

CAUTION

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pFUSEN-Lucia-mG2AFc

Plasmid designed for Lucia::Fc fusion to the N-terminus of a protein of interest

Catalog # pfcn-lcmg2a

For research use only

Version 20K09-MM-v36

PRODUCT INFORMATION

Content:

- 20 μ g of pFU SEN-Lucia-mG2AFc 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.

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

pFUSEN-Fc / pFUSEN-Lucia-Fc is a family of plasmids developed to facilitate the construction of Fc-fusion proteins where the immunoglobulin G (IgG) Fc-domain is fused to the N-terminus of the protein of interest.

pFUSEN-Fc / pFUSEN-Lucia-Fc plasmids yield high levels of Fc-fusion proteins. The level of expression is usually in the μ g/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.

pFUSEN-Lucia-mG2AFc plasmid allows the production of Lucia-Fc fusion proteins. This plasmid can be used to make recombinant Lucia-Fc fusion proteins or can be used as a transfection control in experiments with other pFUSEN-Fc constructs. Quantification of Lucia-Fc expression can be determined utilizing InvivoGen's QUANTI-Luc™ (rep-qlc1 or rep-qlc2).

A choice of cloning sites is provided to allow flexibility in the design of the fusion linker: either use pFUSEN linker, or bring forth your own linker with the protein of interest.

InvivoGen provides pFUSEN-Lucia-Fc vectors featuring Fc regions from different species and isotypes. In mouse, the IgG2a isotype is available. In humans, three options are available: IgG1, IgG1e2, or IgG2. The Fc region mediates effector functions, such as antibody-dependent cellular cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC). IgG isoforms exert different levels of effector functions. The engineered IgG1e2 contains mutations in the FcRn binding sites leading to higher FcRn binding affinity and reduced pH dependence.

PLASMID FEATURES

- **Lucia luciferase** is a secreted coelenterazine-utilizing luciferase reporter protein with advantageous characteristics when associated with Fc-fusion proteins. It possesses superior carrier ability for excellent secretion of the chimeric protein. It provides a simple means to screen for recombinant clones and it minimally affects the activity of the protein of interest.

- **Mouse IgG2a-Fc** : The Fc region comprises the CH2 and CH3 domains of the IgG2a heavy chain, with the hinge region. The last amino acid (lysine) of the Fc region has been replaced by an alanine for better fusion result. The Fc region of mouse IgG2a mediates high ADCC and CDC.

- **hEF1-HTLV prom** is a composite promoter comprising the Elongation Factor-1 α (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.

- **Cloning sites & fusion linker:** The protein of interest can be cloned either as a BamHI—NheI fragment, or as a BsiWI—NheI fragment. With BamHI cloning, the protein of interest will be separated from the Fc-domain by a flexible linker (Gly4Ser dimer).

With BsiWI cloning, the flexible linker will not be retained, allowing for a different fusion design. The provided cloning sites 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*.

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

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

TECHNICAL SUPPORT

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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.

