

pSELECT-zeo-GFP::Sh

A plasmid encoding a CpG-free GFP-Zeocin resistance fusion gene

Catalog code: psetz-zgfpsh

For research use only

Version 23F26v03-JC

PRODUCT INFORMATION

Content:

- 20 µg of pSELECT-zeo-GFP::Sh plasmid provided as lyophilized DNA.
- 1 ml of Zeocin™ (100 mg/ml)

Storage and Stability:

Product is shipped at room temperature. Lyophilized DNA should be resuspended upon receipt and stored at -20°C. Lyophilized DNA is stable 3 months at -20°C. Resuspended DNA is stable more than one year at -20°C.
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.

GENERAL PRODUCT USE

pSelect-zeo plasmids contain genes that have been chemically synthesized. The DNA sequence of these genes was modified by optimizing the codon usage, reducing or eliminating the CpG motifs and avoiding secondary DNA structures without changing the amino acid sequence of the wild type proteins.

pSelect-zeo plasmids may be used:

To subclone the synthetic gene into another vector. To facilitate subcloning, the GFP::Sh gene is flanked by two unique restriction sites: Nco I at the 5' end that encompasses the Start codon, and Nhe I at the 3'end.

As a gene reporter plasmid. pSelect-zeo is a mammalian expression plasmid selectable in *E. coli* and mammalian cells with Zeocin™, as the *Sh ble* gene in the second expression cassette is driven by the eukaryote CMV enhancer/promoter in tandem with the bacterial EM7 promoter.

PLASMID FEATURES

First expression cassette

• **hEF1-HTLV prom** is a composite promoter comprising the Elongation Factor-1alpha (EF-1 α) core promoter¹ and the R segment and part of the U5 sequence (R-U5') of the Human T-Cell Leukemia Virus (HTLV) Type 1 Long Terminal Repeat². The EF-1 α promoter exhibits a strong activity and yields long lasting expression of a transgene *in vivo*. The R-U5' has been coupled to the EF-1 α core promoter to enhance stability of RNA.

• **GFP::Sh CpG-free:** InvivoGen has engineered a fusion gene between the red-shifted variant of the jellyfish GFP gene that encodes a green fluorescent protein and the *Sh ble* gene conferring Zeocin™ resistance. Both genes have been modified and contain no CpG motifs, whereas their wildtype counterparts contain 60 and 50 CpG motifs respectively. This GFP::Sh fusion protein absorbs blue light (major peak at 480 nm) and emits green light (major peak at 505 nm).

• **SV40 pAn:** the Simian Virus 40 late polyadenylation signal enables efficient cleavage and polyadenylation reactions resulting in high levels of steady-state mRNA³.

• **ori:** a minimal *E. coli* origin of replication to limit vector size, but with the same activity as the longer Ori.

Second expression cassette

• **CMV enh/prom:** The human cytomegalovirus immediate-early gene 1 promoter/enhancer was originally isolated from the Towne strain and was found to be stronger than any other viral promoters.

• **EM7** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*.

• **Zeo:** Resistance to Zeocin™ is conferred by the *Sh ble* gene from *Streptallosteichus hindustanus*. The *Sh ble* gene is driven by the CMV enhancer/promoter in tandem with the bacterial EM7 promoter allowing selection in both mammalian cells and *E. coli*.

• **BGlo pAn:** The human beta-globin 3'UTR and polyadenylation sequence allows efficient arrest of the transgene transcription⁴.

1. Kim, D.W. *et al.* (1990). Gene 2: 217-223.

2. Takebe, Y. *et al.* (1988). Mol. Cell Biol. 1: 466-472.

3. Carswell, S. & Alwine, J.C. (1989). Mol. Cell Biol. 10: 4248-4258.

4. Yu J & Russell JE. (2001). Mol Cell Biol, 21(17):5879-88.

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.

Plasmid amplification and cloning

Plasmid amplification and cloning can be performed in *E. coli* GT116 other commonly used laboratory *E. coli* strains, such as DH5α.

