

pSELECT-NHis-blasti

Plasmid for the expression of polyhistidine (His)-N-terminal tagged proteins

Catalog code: psetb-nhis

<https://www.invivogen.com/pselect-his-tag>

For research use only

Version 20I17-MM

PRODUCT INFORMATION

Contents

- 20 µg of pSELECT-NHis-blasti plasmid provided as lyophilized DNA
- 2 x 1 ml blasticidin at 10 mg/ml

Storage and stability

- Product is shipped at room temperature.
- Upon receipt, store lyophilized DNA at -20°C.
- Resuspended DNA should be stored at -20°C.
- Store blasticidin at 4°C or -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 plasmids are specifically designed for strong and constitutive expression of a gene of interest in a wide variety of cell lines. They allow the selection of stable transfecants and offer a variety of selectable markers. pSELECT plasmids contain two expression cassettes: the first drives the expression of the gene of interest and the second drives the expression of a large choice of dominant selectable markers for both *E. coli* and mammalian cells. They are both terminating with a strong polyadenylation signal (polyA) that separates the two expression cassettes thus preventing any transcription interference. The late SV40 polyA terminates the transcription of the gene of interest while the human β-globin polyA terminates the transcription of the selectable marker.

pSELECT-Tag is a new family of expression plasmids designed to generate tagged proteins in order to facilitate their detection and/or purification. pSELECT-Tag features three well-known tags: the green fluorescent protein (GFP) gene, the human influenza hemagglutinin (HA) epitope and the polyhistidine (His) tag. pSELECT-Tag plasmids allow to add the tag either at the N or C terminus of the protein of interest. The N-terminal tag: the tag encompasses the Start codon and is followed by a multiple cloning site (MCS).

PLASMID FEATURES

First expression cassette

- hEF1-HTLV prom is a composite promoter comprising the Elongation Factor-1α (EF-1α) 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-1α promoter exhibits a strong activity and yields long lasting expression of a transgene *in vivo*. The R-U5' has been coupled to the EF-1α core promoter to enhance stability of RNA.

- MCS: The multiple cloning site contains the following restriction sites: 5' - Age I, Sal I, Eco47 III, Nco I, BamH I - 3'

Each restriction site is compatible with many other enzymes.

- 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*.

- Blasti: Resistance to Blasticidin S is conferred by the *bsr* gene from *Bacillus cereus*. The *bsr* 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.

CLONING STRATEGY

For expression of a tagged protein, it is important to ensure to clone your gene-of-interest into the correct reading frame. In general it is recommended to use BamHI/NheI restriction site combination for cloning into plasmids with the N-Tag. For the plasmids with N-Tag, check whether your gene of interest and the start codon of the N-tag are in the correct reading frame.

Note: The Bam HI restriction site is compatible with BgIII.



If it is not possible to use the NheI restriction site, it is possible to use another restriction site such as Ncol.



Alternatively, blunt end cloning can be achieved using the Eco47III site, which is downstream of the BamHI site. Blunt end PCR fragments can be obtained using T4 DNA polymerase or Klenow.



TECHNICAL SUPPORT

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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 or other commonly used laboratory *E. coli* strains, such as DH5α.

Blasticidin usage

Blasticidin should be used at 25-100 µg/ml in bacteria and 1-30 µg/ml in mammalian cells. Blasticidin is supplied at 10 mg/ml in HEPES buffer.

References:

1. Kim, D.W. et al., 1990. Use of the human elongation factor 1 alpha promoter as a versatile and efficient expression system. *Gene* 2: 217-223.
2. Takebe, Y. et al., 1988. SR alpha promoter: an efficient and versatile mammalian cDNA expression system composed of the simian virus 40 early promoter and the R-U5 segment of human T-cell leukemia virus type 1 long terminal repeat. *Mol. Cell Biol.* 1: 466-472.
3. Carswell, S., & Alwine, J.C., 1989. Efficiency of utilization of the simian virus 40 late polyadenylation site: effects of upstream sequences. *Mol. Cell Biol.* 10: 4248-4258.
4. Yu J & Russell JE., 2001. Structural and functional analysis of an mRNP complex that mediates the high stability of human beta-globin mRNA. *Mol Cell Biol*, 21(17):5879-88.