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

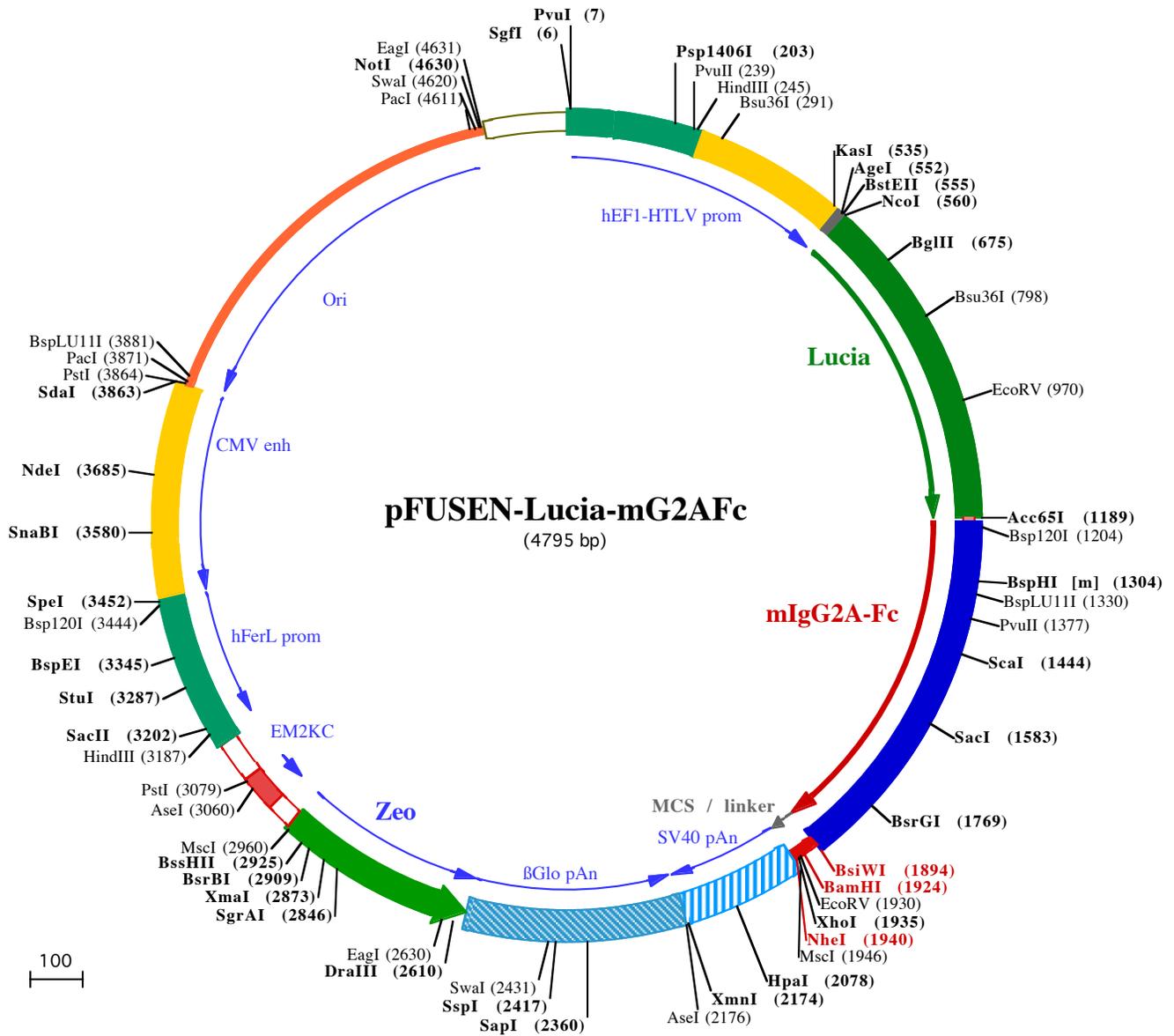
This antibiotic can be used for *E. coli* at 25 µg/ml in liquid or solid media and at 50-200 µg/ml to select Zeocin™-resistant mammalian cells.

RELATED PRODUCTS

Product	Catalog Code
Zeocin™	ant-zn-1
QUANTI-Luc™	rep-qlc1

TECHNICAL SUPPORT

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PvuI (7)
SgfI (6)
 1 GGATCTGCATCGCTCCGGTGCCCGTCAGTGGGAGAGCGCACATCGCCACAGTCCCGGAGAAGTTGGGGGAGGGGTCGGCAATTGAACGGGTGCCTA
 101 GAGAAGTGGCGCGGGTAAACTGGAAAGTGATGCTGTACTGGCTCCGCCTTTTCCGAGGGTGGGGGAGAACCCTATATAAGTGCAGTAGTCGCC

Psp1406I (203) **HindIII (245)** **PvuII (239)** **Bsu36I (291)**
 201 GTGAACGTTCTTTTTCGCAACGGGTTTGCCGCCAGAACACAGCTGAAGCTTCGAGGGGCTCGCATCTCTCTTACACGCCGCCGCCCTACCTGAGGCC
 301 GCCATCCACGCCGGTTGAGTCGCGTTTCTGCCGCTCCCGCCTGTGGTGCCTCCTGAAGCTGCGTCCGCCGTCTAGGTAAGTTTAAAGCTCAGGTCGAGACC
 401 GGGCCTTTGTCCGGCGCTCCCTTGAGGCTACCTAGACTCAGCCGGCTCTCCACGCTTTGCTGACCCTGCTTCTCAACTCTACGTTCTTTGTTTCGTTT

