

pFUSE-mIgG2A-Fc2

Plasmid containing a mouse IgG2A Fc region

Catalog # pfuse-mg2afc2

For research use only

Version 20K05-MM

PRODUCT INFORMATION

Content:

- 20 µg of pFUSE-mIgG2A-Fc2 (IL2ss) 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 stored at -20°C and is stable 3 months.
- Resuspended DNA should be stored at -20°C and is stable up to 1 year.
- Store Zeocin™ at 4 °C or at -20 °C. The expiry date is specified on the product label.
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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

pFUSE-Fc is a family of plasmid developed to facilitate the construction of Fc-fusion proteins by fusing the effector region of a protein to the Fc region of an immunoglobulin G (IgG).

pFUSE-Fc plasmids yield high levels of Fc-fusion proteins. The level of expression is usually in the µg/mL range. They can be transfected in a variety of mammalian cells, including myeloma cell lines, CHO cells, monkey COS cells and human embryonic kidney (HEK)293 cells, cells that are commonly used in protein purification systems.

pFUSE-Fc2 (IL2ss) plasmids allow the secretion of Fc-Fusion proteins. They contain the IL2 signal sequence (IL2ss) for the generation of Fc-Fusion proteins derived from proteins that are not naturally secreted. As Fc-Fusion proteins are secreted, they can be easily detected in the supernatant of pFUSE-Fc-transfected cells by SDS-PAGE. Furthermore, functional domains can be identified by immunoblotting and ligand blotting.

Fc-Fusion proteins can be easily purified by single-step protein A or protein G affinity chromatography.

InvivoGen provides pFUSE-Fc vectors featuring Fc regions from different species and isotypes. Three murine isotypes are available: IgG1, IgG2a and IgG3. The Fc region mediates effector functions, such as antibody-dependent cellular cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC). In ADCC, the Fc region of an antibody binds to Fc receptors (FcγRs) on the surface of immune effector cells such as natural killers and macrophages, leading to the phagocytosis or lysis of the targeted cells. In CDC, the antibodies kill the targeted cells by triggering the complement cascade at the cell surface. IgG isoforms exert different levels of effector functions increasing in the order of mIgG1<mIgG3<mIgG2a.

PLASMID FEATURES

- **mIgG2A Fc (mouse):** The Fc region comprises the CH2 and CH3 domains of the IgG heavy chain and the hinge region. The hinge serves as a flexible spacer between the two parts of the Fc-fusion protein, allowing each part of the molecule to function independently. The Fc region of mouse IgG2A mediates high ADCC and CDC.
- **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.
- **IL2 ss:** The IL2 signal sequence contains 20 amino acids and share common characteristics with signal peptides of other secretory proteins. The intracellular cleavage of the IL2 signal peptide occurs after Ser20 and leads to the secretion of the antigenic protein.
- **MCS:** The multiple cloning site contains several restriction sites that are compatible with many other enzymes, thus facilitating cloning.
- **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.
- **CMV enh / hFerL prom:** This composite promoter combines the human cytomegalovirus immediate-early gene 1 enhancer and the core promoter of the human ferritin light chain gene. This ubiquitous promoter drives the expression of the Zeocin™-resistance gene in mammalian cells.
- **EM2KC** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*. EM2KC is located within an intron and is spliced out in mammalian cells.
- **Zeo:** Resistance to Zeocin™ is conferred by the *Sh ble* gene from *Streptoalloteichus hindustanus*. The same resistance gene confers selection in both mammalian cells and *E. coli*.
- **βGlo pAn:** The human beta-globin 3'UTR and polyadenylation sequence allows efficient arrest of the transgene transcription⁴.

References:

1. Kim DW *et al.* 1990. Use of the human elongation factor 1 alpha promoter as a versatile and efficient expression system. 91(2):217-23.
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. 8(1):466-72.
3. Carswell S. & Alwine JC. 1989. Efficiency of utilization of the simian virus 40 late polyadenylation site: effects of upstream sequences. Mol Cell Biol. 9(10):4248-58.
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.

