

# pFUSE-CHIg-mD

Plasmid featuring the constant region of the mouse IgD heavy chain

Catalog code: pfuse-mchd

## For research use only

Version 24J28-MM-v37

## PRODUCT INFORMATION

### Content:

- 20 µg of pFUSE-CHIg-mD 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.
- 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.

### Quality control:

- Plasmid construct has been confirmed by restriction analysis and sequencing.
- Plasmid DNA was purified by ion exchange chromatography.

### Materials required for antibody generation & isotype switching

- pFUSE2-CL Ig plasmid that features the constant region of the kappa or lambda light chains. pFUSE2-CL Ig plasmids are selectable with blasticidin (sold separately, see RELATED PRODUCTS).
- pFUSE-CHIg plasmid for the constant region of the heavy chain, this plasmid is selectable with Zeocin®.

## GENERAL PRODUCT USE

pFUSE-CL Ig and pFUSE-CHIg plasmids are designed to change a monoclonal antibody from one isotype to another, therefore, enabling the generation of antibodies with the same antigen affinity but with different effector functions (increased or reduced ADCC and CDC). Furthermore, they can be used to produce entire antibodies from Fab or scFv fragments.

pFUSE-CHIg and pFUSE2-CL Ig express the constant regions of the heavy (CH) and light (CL) chains, respectively. They contain a multiple cloning site (MCS) upstream of these constant regions to enable the cloning of the variable (VH and VL) regions of a given antibody. Transfection of mammalian cell lines with the recombinant pFUSE-CHIg and pFUSE2-CL Ig pair allows to generate an antibody that can be purified from the supernatant using the appropriate affinity chromatography.

### Features of pFUSE-CHIg and pFUSE2-CL Ig plasmids

- **hEF1-HTLV prom**: is a composite promoter comprising the Elongation Factor-1α (EF-1 $\alpha$ ) core promoter<sup>1</sup> and the R segment and part of the U5 sequence (R-U5') of the Human T-Cell Leukemia Virus (HTLV) Type 1 Long Terminal Repeat<sup>2</sup>. The EF-1 $\alpha$  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 $\alpha$  core promoter to enhance stability of RNA.
- **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 mRNA<sup>3</sup>.
- **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.
- **βGlo pAn**: The human beta-globin 3'UTR and polyadenylation sequence allows efficient arrest of the transgene transcription<sup>4</sup>.

### pFUSE-CHIg-mD specific features

- **Mouse IGHD (IgD heavy chain constant region)**: When cloning your heavy chain variable region of choice in the MCS, care must be taken to insert the gene in-frame and to preserve the integrity of the heavy chain constant region.
- **Zeo**: Resistance to Zeocin® is conferred by the *Sh ble* gene from *Streptallotheichus hindustanus*. The same resistance gene confers selection in both mammalian cells and *E. coli*.

### 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

InvivoGen USA (Toll-Free): 888-457-5873

InvivoGen USA (International): +1 (858) 457-5873

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

InvivoGen Asia: +852 3622-3480

E-mail: info@invivogen.com

## PROTOCOL

### Obtaining VH and VL sequences

The antibody sequence can be obtained by phage display or from an antibody producing hybridoma. To obtain the cDNA sequence of the VH and VL regions from an antibody producing hybridoma, total RNA or mRNA is extracted and reverse transcribed to cDNA. PCR is performed with 5' degenerate primers to anneal to the unknown VH and VL regions and the 3' primers designed to anneal to the “known” CH and CL regions. Alternatively 5' RACE can be used. The resulting amplicons must be sequenced.

### Plasmid resuspension

Quickly spin the tube containing the lyophilized plasmid to pellet the DNA. To obtain a plasmid solution at 1  $\mu$ g/ $\mu$ l, resuspend the DNA in 20  $\mu$ l of sterile H<sub>2</sub>O. Store resuspended plasmid at -20°C.

### Cloning into pFUSE-CHIg and pFUSE2-CL Ig

Once the VH and VL sequence are known, inserts for cloning into the plasmids can be generated. In pFUSE-CHIg-mD, the constant region of the mouse IgD heavy chain is preceded by a multiple cloning site containing six restriction sites: AgeI, EcoRI, EcoRV, XhoI, NheI, and Eco47III. The first five restriction sites can be used for insertion of the 5' end of the variable region including the native signal sequence. If the immunoglobulin signal sequence is unknown, pFUSEss plasmids containing a signal sequence should be used. In pFUSE-CHIg-mD, Eco47III must be used for insertion of the 3' end of the variable region. Eco47III must be reconstituted to maintain the integrity of the constant region. Therefore we recommend to introduce by PCR an Eco47III site at the 3' end of the variable region in frame with the constant region.