RELATED PRODUCTS

Product	Catalog Code
pSELECT-NGFP-blasti	psetb-ngfp
pSELECT-CGFP-blasti	psetb-cgfp
pSELECT-NGFP-zeo	psetz-ngfp
pSELECT-CGFP-zeo	psetz-cgfp
pSELECT-NHA-blasti	psetb-nha
pSELECT-CHA-blasti	psetb-cha
pSELECT-NHA-zeo	psetz-nha
pSELECT-CHA-zeo	psetz-cha
pSELECT-NHis-zeo	psetz-nhis
pSELECT-CHis-blasti	psetb-chis

TECHNICAL SUPPORT

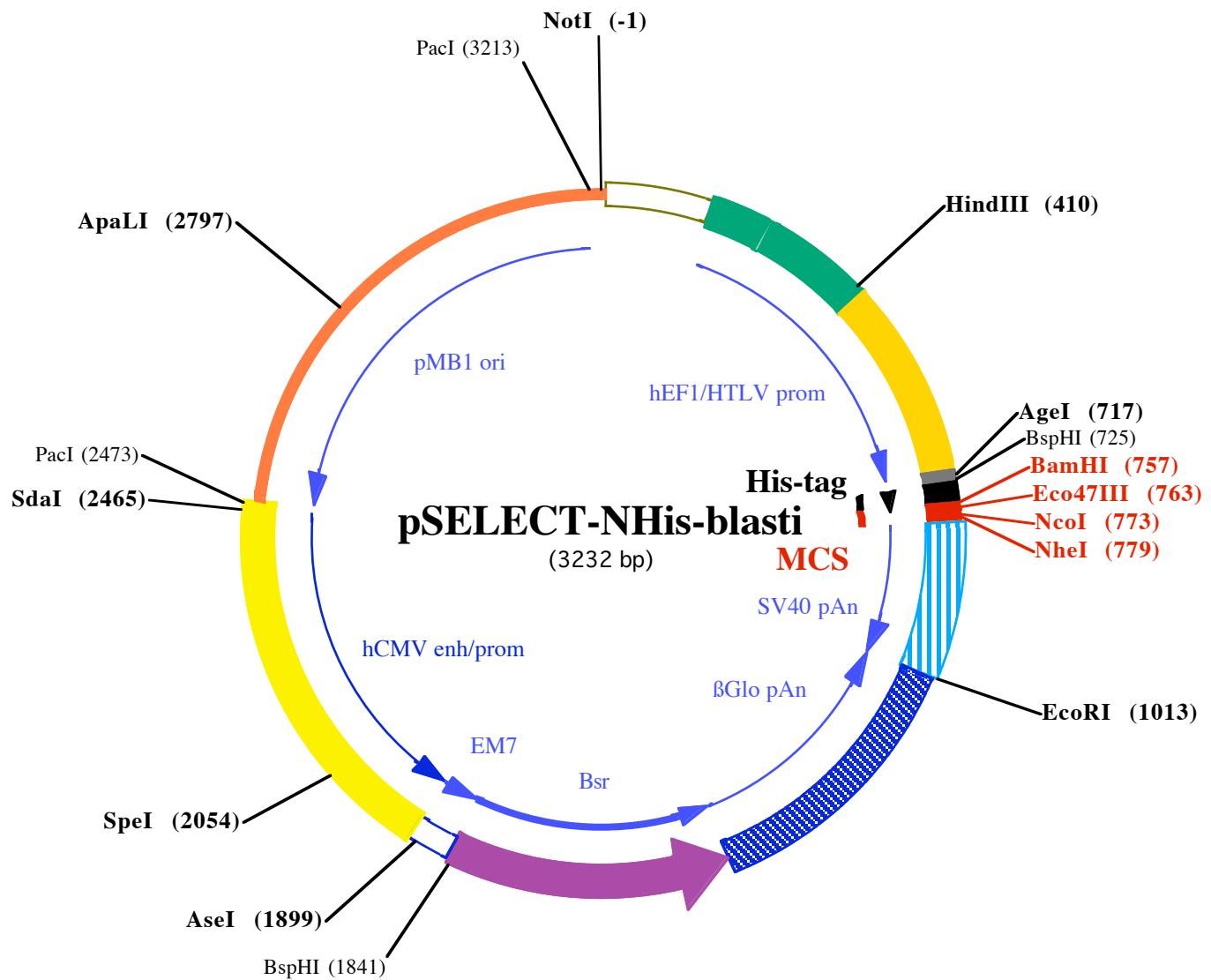
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NotI (-1)