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

Zeocin™ usage

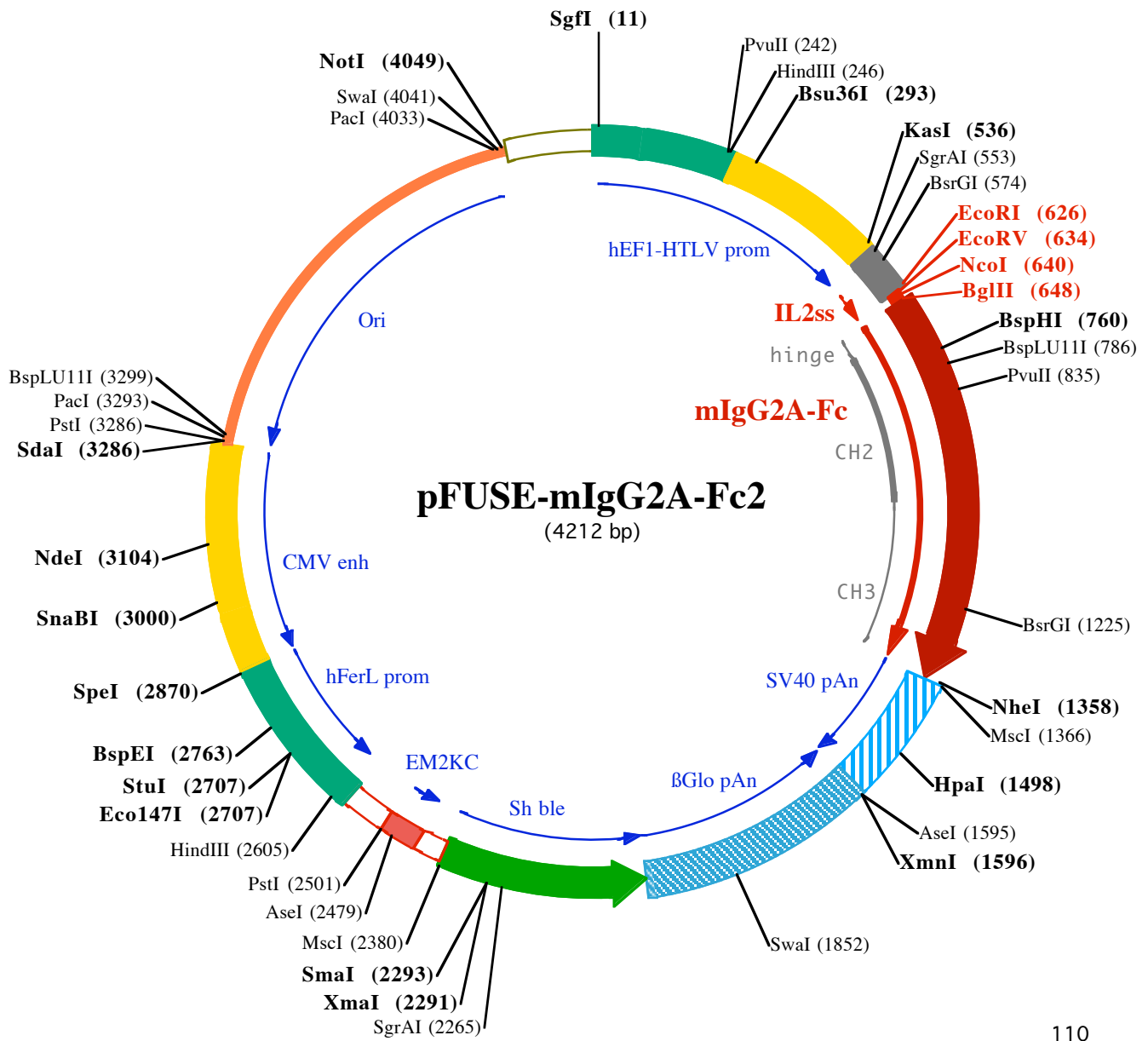
This antibiotic can be used for *E. coli* at 25 µg/ml in liquid or solid media and at 50-200 µg/ml to select Zeocin™-resistant mammalian cells.