NcoI (560) **BstEII (555)**
KasI (535) **AgeI (552)**
 501 TCTGTTCTGCGCCGTTACAGATCCAAGCTGTGACCGCGCGCTACCTGAGATCACCGGTACCATGAAATCAAGGTGCTGTTTGCCTCATCTGTATTGC
 1 M E I K V L F A L I C I A
BglIII (675)
 601 TGGTGTGAGGCAAACCCACTGAAATCAATGAAGACCTCAATATAGCTGCTGTGGCCTCCAACCTTGGCCACCACAGATCTTGAGACTGACCTGTTCCACC
 13 V A E A K P T E I N E D L N I A A V A S N F A T T D L E T D L F T
Bsu36I (798)
 701 AACTGGGAGACCATGAATGTGATTAGCACTGACACAGAGCAGGTGAACACAGATGCTGACAGGGGCAAGCTGCCTGGCAAAAACTCCCCCAGATGTCC
 47 N W E T M N V I S T D T E Q V N T D A D R G K L P G K K L P P D V
 801 TGAGGGAGCTGGAGGCCAATGCCAGAAGGGCTGGTGCACAAGAGGCTGCCTCATTGGCTCTCCACATTAAGTGACCCCTAAGATGAAGAAATTTAT
 80 L R E L E A N A R R A G C T R G C L I C L S H I K C T P K M K K F I
EcoRV (970)
 901 CCCTGGCAGGTGCCACACTTATGAAGGTGAAAAGGAGTCTGCTCAGGGAGGGATTGGAGAGGCAATTGTTGATATCCCAGAGATTCTGGCTTCAAGGAT
 113 P G R C H T Y E G E K E S A Q G G I G E A I V D I P E I P G F K D
 1001 AAGGAGCCACTGGACCACTTTATTGCTCAAGTGGACCTCTGTGCTGATTGCACCACTGCTGTCTGAAGGGCCTTGCCAAATGTCCAGTCTCTGACCTCC
 147 K E P L D Q F I A Q V D L C A D C T T G C L K G L A N V Q C S D L
Acc65I (1189)
 1101 TGAAGAAGTGGCTTCCCCAGAGGTGACCACTTTTCCAGCAAGATTGAGGGTAGGGTGGACAAAATCAAGGGTCTGGCTGGGACAGAGGTACCGAGCC
 180 L K K W L P Q R C T T F A S K I Q G R V D K I K G L A G D R G T E P

Bsp120I (1204)
 1201 CAGAGGGCCACAATCAAGCCCTGTCTCCATGCAATGCCAGCACCTAACCTTTGGGTGGACCATCCGTCTTCATCTTCCCTCCAAGATCAAGGAT
 2 R G P T I K P C P P C K C P A P N L L G G P S V F I F P P K I K D
BspHI [m] (1304) **BspLU11I (1330)** **PvuII (1377)**
 1301 GTACTCATGATCTCCCTGAGCCCATATGCACATGTGTGGTGGATGTGAGCGAGGATGACCCAGATGTCCAGATCAGCTGGTTTGTGAACAACGTGG
 36 V L M I S L S P I V T C V V V D V S E D D P D V Q I S W F V N N V
ScaI (1444)
 1401 AAGTACACACAGCTCAGACACAAACCCATAGAGAGGATTACAACAGTACTCTCCGGTGGTCACTGCCCCTCCCATCCAGCACCAGGACTGGATGAGTGG
 69 E V H T A Q T Q T H R E D Y N S T L R V V S A L P I Q H Q D W M S G
SacI (1583)
 1501 CAAGGAGTCAAATGCAAGGTCAACAACAAGACCTCCAGCGCCATCGAGAGAACCATCTCAAACCAAAGGGTCAGTAAGAGCTCCACAGGTATAT
 102 K E F K C K V N N K D L P A P I E R T I S K P K G S V R A P Q V Y
 1601 GTCTTGCCTCCACCAGAAGAAGAGATGACTAAGAAACAGGTCACTCTGACCTGCATGGTCACAGACTTCATGCCTGAAGACATTTACGTGGAGTGGACCA
 136 V L P P P E E E M T K K Q V T L T C M V T D F M P E D I Y V E W T
BsrGI (1769)
 1701 ACAACGGGAAAACAGAGCTAAACTACAAGAACACTGAACCAGTCTGGACTCTGATGGTCTTACTTTCATGTACAGCAAGCTGAGAGTGGAAAAGAAGAA
 169 N N G K T E L N Y K N T E P V L D S D G S Y F M Y S K L R V E K K N
BsiWI (1894)
 1801 CTGGTGGAAAGAAATAGCTACTCTGTTCACTGTTGAGTGGTCCAGGGTCTGCACAATCACCACACGACTAAGAGCTTCTCCGGACTCCGGGTGCACTGACG
 202 W V E R N S Y S C S V V H E G L H N H H T T K S F S R T P G A
 Lys changed to Ala (1891)
 1 R T