Zeocin™ usage

This antibiotic can be used for *E. coli* at 25 µg/ml in liquid or solid media and at 50-200 µg/ml to select Zeocin™-resistant mammalian cells.

TECHNICAL SUPPORT

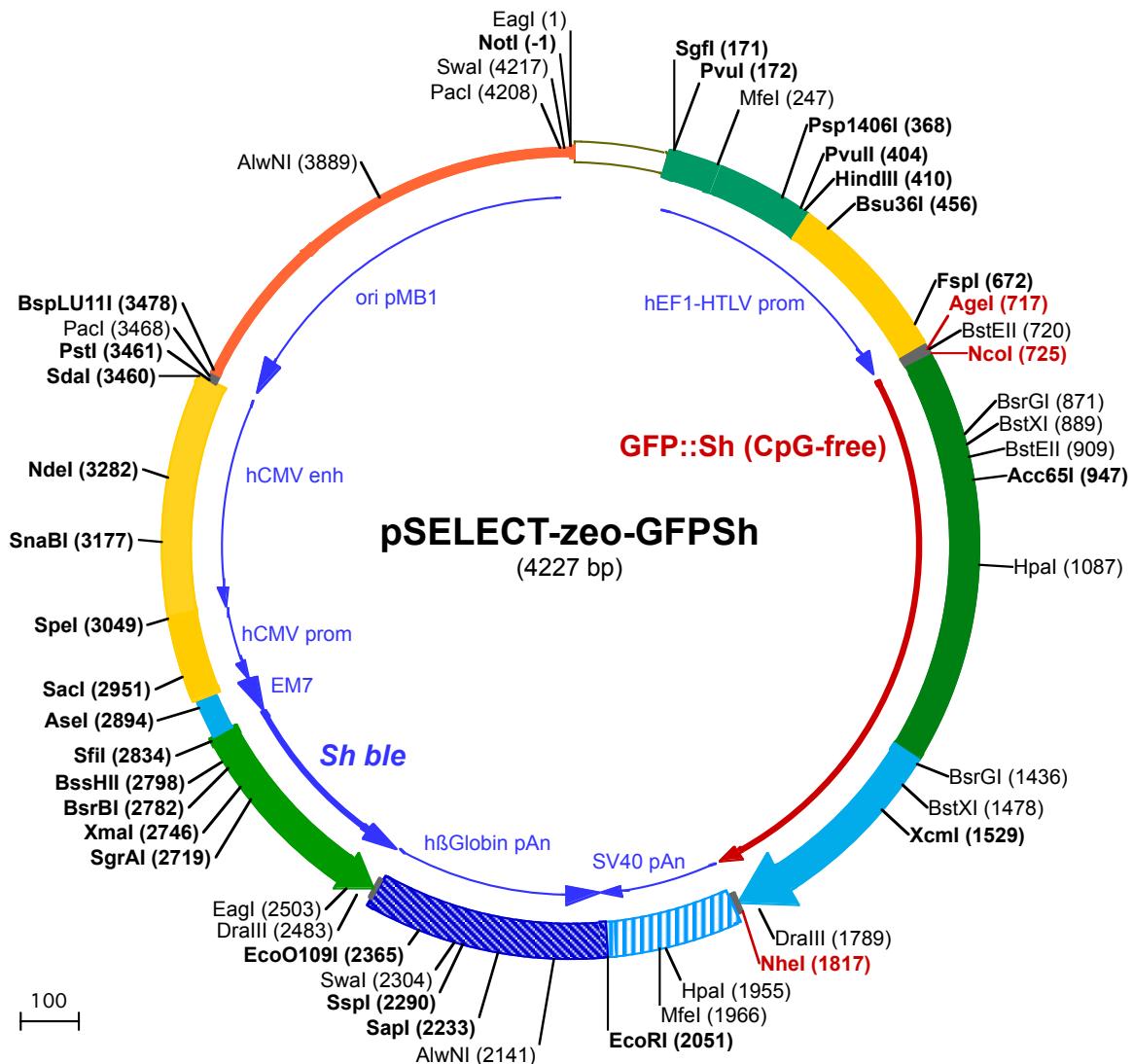
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InvivoGen Europe: +33 (0) 5-62-71-69-39