RELATED PRODUCTS

Product	Catalog Code
Zeocin™	ant-zn-1

TECHNICAL SUPPORT

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SgfI (11)
1 GGATCTGCGATCGCTCCGGTCCCGTCAGTGGGAGGCGCACATCGCCACAGTCCCGAGAAGTTGGGGGAGGGTTCGCAATTGAACGGGTGCCTA
101 GAGAAAGTGGCGGGGTAACCTGGGAAAGTGTCTGTACTGGCTCCGCTTTTCCCGAGGGTGGGGGAGAACCCTATATAAGTGCAGTAGTCGCC

HindIII (246) **PvuII (242)** **Bsu36I (293)**
201 GTGAACGTTCTTTTCGCAACGGGTTTCCGCCAGAACACAGCTGAAGCTTCGAGGGGCTCGCATCTCTCTTACGCGCCCGCCCTACCTAGAGGCC
301 GCCATCCACGCGGGTTGAGTGCCTCTCCGCTCCCGCTGTGGTGCCTCTGAACTGCGTCCGCGTCTAGGTAAGTTAAAGCTCAGGTCGAGACC
401 GGGCCTTTGTCCGGCTCCCTTGGAGCTACCTAGACTCAGCGGCTCTCCACGCTTTGCTGACCCTGCTTCTCAACTCTACGTTCTTTGTTCTGTT

KasI (536) **SgrAI (553)** **BsrGI (574)**
501 TCTGTTCTGCGCGTTACAGATCCAAGCTGTGACCGGGCTACTCTGAGATCACCGCGAAGGAGGCCACCATGTACAGGATGCAACTCTGCTTTGCA
1MetTyrArgMetGlnLeuLeuSer CysI

EcoRV (634) **BglII (648)**
EcoRI (626) **NcoI (640)**
601 TTGCACTAAGTCTTGCATTGTACGAATTCGATATCGGCATGGTTAGATCTCCAGAGGGCCCAATCAAGCCCTGTCTCCATGCAATGCCAGC
10PLeuAlaLeuSerLeuAlaLeuValThrAsnSer 1ProArgGlyProThrIleLysProCysProProCysLysCysProAl

BspHI (760) **BspLU11I (786)**
701 ACCTAACCTTTGGGTGACCATCCGCTTTCATCTTCCCTCAAAGATCAAGGATGTACTCATGATCTCCCTGAGCCCATAGTCACATGTGTGGTGGTG
16PAsnLeuLeuGlyProSerValPheIlePheProProLysIleLysAspValLeuMetIleSerLeuSerProIleValThrCysValValVal

PvuII (835)
801 GATGTGAGCGAGGATGACCCAGATGTCCAGATCAGCTGGTTTGTGAACAACGTGGAAGTACACAGCTCAGACACAAACCCATAGAGAGGATTACAACA
50PAspValSerGluAspAspProAspValGlnIleSerTrpPheValAsnAsnValGluValHisThrAlaGlnThrGlnThrHisArgGluAspTyrAsnS

901 GTACTCTCCGGTGGTCAAGTCCCTCCATCCAGCACCAGACTGGATGAGTGGCAAGGAGTTCAAATGCAAGTCAACAACAAAGCTCCAGCGCC
83PThrLeuArgValValSerAlaLeuProIleGlnHisGlnAspTrpMetSerGlyLysGluPheLysCysLysValAsnAsnLysAspLeuProAlaPr