When generating the insert for VL, a BstAPI (pFUSE2-CL Ig-mk; mouse kappa), or AvrII (pFUSE2-CL Ig-ml1 / pFUSE2-CL Ig-ml2; mouse lambda) site must be introduced at the 3' end. There is a choice of restriction sites at the 5' end.

Note: The 5'end of the variable region should encompass the native ATG initiation codon and the region immediately after which corresponds to the signal sequence. For proper initiation of translation, make sure that your insert contains a Kozak translation initiation sequence upstream of the ATG initiation codon such as (G/A)NNATG.

### Choice of strategies for the transfection

Transfect cells using a transfection agent, such as LyoVec™, with the plasmid coding for light chain and select the best clone. Following selection of the best clone, the plasmid coding for the heavy chain clone can be transfected into this clone.

OR

A cotransfection can be performed with the plasmid coding for the light chain and the plasmid coding for the heavy chain. Since the pFUSE2-CL Ig and pFUSE-CHIg plasmids share the same plasmid backbone, the appropriate heavy chain to light chain ratio can be easily determined by varying the quantities of pFUSE2-CL Ig and pFUSE-CHIg plasmids. We recommend using a ratio of 3:2 of pFUSE2-CL Ig:pFUSE-CHIg plasmids. pFUSE2-CL Ig plasmids feature the constant region of a kappa or lambda light chain. pFUSE2-CL Ig plasmids are selectable with blasticidin. pFUSE-CHIg plasmids are selectable with Zeocin®.