EcoRV (1930) **NheI (1940)**
BamHI (1924) **XhoI (1935)** **MscI (1946)**
 1901 GGTGGTGGCGGTAGCGGTGGTGGCGGATCCGATATCTCGAGCTAGCTGCCAGACATGATAAGATACATTGATGAGTTTGACAAACCACAACCTAGAATG
 3 G G G G S G G G G S D I S S •

HpaI (2078)
 2001 CAGTGAATAAATGCTTTATTTGTGAAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAACAAGTTAAACAACAACATTGCATTC

AseI (2176) **XmnI (2174)**
 2101 ATTTTATGTTTCAGGTTCCAGGGGAGGTGTGGGAGTTTTTAAAGCAAGTAAACCTCTACAAATGTGGTATGGAATTAATTCTAAAATACAGCATAGC
 2201 AAACTTTAACCTCCAATCAAGCCTCTACTTGAATCCTTTTCTGAGGGATGAATAAGGCATAGGCATCAGGGGCTGTTGCCAATGTGCATTAGCTGTTT

SapI (2360)
 2301 GCAGCCTCACCTTCTTTCATGGAGTTTAAAGATATAGTGTATTTTCCAAAGTTTTGAAGTACTCTTCACTTTTATGTTTTAAATGCACTGACCTCCCA

2401 **SspI (2417)** SwaI (2431) CATTCCCTTTTTAGTAAAATATTCAGAAATAATTTAAATACATCATTGCAATGAAAATAAATGTTTTTTATTAGGCAGAATCCAGATGCTCAAGGCCTT
 2501 CATAATATCCCCAGTTTAGTAGTTGGACTTAGGGAACAAAGGAACCTTTAATAGAAATTGGACAGCAAGAAAGCGAGCTTCTAGCTTATCCTCAGTCTC
 1254 • D Q

2601 **DraIII (2610)** EagI (2630) GCTCCTCTGCCACAAAGTGCACGCAGTTGCCGGCCGGTTCGCGCAGGGCGAACTCCCGCCCCACGGCTGCTCGCCGATCTCGGTCATGGCCGGCCCGGA
 1224 E E A V F H V C N G A P D R L A F E R G W P Q E G I E T M A P G S
 2701 GCGTCCCGGAAGTTCGTGGACACGACCTCCGACCACTCGGCGTACAGCTCGTCCAGGCCGCGCACCCACACCCAGGCCAGGGTGTGTCCGGCACCACC
 894 A D R F N T S V V E S W E A Y L E D L G R V W V W A L T N D P V V

2801 **SgrAI (2846)** **XmaI (2873)** TGGTCTGGACCGCGCTGATGAACAGGGTACGTCGTCGCCGACACACCGGGGAAAGTCTCCTCCAGAAAGTCCCGGAGAACCCGAGCCGGTTCGGTCC
 554 Q D Q V A S I F L 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

2901 **BsrBI (2909)** **BssHII (2925)** MscI (2960) AGAACTCGACCGCTCCGGCGACGTCGCGCGGGTGAGCACCAGGAAAGCGGACTGGTCAACTTGGCCATGATGGCTCTCctgtcaggagaggaagagaag
 224 F E V A G A V D R A T L V P V A S T L K A M