InvivoGen Hong Kong: +852 3622-3480

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EagI (1)
NotI (-1)

1 GCGGCCGAATAAAATCTTATTTATTACATCTGTGTTGGTTGGTGTGAATCGTAACATACGCTCTCCATCAAAACAAAACGAAACA

PvuI (172)
SgfI (171)

101 AAACAAACTAGCAAAATAGGCTGCCCCAGTCAAGTGCAGGTGCCAGAACATTCTCATCGAAGGATCTGATCGCTCCGGTGCCCGTCAGTGGGCA

MfeI (247)

201 GAGCGCACATCGCCCACAGTCCCCGAGAAGTTGGGGGAGGGTGGCAATTGAACGGGTGCTAGAGAAGGTGGCGGGTAAACTGGAAAGTGATG

Psp1406I (368)

301 TCGTGTACTGGCTCCGCTTTCCGAGGGTGGGGAGAACCGTATATAAGTGCAGTAGTCGCCGTGAACTGTTCTTCGCAACGGTTGCCAG

HindIII (410)

PvuII (404)
Bsu36I (456)

401 AACACAGCTGAAGCTTCGAGGGGCTCGATCTCTCCTCACGCGCCGCCACTGAGGCCCATCCACGCCGGTTGAGTCGCTCTGCCGCT

501 CCCGCCTGTTGCTCTGAACTCGCTCCGCTAGGTAAGTTAAAGCTCAGGTCGAGACCGGGCCTTGTCCGGCCTCCGGACCTACCTA

FspI (672)

601 GACTCAGCCGGCTCCACGCTTGCTGACCCCTGCTGCTCAACTCTACGTCTTGTTCGTTCTGTCGCAAGTACAGATCCAAGCTGTGACC

BstEII (720)

AgeI (717) NcoI (725)

701 GGCCTACCTGAGATCACCGTACCAATGGTTCTAAGGGAGAAGAACCTTTACTGGTGTGCAATTCTGGTTGAGCTGGATGGTATGTGAATG

1▶ M V S K G E E L F T G V V P I L V E L D G D V N

BsrGI (871) BstXI (889)

801 GCCACAAATTCTCTGTGCTGGTGAAGGTGAAGGGAGATGCAACTTATGAAAGCTGACTCTGAAGTTCAATTGTCAGACAGGAAAGCTGCCAGTGCCTTG

25▶ G H K F S V S G E G E G D A T Y G K L T L K F I C T T G K L P V P W

BstEII (909) Acc65I (947)

901 GCCAACTCTGGTGAACCCCTGACTTATGGTGTCAATGTTCACTGGTGTGCAATTGTCAGGATCCACATGAAGCAGCATGACTCTTAAATCTGCAATGCCAGAA

58▶ P T L V T T L T Y G V Q C F S R Y P D H M K Q H D F F K S A M P E

HpaI (1087)

1001 GTTATGTCAGGAGAGAACATTCTTAAGGATGATGAAATTATAAGACAAGGGCAGAAGTGAAGTTGAAGGTGATACTGGTAAACAGAAATTG

92▶ G Y V Q E R T I F F K D D G N Y K T R A E V K F E G D T L V N R I

1101 AGCTGAAAGGCATTGATTTAAGGAAGATGAAACATTCTGGGTACAAGCTGGAGTACAACATAATTCTACAATGTTACATTGGCAGATAAGCA

125▶ E L K G I D F K E D G N I L G H K L E Y N Y N S H N V Y I M A D K Q

1201 GAGGAATGAAATTAAAGGCTAATTCAAGATTAGACACAAACATTGAGGATGGATCTGCAACTGGCAGACCAATTACAGCAGAACACCCCTATTGGTGT

158▶ R N G I K A N F K I R H N I E D G S V Q L A D H Y Q Q N T P I G D

1301 GGCCAGTTCTCTCCAGATAATCACTATCTCAGCACTCAATCTGTCCTGCAAAGACCTAATGAGAAAAGAGACCATGGTCTCTGGAGTTG

192▶ G P V L L P D N H Y L S T Q S A L S K D P N E K R D H M V L L E F

BsrGI (1436) BstXI (1478)