### TECHNICAL SUPPORT

InvivoGen USA (Toll-Free): 888-457-5873

InvivoGen USA (International): +1 (858) 457-5873

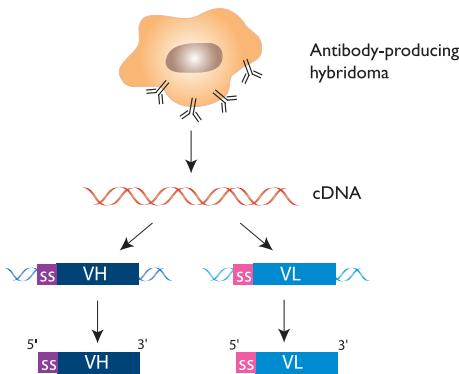
InvivoGen Europe: +33 (0) 5-62-71-69-39

InvivoGen Asia: +852 3622-3480

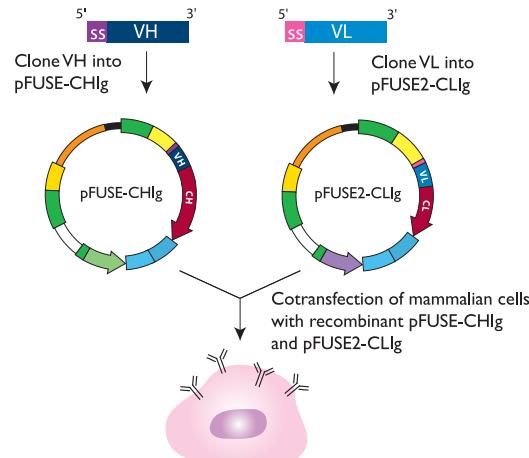
E-mail: info@invivogen.com

### Antibody generation using pFUSE-CHIg & pFUSE-CL Ig

#### I- Obtention of VH and VL sequences



#### 2- Cloning into pFUSE-CHIg and pFUSE2-CL Ig

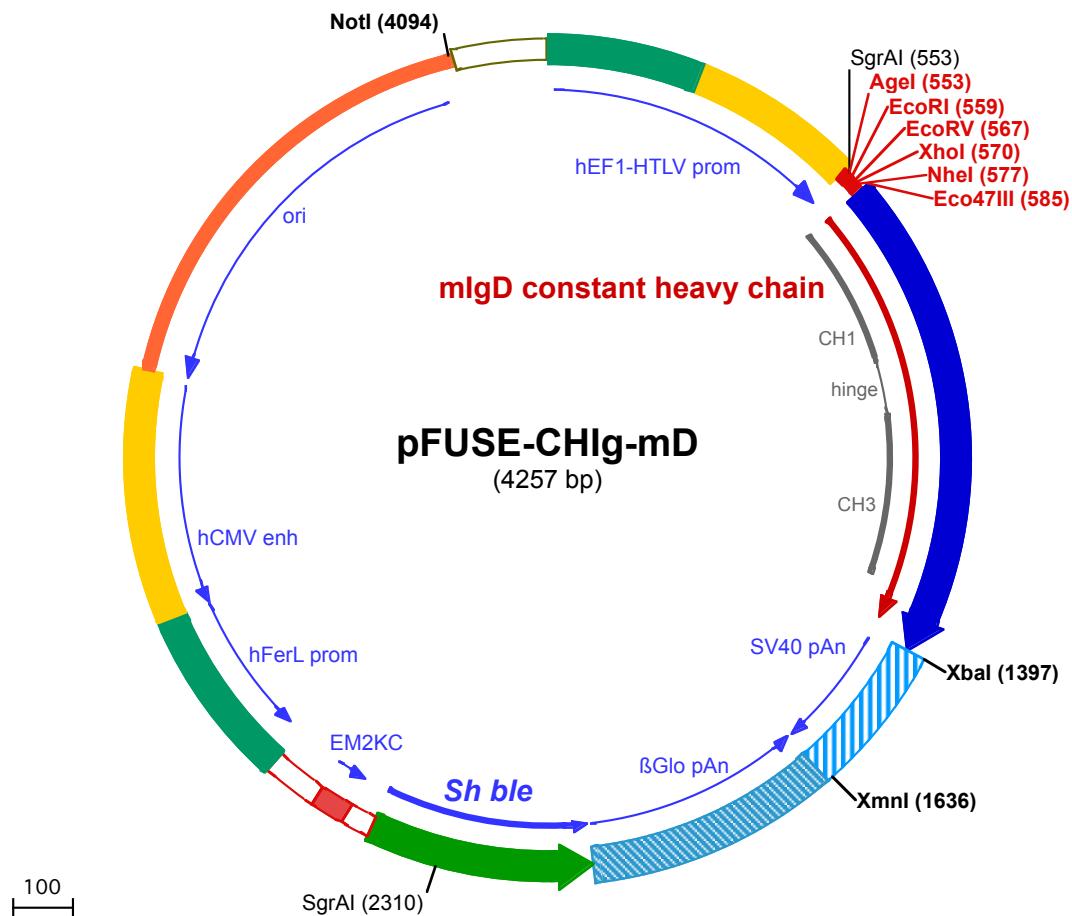


To check for production of your antibody after transfection, you may take an aliquot of growth medium and perform SDS-PAGE, protein-specific ELISA, or the bioactivity assay of choice to determine that your cells are producing your antibody of interest.

The resulting IgD antibody that can be purified from the supernatant using the appropriate affinity chromatography.

## RELATED PRODUCTS

Product	Catalog Code
pFUSE2-CL Ig-mk	pfuse2-mcl1k
pFUSE2-CL Ig-ml1	pfuse2-mcl1l
pFUSE2-CL Ig-ml2	pfuse2-mcl1l2
LyoVec™	lyec-12
Protein L / Agarose	gel-protl-2
Zeocin®	ant-zn-1



1 GGATCTGCATCGCTCCGGTCCCCTCAGGGCAGAGCGCACATGCCACAGTCCCAGAAGTTGGGGGAGGGTCGCAATTGAACGGGTGCTAG  
 102 AGAAGGTGGCGCGGGTAAACTGGAAAGTGATGTCGTACTGGCTCCGCTTTCCGAGGGTGGGGAGAACCGTATAAGTCAGTAGTCGCCGT  
 203 GAACGTTCTTTCGCAACGGTTGCCAGAACACAGCTGAAGCTCGAGGGGCTCGATCTCCTCACCGCCGCCCTACCTGAGGCCGCC  
 304 ATCCACGCCGGTTGAGTCGCGTTCTGCCGCCCTGGTGCCTCTGAACCTCGTCCGCCGTAGGTAAGTTAAAGCTCAGGTCAGACCGGGC  
 405 CTTTGTCCGGCTCCCTGGAGCCTACCTAGACTCAGCCGCTCTCCACGCTTGCTGACCCCTGCTCAACTTACGCTTTGTTCTGTTCTG  
  
 EcoRI (559)  
 AgeI (553) Xhol (570) Eco47III (585)  
 SgrAI (553) EcoRV (567) NheI (577)  
 506 TCTGCGCCGTTACAGATCCAAGCTGTGACCGGCGCTACCTGAGATACCGGTGAATTGATATCTGAGTCTAGCAGCCTGGTATAAAAAGGAACCT  
 1 ▶ G D K K E P  
  