1001 CATCGAGAGAACCATCTCAAACCCAAAGGGTCAAGAGCTCCACAGGTATATGCTTGCCTCCACCAGAAGAAGATGACTAAGAAACAGTCACT
116PAlaIleGluArgThrIleSerLysProLysGlySerValArgAlaProGlnValTyrValLeuProProProGluGluGluMetThrLysLysGlnValThr

1101 CTGACCTGCATGGTCAAGACTTTCATGCTGAAGACATTTACGTGGAGTGGACCAACACGGGAAAACAGAGCTAAACTACAAGAACTGAACAGTCC
150PLeuThrCysMetValThrAspPheMetProGluAspIleTyrValGluTrpThrAsnAsnGlyLysThrGluLeuAsnTyrLysAsnThrGluProValL

BsrGI (1225)
1201 TGGACTCTGATGGTCTTACTTCTACAGCAAGCTGAGAGTGGAAAAGAACTGGGTGGAAAAGAACTACTCTCTTCAAGTGGTCCACGAGGG
183PeuAspSerAspGlySerTyrPheMetTyrSerLysLeuArgValGluLysLysAsnTrpValGluArgAsnSerTyrSerCysSerValValHisGluGlu

MscI (1366) **NheI (1358)**
1301 TCTGCACAATCACACAGACTAAGAGCTTCTCCCGACTCCGGTAAATGAGCTCAGCTAGCTGGCCAGACATGATAAGATACATTGATGAGTTGGAC
216PLeuHisAsnHisHisThrThrLysSerPheSerArgThrProGlyLys

HpaI (1498)
1401 AAACCACAAC TAGAATGCAGTGAATAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAA

AseI (1595) **XmnI (1596)**
1501 CAACAACAATTGCATTCAATTTATGTTTCAGGTTCAAGGGGAGGTGTGGGAGGTTTTTAAAGCAAGTAAACCTCTACAAATGTGGTATGGAATTAATT
1601 CTAATAACAGCATAGCAAACTTTAACCTCCAAATCAAGCCTCTACTTGAATCTTTTCTGAGGGATGAATAAGGCATAGGCATCAGGGCTGTTGCCA
1701 ATGTGCATTAGCTGTTTGCAGCCTCACCTTCTTTCATGGAGTTAAGATATAGTGTATTTTCCAAAGGTTTGAAC TAGCTCTTCATTTCTTATGTTTTA

SwaI (1852)
1801 AATGCACTGACCTCCCACTTCCCTTTTAGTAAATATTCAGAATAAATTAATATACATCATTGCAATGAAATAAATGTTTTTATTAGGCAGAATCC
1901 AGATGCTCAAGGCCCTCATAATATCCCGAGTTTAGTAGTTGACTTAGGGAACAAAGAACCTTTAATGAAATTTGGACAGCAAGAAAGCGAGCTCT

2001 AGCTTATCCTCAGTCTGCTCCTGCCACAAAGTGCACGAGTTGCCGGCGGGTCCGCGAAGTCCCGCCCCACGGCTGCTCGCGATCTCG
125PAspGlnGluGluAlaValPheHisValCysAsnGlyAlaProAspArgLeuAlaPheGluArgGlyTrpProGlnGluGlyIleGluT
2101 GTCATGGCGCGCCGAGGCGTCCGGAAGTTCGTGGACACGACCTCCGACCTCGCGGTACAGCTGTCAGGCGCGCACCCACACCCAGGCCAGGG
94PThrMetAlaProGlySerAlaAspArgPheAsnThrSerValValGluSerTrpGluAlaTyrLeuGluAspLeuGlyArgValTrpValTrpAlaLeuTh

XmaI (2291) **SmaI (2293)**
2201 TGTTGTCCGGCACCTGCTGACCGCTGATGAACAGGGTCAAGTCTGCTCCAGGACCAACCGCGAAGTCTCCAGGAGTCCCGGGAGAA
61PAsnAspProValValGlnAspGlnValAlaSerIlePheLeuThrValAspAspArgValValGlyAlaPheAspAspGluValPheAspArgSerPhe