1401 TGACAGCAGCAGGAAATTACTCTGGATGGATGAGCTGACAAGGCAAGTTGACAGTGCTGCTCCAGTGCTCACAGCAGGGATGGCTGGAGCTGT

225▶ V T A A G I T L G M D E L Y K A K L T S A V P V L T A R D V A G A V

XcmI (1529)

1501 TGAGTTGGACTGACAGGTTGGGTTCTCAGAGATTGGAGGATGACTTGCAAGGTGAGATGATGTCACCCGTTCATCTCAGCAGTC

258▶ E F W T D R L G F S R D F V E D D F A G V V R D D V T L F I S A V

1601 CAGGACCAAGGGTGGTGCCTGACAACACCCCTGGCTGGGTGGAGAGGACTGGATGAGCTGTGAGTGGAGTGGAGTGGCTCCACCAACTTC

292▶ Q D Q V V P D N T L A W V W V R G L D E L Y A E W S E V V S T N F

DraIII (1789)

1701 GGGATGCCAGTGGCCCTGCCATGACAGAGATTGGAGAGCAGCCCTGGGGAGAGAGTTGCCCTGAGAGACCCAGCAGGCACTGTGCACTTGTG

325▶ R D A S G P A M T E I G E Q P W G R E F A L R D P A G N C V H F V A

NheI (1817)

1801 AGAGGAGCAGGACTAAAGCTAGCTGGCCAGACATGATAAGATAACATTGATGAGTTGGACAAACCAACTAGAATGCACTGAGTAAAAAATGCTTATTG

358▶ E E Q D •

HpaI (1955) MfeI (1966)

1901 TGAATTGTGATGCTATTGTTATTGTAACCATTAAAGCTGCAATAAACAAAGTTAACAAACATTGATTCTGTTATGTTCAAGTTCAAGGG

EcoRI (2051)

2001 GAGGTGTGGAGGTTTTAAAGCAAGTAAACCTCTACAAATGTGGATGGATTCTAAACATACGGCATAGCAAAACTTAACTCCAAATCAAGCCTC

2101 TACTTGAATCCTTCTGAGGGATGAATAAGGCATAGGCATCAGGGCTTGTGCAATGTGATTAGCTGTTGAGCCTCACCTCTTATGGAGTT

SapI (2233)
SspI (2290)

2201 AAGATATAGTGTATTCCCAAGGTTGAACTAGCTCTCATTTCTTATGTTAAATGCACTGACCTCCACATTCCCTTTAGTAAAATATTCAAG

Swal (2304)
EcoO109I (2365)

2301 AATAATTAAATACATCATTGCAATGAAATAATGTTTTATTAGGCAGAATCCAGATGCTCAAGGCCCTCATAATATCCCCAGTTAGTTAGTGG

DraIII (2483)

2401 ACTTAGGAACAAAGGAACCTTAATAGAAATTGGACAGCAAGAAAGCGAGCTCTAGCTTATCCTCAGTCCTGCTCCACCAAAGTCACGCAGT
1254 • D Q E E A V F H V C N

EagI (2503)

2501 TGCCGGCCGGGTCGCGAGGGCGAACTCCGCCACGGCTGCTGCCATCTCGGTATGGCCGGCCGGAGGCCTCCAGGAAAGTTCTGGACACGAC
1134 G A P D R L A F E R G W P Q E G I E T M A P G S A D R F N T S V V

