTTTAGTGGTAGTGTATTCTCTTCTATATAT

301 ACACACACATGTGTGATTCTATAATATACAATTGGAAATAAAATTAGCAATCAATTGAAACACTGATTGGTTATGTGAGCAA

Bsp120I (426) Ppu10I (439)
SdaI (420) NsiI (439) BamHI (452)
EcoRI (415) AvrII (433) ScaI (445) SpeI (466) HindIII (480)

401 ACAGCAGATTAAAAGGAATTCTGCAGGGGCCACCTAGGATCATGACTAGGATCCAACATGTAACTAGTAGCATGCAAAGCTTAGAAAttgtactaac

501 cttcttccttcctccatcag GTTGGTGTACAGTAGCTTCCACATGTGGGTGCCTGCTATTGCTGGCTTAAGTCTCAAGTTGCCA
1 ▶ Me tT rpG lyA laCysLeuLeuLeuLeuG lyLeuSe rLeuG InVa lCysP roS

601 GTGTCATTCTGTGGAGGAGGAATCCTGCTTTGGAAATAGGAAGGGCAGCTGAAGCCTGGATGCAGCCAAGAACGCTCAAGCCCATTAGACATCTGC
18 ▶ e rVa I leP roVa IG luG luG luAsnP roA laPheT rpAsnArgLysA laA laG luA laLeuAspA laA laLysLysLeuLysPro I leG InTh rSe rA I

701 AAAGAATCTTGTTCATCTCATGGGTGATGGAATGGGTGTCCTCAGTAAACAGCCACAGGATTCTGAAGGGCCAGCAACAAGGTCATCTAGGCCAGAG
51 ▶ aLysAsnLeuVa I leLeuMe tG lyAspG lyMe tG lyVa I se rTh rVa I Th rA laTh rArg I leLeuLysG lyG InG InG lyH isLeuG lyP roG lu

801 ACCCAGTTGCAATGGACAGGTTCCCTACATGGCCCTTCCAAGACTACACAGTACAAGCAGATTCTGACTCTGCTGGACAGGCACAGCATTCT
85 ▶ Th rG InLeuA laMe tAspArgPheP roH isMe tA laLeuSe rLysTh rTy rAsnTh rAspLysG In I leP roAspSe rA laG lyTh rG lyTh rA laPheL

901 TGTGTGGAGTAAAACCAACATGAAAGTCATTGGTCTTCAGCTGCTGCCAGATTCAACAGTGCAACACCACATGGGCAATGAAGTGGTCTCTGTAAT
118 ▶ euCysG lyVa I LysTh rAsnMe tLysVa I leG lyLeuSe rA laA laA laArgPheAsnG InCysAsnTh rTh rTpG lyAsnG luVa Ma I se rVa Me

1001 GCACAGGGCAAAAGCTGGGAAAGTGTGGGTGTTGACAACCACCTCTGTCAGCATGCCTCTCTGCTGGAACTTACAGGACATGCCACACAGTGAACAGA
151 ▶ th isArgA laLysLysA laG lyLysSe rVa IG lyVa Ma I Th rTh rH rSe rVa IG InH isA laSe rP roA laG lyTh rTy rA laH isTh rVa AsnArg

1101 GGTTGGTACTCTGATGCTCAGATGCCCTCAGTTACAAGATGGCTGCAAGGACATCAGCACCCAGCTCATCTAAACATGGACATAGATGTCATCT
185 ▶ G lyT rpTy rSe rAspA laG InMe tP roA laSe rA laLeuG InAspG lyCysLysAsp I leSe rTh rG InLeu I leSe rAsnMe tAsp I leAspVa I leL

1201 TAGGGGGTGGAGAAAGTCATGTTCCAAAGGGACTCTGACCAAGGAGTACCCACAGACACAAAGCAGGCTGGCACAAAGATTAGTGTAGGAACCT
218 ▶ eug lyG lyA ArgLysPheMe tPheP roLysG lyTh rP roAspG InG luTy rP roTh rAspTh rLysG InA laG lyTh rA rgLeuAspG lyA rgAsnLe

1301 TGTGCAAGAGTGGCTTGCAGCATCAGGGAGCAAGGTATGCTGGACAGGAGTACCTGGCTTGAACAGGTGTCACCTCACCTAATG
251 ▶ uVa IG InG luT rpLeuA laLysH isG InG lyA laArgTy rVa I T rpAsnArgSe rG luLeu I leG InA laSe rLeuAsnA rgSe rVa I Th rH isLeuMe t

1401 GGTTATTTGAGCCAATGACATGAAGTATGAGATACACAGGGACCTGGCCAGGACCCCTCTAGCAGAAATGACTGAAGTTGCTGTGAGGATGTTG
285 ▶ G lyLeuPheG luP roAsnAspMe tLysTy rG lu I leH isArgAspP roA laG InAspP roSe rLeuA laG luMe tTh rG luVa IA laVa I A rgMe tLeuS

1501 CCAGAAATCCAAAGGGTTACCTCTTGTGAGGGGGAGGATTGATCATGGTACCATGAGACAGTTGCTTACAGAGCTTAAC TGAGGCTGTGAT
318 ▶ e rArgAsnP roLysG lyPheTy rLeuPheVa IG luG lyG lyArg I leAspH isG lyH isH isG luTh rVa IA laTy rArgA laLeuTh rG luA laVa Me

PshAI (1670)

1601 GTTGTGATTCTGTGGACAAGGCTGACAAACTGACCTGAGCAGGACACAATGATTCTAGTGACTGCTGACCAAGCTCATGTTCTCTTGGGGC
351 ▶ tPheAspSe rA laVa lAspLysA laAspLysLeuTh rSe rG luG InAspTh rMet I leLeuVa I Th rA laAspH isSe rH isVa lPheSe rPheG lyG ly