MscI (2380)
2301 CCCGAGCGGTCGGTCCAGAACTCGACCGCTCCGCGCAGCTCGCGCGGTTGAGCACCAGAACGGCACTGGTCAACTTGGCCATGATGGCTCTCctgtc
28PLeuLysArgAspThrTrpPheGluValAlaGlyAlaValAspArgAlaThrLeuValProValAlaSerThrLeuLysAlaMet

AseI (2479) **PstI (2501)**
2401 aggagaggaaagagaagaaggttagtacaattgCTATAGTGTGTTGATTATACATATGAGATATACTATGCAATGATAAATGTGCAAACTAGGGCTGC
2501 AgggttcatatgcccacttttctgactgccccactctctgccacccttccaggcatagacagtcagtgacttacCAAACCTCACAGGAGGGAGAAG
HindIII (2605)
2601 GCAGAAGCTTGAGACAGACCCGCGGACCGCCGAAGTCCGAGGGGACGTGGCTAGGGCGGCTTTTATGGTGGCGCGCCCTCGGAGGCAGGGCGCTC

2701 **StuI (2707)**
Eco147I (2707) GGGGAGGCCTAGCGCCAATCTGCGGTGGCAGGAGCGGGGCCGAAGCCGTGCTGACCAATCCGGAGCACATAGGAGTCTCAGCCCCCGCCCAAG
BspEI (2763)

2801 CAAGGGGAAGTCACGCGCTGTAGCGCCAGCGTGTGTGAAATGGGGCTTGGGGGGTGGGGCCCTGACTAGTCAAAACAAACTCCCAATTGACGTCAA
SpeI (2870)

2901 TGGGGTGGAGACTTGAAATCCCGTGAGTCAAACCGCTATCCACGCCATTGATGTACTGCCAAAACCGCATCATCATGGTAATAGCGATGACTAATAC
SnaBI (3000)

3001 GTAGATGTACTGCCAAGTAGGAAAGTCCATAAGGTCATGTACTGGGCATAATGCCAGGCGGGCCATTTACCGTCATTGACGTCAATAGGGGGCGTACTT

3101 **NdeI (3104)**
GGCATATGATACACTTGATGTACTGCCAAGTGGGCAGTTTACCGTAAATACTCCACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAA

3201 CATACTGTCATTATTGACGTCAATGGGCGGGGTCGTTGGGGCGTACGCCAGGCGGGCCATTACCCTAAGTTATGTAACGCCGCTGACAGGTTAATTAAGAAC
PacI (3293)
PstI (3286)
SdaI (3286) BspLU11I (3299)

3301 ATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAAAAGCCGCGTTGCTGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATC

3401 GACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGCGTTTTCCCTGGAAGCTCCCTCGTGGCTCTCCTGTTCCGACCTGCCC

3501 GCTTACCGGATACCTGTCCGCTTTCTCCCTTGGGAAGCGTGGCGTTTTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTGCTTCCGCTCC

3601 AAGCTGGGCTGTGTGCACGAACCCCGTTCCAGCCGACCGCTGCGCTTATCCGGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATCGC

3701 CACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGCGGTGCTACAGAGTTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAG

3801 AACAGTATTTGGTATCTGCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAACCCAGCTGGTAGCGGTGGT

3901 TTTTTTGGTTCGCAAGCAGCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAAACGAAAAC

4001 CACGTTAAGGGATTTTGGTCAAGGCTAGTTAATTAACATTTAAATCAGCGGCCGCAATAAAATATCTTTATTTTTCATTACATCTGTGTGTTGGTTTTTTG
PacI (4033) **Swal (4041)** **NotI (4049)**

4101 TGTGAATCGTAACTAACATACGCTCTCCATCAAAACAAAACGAAACAAAACAACTAGCAAAATAGGCTGTCCCAGTGCAAGTGCAGGTGCCAGAACAT

4201 TTCTCTATCGAA