2601 CTCCGACCACCTCGCGTACAGCTCGCCAGGCCGCGACCCACACCCAGGCCAGGGTGTGCTGGCACACCTGGCTCTGGACCGCGCTGATGAACAGG
804 E S W E A Y L E D L G R V W V W A L T N D P V V Q D Q V A S I F L

SgrAI (2719) XmaI (2746) BsrBI (2782) BssHII (27 98)
2701 GTACGTCGTCCGGACCACCCGGCGAAGTCGTCTCCACGAAGTCCGGAGAACCCGAGCCGGTCCAGAACCTGACCGCTCCGGCACGTCGC
464 T V D D R V V G A F D D E V F D R S F G L R D T W F E V A G A V D R

SfiI (2834) Asel (2894)
2801 GCGCGGTGAGCACCGAACGGCACTGGTCAACTTGCCATGATGGCCCTCTATAGTGAGTCGTATTATACTATGCCGATATACTATGCCGATGATTAAT
134 A T L V P V A S T L K A M ← SacI (2951)

2901 TGTCAAACAGCGTGGATGGCGTCTCCAGCTTATCTGACGGTTACTAAACGAGCTGCTTATAGACCTCCCACCGTACACGCCCTACCGCCATTG
← Spel (3049)

3001 CGTCAATGGGGCGGAGTTTACGACATTGGAAAGTCCCGTTGATTACTAGTCAAACAAACTCCATTGACGTCAATGGGTGGAGACTGGAAAT
← SnaBI (3177)

3101 CCCCGTGAGTCAAACCGCTATCCACGCCATTGATGTACTGCCAAAACCGCATCATGGTAATAGCGATGACTAATACGTAGATGACTGCCAAGTAG
← NdeI (3282)
3201 GAAAGTCCCATAAGGTATGTACTGGGCATAATGCCAGGGGGCATTACCGTATTGACGTCAATAGGGGGTACTTGGCATATGATACATTGATG
3301 TACTGCCAAGTGGCAGTTACCGTAAATACTCCACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAACATACGTATTGACGTC
← Pacl (3468)
3401 AATGGGCAGGGGCTGGGGCGTCAGCCAGGCCATTACCGTAAGTTATGTAACGCTTGAGCTTAAAGACATGTGAGCAAAGGCCAGCA
PstI (3461) Sdal (3460) BspLU11I (3478) ← AlwNI (3889)
3501 AAAGGCCAGGAACCGTAAAAGGCCGTTGCTGGCTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAATCGACGCTCAAGTCAGAGGTGG
3601 CGAAACCCGACAGGACTATAAGATACCAGGCCTTCCCTGGAAGCTCCCTGCGCTCTCTGTTCCGACCCCTGCCCTACCGGATACCTGCTCG
3701 CCTTCTCCCTCGGAAGCGTGGCCTTCATAGCTCACGCTGTAGGTATCTCAGTCGGTAGGTCTGCTCCAAGCTGGCTGTGACGA
3801 ACCCCCCGTTCAGCCGACCGCTGCCCTATCCGTAACTATCGTCTGAGTCCAACCCGTAAGACACGACTTATGCCACTGGCAGCAGCCACTGGT
3901 AACAGGATTAGCAGAGCGAGGTATGTTAGGGCGTACAGAGTTGAAGTGGGGCTAACTACGGCTACACTAGAAGAACAGTATTGGTATCTGCG
4001 CTCTGCTGAAGCCAGTTACCTCGGAAAAAGAGTTGGTAGCTTGTACCGGCAAACAAACCACCGCTGGTAGCGGTGTTTGTGCAAGCAGCA
4101 GATTACGCGCAGAAAAAAAGGATCTAAGAAGATCCTTGATCTTGTACGGGTCTGACGCTCAGTGGAAACAAACTACGTTAAGGGATTTGGTC
← Pacl (4208) Swal (4217)
4201 ATGGCTAGTTAATTAACATTAAATCA