1701 TACACCCAGAGGGTGCTTCAATCTTGGCTGGCCCTTCAAGGCAGAAGATGGAAAGAGTTTACCTCCATCTATGGGAATGGCTGGTACA
385 ▶ Ty rTh rG InArgG lyA laSe rI lePheG lyLeuA laP roPheLysA laG luAspG lyLysSe rPheTh rSe rI leLeuTy rG lyAsnG lyP roG lyTy rL

SacI (1840)

1801 AGCTGCACAATGGGCCAGAGCTGATGTGACAGAAGAGGAGGCTCAACCAACCTACAGCAGCAAGCAGCAGTCCCTTTCTTCAAGAAACCAACTC
418 ▶ ysLeuH isAsnG lyA laArgA laAspVt I Th rG luG luG luSe rSe rAsnProTh rTy rG InG InG InA laA laVa I P roLeuSe rSe rG luTh rH isSe

1901 TGGGAAGATGTGGCATATTGCCAGAGCCCCCAAGCCCCTGGTGCATGGAGTTCAAGGAGCAGAAATTACATAGCTCATGTAATGGCTTTGCTGCT
451 ▶ rG lyG luAspVa IA la I lePheA laArgG lyP roG luA laH isLeuVa H isG lyVa IG InG luG InAsnTy r I leA laH isVa Me tA laPheA laA la

NheI (2073)

2001 TGCTGGAGCCCTACACAGACTGTGGCTAGCCAGGCCAGCAGGCCAGTCCTCTGAGTAAGCCCAGGCTAGAGCTAGCTGGCCAGACATGATAAGATAC
485 ▶ CysLeuG luP roTy rTh rAspCysG lyLeuA laSe rP roA laG lyG InSe rSe rA laVa I se rP roG ly***

2101 ATTGATGAGTTGGACAAACACAAACTAGAATGCACTGAGTAAAAAAATGCTTATTGTGAAATTGTGATGCTATTGCTTATTGTAACCTATAAGCT

2201 GCAATAAACAGTTAACACAAATTGCAATTCTATTGTTAGGTTAGGGGAGGTGGAGGTTAAAGCAAGTAAACCTCTACAAATG

EcoRI (2307)

2301 TGGTATGGAATTCACTGCAATATGTTACCCCCAAAAAGCTGTTAAGTGCACCTCATTCTAAATGTATAGAAGCCAAAAGACAATAACAA

2401 AAATATTCTGTAGAACAAATGGAAAGAATGTTCACTAAATATCAAGATTAGAGCAAGCATGAGATGTGGGGATAGACAGTGGCTGATAAA

SacI (2508)

2501 ATAGAGTAGAGCTAGAACAGACCCATTGATATATGTAAGTGACCTATGAAAAAAATGGCATTTACAATGGAAAATGATGGCTTTCTTTT

2601 AGAAAAACAGGAAATATTTATATGAAAAAAATAAAAGGGACCCATATGTCATACCATACACACAAAAAAATCCAGTGAATTATAAGCTAAATGG

2701 AGAAGGCAAACCTTAAATCTTTAGAAAATATAGAACGATGCCATCAAGACTTCAGTGTAGAGAAAAATTCTTATGACTCAAAGTCTAACACAC

2801 AAGAAAAGATTGTTAATTAGATTGCAATATAGACTTATTAAAATAAAAACCTTAAGAAAAGTCAGGCCATAGAATGACAGAAAATTT

2901 GCAACACCCAGTAAAGAGAATTGTAATATGCAATTAAAAAGAAGTCTTACAAATCGTAAAAAAACTAGACAAAATTGACAGATGAAAG

3001 AGAAACTCTAATAATCATTACACATGAGAACTCAATCTCAGAAATCAGAGAACTATTCATGATACACTAAATTAGAGAAATTAAAAGGCTAAG

PacI (3113)

3101 TAACATCTGTGGCTTAATTAAATCAGCAGTTAACCTGTTGATAGTGTACTAAGCTCATGTTAATGTTACTAAGCTCTATGTTAATGAACAA

AseI (3288)

3201 ACCCTCATGGCTAATGTTACTAAGCTCTATGGCTAATGTTACTAAGCTCTATGTTCATGTTACTAAGCTCTATGTTGAACAATAAAATTAAAT

3301 CAGCAACTAAATAGCTCTAAGGTTTAAGTTATAAGAAAAAAAAGAATATATAAGGTTAAAGGTTAAAGGTTCTAGCTTAGTCCTGTT

3401 CTCAGCTACAAATGGACACAATTCCAGCAGGGCTCTGAGGGCAAATCCCTCCCCAAGGTTGTCACCAATTCTGTATGGCTGGGCCAGAGGCA

3501 TCCCTGAAATTGTGCTGACTACTTGACCATCTGCATAAAGCTCATCTAGGCTCTGACCCAGACCAAGCAAGGGTTGTCAGGGACAACCTGGT

3601 CCTGAAC TGCTGAGATGAAGAGGGTGACATCATCTGACAACACCAGAAAATCATTTCAACAAAGTCTGGAGAATCTTAATCTGTCAGTCCAGAA

SfiI (3757)

3701 CTCTACAGCCCTGCAACATCCCTGCTGTGAGGACTGGACTGCAGAAGTGA GATGGCCCTCTAGTGA GTTGATTATACTATGCA

3801 GATATACTATGCCAATGTTAATTGTCAACTACCTGTT