3001 **AseI (3060)** **PstI (3079)** aaggttagtacaattgCTATAGTGAGTTGATTATACTATGCAGATATACTATGCCAATGATTAATTGTCAAAGTGGGCTGCAGgggttcattagtccac
 HindIII (3187)

3101 tttcctgcaactgccccatctcctgcccacccttccaggcatagacagtcagtgacttacCAAAGTACAGGAGGGAGAAGGCAGAAGCTTGAGACAG
 SacII (3202) **StuI (3287)**

3201 ACCCGCGGACCGCCGAAGTGCAGGGGACGTGGCTAGGGCGGCTTCTTTTATGTTGCGCGCCCTCGAGGCAGGGGCTCGGGGAGCCCTAGCGGCC

3301 **BspEI (3345)** AATCTGCGGTGGCAGGAGCGGGGCCGAAGCCGTGCCTGACCAATCCGGAGCACATAGGAGTCTCAGCCCCCGCCCAAAGCAAGGGGAAGTCACGCG

3401 **SpeI (3452)** **BspI20I (3444)** CCTGTAGCGCCAGCGTGTGTGAAATGGGGCTTGGGGGGTTGGGGCCCTGACTAGTCAAAACAAACTCCATTGACGTCAATGGGGTGGAGACTTGGGA
 SnaBI (3580)

3501 AATCCCCGTGAGTCAAACCGCTATCCACGCCCATTTGATGTACTGCCAAAACCGCATCATCATGGTAATAGCGATGACTAATACGTAGATGTACTGCCAAG

3601 **NdeI (3685)** TAGGAAAGTCCATAAGGTCATGTACTGGGCATAATGCCAGGCGGGCCATTTACCGTCATTGACGTCAATAGGGGGCGTACTTGGCATATGATACACTTG

3701 ATGTACTGCCAAGTGGGCAGTTTACCGTAAATACTCCACCCATTGACGTCAATGGAAAGTCCCTATTGGCGTTACTATGGGAACATACGTCATTATTGAC

3801 **PacI (3871)** **PstI (3864)** **SdaI (3863)** **BspLU11I (3881)** GTCAATGGGCGGGGTCGTTGGCGGTCAGCCAGGCGGGCCATTTACCGTAAGTTATGTAACCGCTGCAGGTTAATTAAGAACATGTGAGCAAAGGCCA
 3901 GCAAAAGGCCAGGAACCGTAAAAAGCCGCTTGTGCGGTTTTCCATAGGCTCCGCCCCCTGACGAGCATCAAAAAATCGACGCTCAAGTCAGAGG

4001 TGGCGAAACCCGACAGGACTATAAAGATACCAGGCTTTCCCCCTGGAAGTCCCTCGTGCCTCTCCTGTTCCGACCCTGCCGTTACCGGATACCTGT

4101 CCGCTTTCTCCCTTCGGGAAGCGTGGCGCTTTCTCATAGCTACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCCGCTCCAGCTGGGCTGTGTGCA

4201 CGAACCCCCGTTACGCCGACCGCTGCGCCTTATCCGGTAACTATCGTCTTGTAGTCCAACCCGTAAGACACGACTTATCGCCACTGGCAGCAGCCACT

4301 GGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTCTTGAAGTGGTGGCCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCT

4401 GCGCTCTGCTGAAGCCAGTTACCTTCGAAAAAGAGTTGGTAGCTCTTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTTGTTTGAAGCA

4501 GCAGATTACGCGCAGAAAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGTCTGACGCTCAGTGGAAACGAAAACCTCACGTTAAGGGATTTG

4601 **EagI (4631)** **PacI (4611)** **SwaI (4620)** **NotI (4630)** GTCATGGCTAGTTAATTAACATTTAAATCAGCGGCCGCAATAAAATATCTTTATTTTATTACATCTGTGTGTTGGTTTTTTGTGTAATCGTAACTAAC
 4701 ATACGCTCTCCATCAAAACAAAACGAAACAAAACAAACTAGCAAAATAGGCTGTCCCAAGTGCAGGTGCCAGAACATTTCTCTATCGAA