 607 GACATGTTCTCTCTCAGAGTGCAGGCCCCAGAGGAAATGAAAAGATAAACCTGGCTGTTAGTAATTGAGTCAGCCACTGAAAATCAGCTGGG  
 7 ▶ D M F L L S E C K A P E E N E K I N L G C L V I G S Q P L K I S W E  
  
 708 GCCAAAGAAGTCAAGTATAGTTGAAACATGCTTCCCTCTGAAATGAGAAATGGCAATTATAACATGGCTCCAGGTCACTGTGCTGGCCTCAGAACTG  
 40 ▶ P K K S S I V E H V F P S E M R N G N Y T M V L Q V T V L A S E L  
  
 809 ACCTCAACCACACTTGACCATAATAACCCAAAAGGAAAGAAAAACCTTCAAGTTCCCTGAGTCATGGGATTCCAGTCTAAGAGAGTCACCTCA  
 74 ▶ N L N H T C T I N K P K R K E K P F K F P E S W D S Q S S K R V T P  
  
 910 ACTCTCAAGCAAAGAATCACTCCACAGAACGCTATTACCAACAAAGGACATAGAAGGGGCATGGCACCCAGCAACCTCACTGTGACAT  
 108 ▶ T L Q A K N H S T E A T K A I T T K K D I E G A M A P S N L T V N I  
  
 1011 CCTGACCACATCCACCCATCTGAGATGTCATCTGGCTCTGTGAGATATCTGGCTCTCCCGAAAATATCCACCTCATGGCTGAGTGTCCACA  
 141 ▶ L T T S T H P E M S S W L L C E V S G F F P E N I H L M W L S V H  
  
 1112 GTAAAATGAAGTCTACAAACTTTGTCAGTGCACCCACCGCCAGCCTGGGGCACATTCCAGACCTGGAGTGTCTGAGACTACCACTGCTGAGC  
 175 ▶ S K M K S T N F V T A N P T A Q P G G T F Q T W S V L R L P V A L S  
  
 1213 TCATCACTTGACACTTACACATGTGTTGAAACATGAGCCCTCAAAGACAAAGCTTAATGCCAGCAAGAGCCTAGCAATTAGTGGATGCTACCACTCT  
 209 ▶ S S L D T Y T C V V E H E A S K T K L N A S K S L A I S G C Y H L L  
  
 XbaI (1397)  
 1314 GCCTGAGTCAGACGGTCCTCCAGGAGACCTGATGGCTCTGCCCTGCCAGACCTTCTAGGCTGAATGGTCATCATGCTCTAGACCTAGCTGGCA  
 242 ▶ P E S D G P S R R P D G P A L A •  
 1415 GACATGATAAGATACATTGATGAGTTGGACAAACCAACTAGAATGCACTGAAATGCTTATTGTGAAATTGATGCTATTGCTTATTG  
  
 1516 AACCAATTATAAGCTGAATAAACAGTTAACACAACAAATTGATTCTATTTATGTTCAGGTTAGGGGAGGTGGAGGTTAAAGCAAGTAA  
  
 XmnI (1636)  
 1617 ACCTCTACAAATGTGGTATGGAATTAACTCTAAACACAGCATAGCAAAACTTAAACCTCAAATCAAGCTCTACTTGAATCCTTCTGAGGGATGAAT  
 1718 AAGGCATAGGCATCAGGGCTTGCCTGCAATTGTCATTAGCTGTTGCAGCCTCACCTCTTCTGAGTTAAAGATATAGTGTATTCCAAGGTTGA  
 1819 ACTAGCTCTCATTCTTATGTTAAATGCACTGACCTCCACATTCCCTTTAGTAAATATTAGAATAATTAAACATCATTGAATGAAA  
 1920 TAAATGTTTTATTAGGCAGAATCCAGATGCTCAAGGCCCTCATAATATCCCCAGTTAGTTGACTTAGGAACAAAGAACCTTAATAGAAA  
  