1 **GC****GCCCCG**CATAAAATATCTTATTTCATTACATCTGTGTGGTTTTGTGAATCGTAAC

71 ATACGCTCTCCATCAAACAAAAGAAACAAAACAACTAGCAAAATAGGCTGCCAGTGCA

141 GGTGCCAGAACATTCTATCGAAGGATCTGCGATCGCTCCGGTCCCAGTGGCAGAGCGCACAT

211 CGCCCACAGTCCCCGAGAAGTTGGGGGAGGGTCGGCAATTGAACGGGTGCCTAGAGAAGGTGGCGCG

281 GGTAAACTGGAAAGTGTGTCGTACTGGCTCCGCCTTTCCGAGGGTGGGGAGAACCGTATATA

HindIII (410)

351 AGTCAGTAGTCGCCGTGAACGTTCTTCGCAACGGTTGCCAGAACACAGCTGAAGCTCGAG

421 GGGCTCGCATCTCCTTCACGCCGCCGCCCTACCTGAGGCCGCCATCCACGCCGGTTGAGTCGCGT

491 TCTGCCGCCCTCCGCCGTGGTGCCTCTGAACCTGCGTCCGCCGTAGGTAAGTTAAAGCTCAGGTG

561 AGACCGGGCCTTGTCCGGCGCTCCCTGGAGCCTACCTAGACTCAGCCGCTCTCCACGCTTGCGCTGA

631 CCCTGCTTGCTCAACTCTACGTCTTGTTCTGTTCTGCGCCGTTACAGATCCAAGCTGTGACC

BspHI (725) Eco47III (763)

AgeI (717) **BamHI (757)**

701 **GGCGCCTAC**CTGAGATCACCGGTCACTAGAGCGGCTCCATCATCACCATCAC**GGATCCAGCGCTG**

↑ Met Ser Gl ySer Hi sHi sHi sHi sHi sHi sHi sHi ySer

NheI (779)

NcoI (773)

771 CAGCCATGGGCTAG**CTGGCCAGACATGATAAGATA**CATTGATGAGTTGGACAAACACA

841 AGTAAAAAAATGCTTATTGTGAATTGTGATGCTATTGCTTATTGTAACCATTATAAGCTGCAA

911 TAAACAAGTTAACACAACAATTGCATTCAATTGTTAGGTTCAAGGGGAGGTGGAGGTTTT

EcoRI (1013)