 2021 TTGGACAGCAAGAAAGCGAGCTCTAGCTTACCTCAGTCCTGCTCTGCCACAAAGTCACCGCAGTGGCCGGCGGGTCGCGCAGGGCAACTCCCC  
 125 ▶ • D Q E E A V F H V C N G A P D R L A F E R  
 2122 CCCACAGGCTGCTCGCGATCTGGTCATGGCCGGCCGGAGGCCTCCGGAAAGTCGAGACACGACCTCCGACCACTGGCGTACAGCTCGTCAGGGC  
 102 ▶ G W P Q E G I E T M A P G S A D R F N T S V V E S W E A Y L E D L G  
 SgrAI (2310)  
 2223 GCGCACCCACACCCAGGCCAGGGTGTGTCGGCACACCTGGCTGGACCGCGCTGATGAACAGGGTCACGTCGCTCCGGACACACCGCGAAGTCG  
 69 ▶ R V W V W A L T N D P V V Q D Q V A S I F L T V D D R V V G A F D D  
 2324 CCTCCACGAAGTCCCGGGAGAACCCGAGCCGGTCGGTCAGAACCTCGACCGCTCCGGCAGCTCGCGCGCGTGGAGCACCAGGGCACTGGTCAACTTG  
 35 ▶ E V F D R S F G L R D T W F E V A G A V D R A T L V P V A S T L K  
 2425 GCCATGATGGCTCCTCctgtcaggagagaaagagaagggtagtacaatttgCTATAGTGAGTTGATTACTATGCAGATATACTATGCCAATGATT  
 1 ▶ A M  
 2526 AATTGTCAAACCTAGGGCTGCAgggtcatagtgcactttccctgcactgccccatctcccccaccccccaggcatagacagtcagtgcattacCA  
  
 2627 AACTCACAGGAGGGAGAACGGAGAACGCTTGAGACAGACCCGGGACGCCGAACCTGCGAGGGACGTGGCTAGGGCGCTCTTATGGTGCAGGGC  
 ←

2728 CTCGGAGGCAGGGCCTCGGGAGGCCATCGGCCAATCTGGGTGCCAGGAGGCGGGCGAAGGCCGCTGCCATCGAACATCCGAGCACATAGGAGTCTC  
2829 AGCCCCCGCCCCAAAGCAAGGGAAAGTCACGCCCTGTAGCGCCAGCGTGTGAATGGGGCTTGGGGGTTGGGCCCTGACTAGTCAAAACAAA  
2930 CTCCCATTGACGTCAATGGGTGGAGACTTGGAAATCCCCTGAGTCAAACCGCTATCCACGCCATTGATGTAUTGCCAAACCGCATCATGGTAAT  
3031 AGCGATGACTAATACGTAGATGTACTGCCAAGTAGGAAAGTCCCATAAGGTATGTAUTGGCATAATGCCAGGGCCATTACCGTATTGACGTCAA  
3132 TAGGGGGCGTACTTGGCATATGATACACTTGATGTAUTGCCAAGTGGCAGTTACCGTAAATACTCCACCCATTGACGTCAATGGAAAGTCCCTATTGGC  
3233 GTTACTATGGGAACATACGTCAATTGACGTCAATGGCGGGGTCGTTGGCGGTAGCCAGGGCCATTACCGTAAGTTATGTAACGCCGCAGG  
3334 TTAATTAAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAGGCCCGTTGCTGGCGTTTTCCATAGGCTCCGCCCTGACGAGA  
3435 TCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAGATACCGAGCGTTCCCTGGAAAGCTCCCTGTGCGCTCTCTGTC  
3536 CGACCCCTGCCGCTTACCGGATACCTGTCCGCCCTTCCCTCGGAAGCGTGGCGCTTCTCATAGCTCACGCTGTAGGTATCTCAGTCGGTAGGTC  
3637 GTTCGCTCCAAGCTGGCTGTGCACGAACCCCCGTTCAAGCCGACCCGCTGCGCCTTACCGTAACTATCGTCTGAGTCAACCCGTAAGACACGA  
3738 CTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAAGCGGTGCTACAGAGTTCTGAAGTGGTGGCTAACTACGGCTACA  
3839 CTAGAAGAACAGTATTGGTATCTGCCTGCTGAAGCCAGTTACCTTCGGAAAAAGAGTTGGTAGCTCTGATCCGCAAACAAACCACCGCTGGTAGC  
3940 GGTGGTTTTTGTGCAAGCAGATTACGCGCAGAAAAAAAGATCTAAGAAGATCTTGTATTTCTACGGGCTGACGCTAGTGGAAC  
4041 AAACTACGTTAAGGATTTGGTATGGTAGTTAACATTAACATTAAATCAGCGGCCGAATAAAATATCTTATTTCATTACATCTGTGTTGGTT  
4142 TTTGTGTGAATCGTAACATACGCTCTCCATAAAACAAAACGAAACAAAACAAACTAGCAAAATAGGCTCCAGTGCAAGTGCAGGTGCCAGAA  
4243 CATTCTCTATCGAA

**NotI (4094)**