981 TAAAGCAAGTAAACCTCTACAAATGTGGTATGGATTCTAAACAGCATAGCAAAACTTAACTCC

1051 AAATCAAGCCTACTTGAATCCTTCTGAGGGATGAATAAGGCATAGGCATAGGGCTGTTGCCAAT

1121 GTGCATTAGCTGTTGCAGCCTCACCTCTTCATGGAGTTAAAGATATAGTGTATTGCCAAGGTTG

1191 AACTAGCTTCAATTCTTATGTTAAATGCACTGACCTCCACATTCCCTTTAGTAAAATATTCA

1261 GAAATAATTAAATACATCATTGCAATGAAAATAATGTTTTATTAGGCAGAATCCAGATGCTCAAGG

1331 CCCTCATAATATCCCCAGTTAGTTAGGGACTTAGGAAACAAAGGAACCTTAATAGAAATTGGACA

1401 GCAAGAAAGCGAGCTCTAGCTTAGTCTGGTACTTGAGGGGATGAGTCCTCAATGGTTTT

141 Asn Arg Thr Tyr Lys Leu Pro Ile Leu Gl u Gl u Ile Thr Thr Lys

1471 GACCAGCTGCCATTCACTCAATGAGCACAAAGCAGTCAGGAGCATAGTCAGAGATGAGCTCTGCAC

125 Val Leu Lys Gl y Asn Met Gl u Ile Leu Val Phe Cys Asp Pro Al a Tyr Asp Ser Ile Leu Gl u Arg Cys M

1541 ATGCCACAGGGCTGACCACCTGATGGATCTGTCACCTCATCAGAGTAGGGTGCCTGACAGCCACAA

101 Met Gl y Cys Pro Ser Val Val Arg Ile Ser Arg Asp Val Gl u Asp Ser Tyr Pro His Arg Val Al a Val I I

1611 TGGTGTCAAAGTCCTCTGCCGTTGCTCACAGCAGACCCAATGGCAATGGCTTCAGCACAGACAGTGAC

78 Ile Thr Asp Phe Asp Lys Gl n Gl y Asn Ser Val Al a Ser Gl y Ile Al a Ile Al a Gl u Al a Cys Val Thr Val

1681 CCTGCCAATGTAGGCCCTCAATGTGGACAGCAGAGATGATCTCCCAGTCTGGTCTGATGGCCGCCCCG

55 Arg Gl y Ile Tyr Al a Gl u Ile His Val Al a Ser Ile Ile Gl u Gl y Thr Lys Thr Arg Ile Al a Al a Gl y V

1751 ACATGGTGCTTGTTCCTCATAGAGCATGGTATCTCTAGTGGCACCTCACCAGCTCCAGATCCT
 314 al His Lys Asn Asp Glu Tyr Leu Met Thr Ile Lys Glu Thr Ala Val Glu Val Leu Glu Leu Asp Glu
 BspHI (1841)
 1821 GCTGAGAGATGTTGAAGGTCTTCATGATGCCCTCTATAGTGAGTCGTATTATACTATGCCGATATACT

←

841 n Glu n Ser Ile Asn Phe Thr Lys Met
AseI (1899)
 1891 ATGCCGATGATTAATTGTCAAAACAGCGTGGATGGCGTCTCCAGCTTATCTGACGGTTCACTAACGAGC

←

1961 TCTGCTTATATAGACCTCCCACCGTACACGCCTACCGCCATTGCGTCAATGGGGGGAGTTGTTACGA

SpeI (2054)

2031 CATTGGAAAGTCCC GTT GATT TACT AGT CAAAACAAACTCCCATTGACGTCAATGGGTGGAGACTT

2100 GGAAATCCCCGTGAGTCAAACCGCTATCCACGCCATTGATGTA C T G C C A A A C C G C A T C A T G G T A A

2170 TAGCGATGACTAATACGTAGATGTA C T G C C A A G T C C A T A A G G T C A T G T A C T G G G C A T A T G

2240 CCAGCGGGCCATT TACCGT CATTGACGTCAATAGGGGGCGTACTTGGCATATGATA C A C T T G A T G T A C T

2310 GCCAAGTGGCAGTTACGTAAATACTCCACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTA

2380 TGGGAACATACGT CATT ATTGACGTCAATGGCGGGGTCGGTGGCGGT CAGCCAGGCGGGCATT TAC

←

PacI (2473)
SdaI (2465)
 2450 CGTAAGTTATGTAACGCC T G C A G G T T A A T T A A G A A C A T G T G A G C A A A A G G C C A G C A A A A G G C C A G G A A

←

2518 CCGTAAAAAGGCCCGTTGCTGGCGTTTCCATAGGCTCCGCCCTGACGAGCATCACAAAATCGA

2588 CGCTCAAGTCAGAGGTGGC GAAACCCGACAGGACTATAAGATA C C A G G C G T T C C C T G G A A G C T C C

2658 TCGTGCCTCTCTGTTCCGACCCCTGCCGTTACCGGATACCTGTCCGCCCTTCTCCCTCGGAAGCGT

2728 GGCCTTCTCATAGCTACGCTGTAGGTATCTCAGTTGGTAGGTCTCGCTCCAAGCTGGCTGT

ApalI (2797)

2798 GTGCACGAACCCCCCGTT CAGCCGACCGCTGCCCTATCGTA ACT ATCGTCTTGAGTCCAACCCGG

2868 TAAGACACGACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGC GG

2938 TGCTACAGAGTTCTGAAGTGGTGGCTAACTACGGCTACACTAGAAGAACAGTATTGGTATCTGC GCT

3008 CTGCTGAAGCCAGTTACCTCGGAAAAAGAGTTGGTAGCTTGATCCGGCAAACAAACCACCGCTGGTA

3078 GCGGTGGTTTTTGTGCAAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCCTTGAT

←

3148 CTTTCTACGGGTCTGACGCTCAGTGGAACGAAACTCACGTTAAGGGATTTGGTATGGCTAGTTAA

3218 TTAACATTAAATCA

←

PacI (3213)