

pDUO2-mcs

A plasmid containing two multiple cloning sites and the hygromycin resistance gene

Catalog code: pduo2-mcs

<https://www.invivogen.com/pduo-mcs>

For research use only

Version 19I24-MM

PRODUCT INFORMATION

Contents

- 20 µg of pDUO2-mcs provided as lyophilized DNA
- 1 ml of Hygromycin B Gold at 100 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 Hygromycin B Gold 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

Toll-Like receptors (TLRs) play a critical role in early innate immunity to invading pathogens by sensing microorganisms. These evolutionary conserved receptors, homologues of the Drosophila Toll gene, recognize highly conserved structural motifs only expressed by microbial pathogens, called pathogen-associated microbial patterns (PAMPs). PAMPs include various bacterial cell wall components such as lipopolysaccharides (LPS), peptidoglycans and lipopeptides, as well as flagellin, bacterial DNA and viral double-stranded RNA. Stimulation of TLRs by PAMPs initiates a signaling cascade that involves a number of proteins, such as MyD88 and IRAK. This signaling cascade leads to the activation of the transcription factor NF-κB which induces the secretion of pro-inflammatory cytokines and effector cytokines that direct the adaptive immune response.

Ten human and twelve murine TLRs have been characterized, TLR1 to TLR10 in humans, and TLR1 to TLR9, TLR11, TLR12 and TLR13 in mice, the homolog of TLR10 being a pseudogene. In many instances, TLRs require the presence of a co-receptor to initiate the signaling cascade. One example is TLR4 which interacts with MD2 and CD14 to induce NF-κB in response to LPS stimulation. **pDUO2** is an expression vector designed to co-express two TLRs or TLR-related genes known to interact with each other.

The genes cloned into pDUO comprise the coding sequence (without introns) from the ATG to the Stop codon.

pDUO2-mcs does not contain a TLR gene and can be used in conjunction with other vectors of the pDUO2 family to serve as experimental controls

PLASMID FEATURES

- **hFerH and hFerL composite promoters:** Ferritin is a 24 subunit protein composed of two subunit types, termed H (heavy) and L (light), which perform complementary functions in the protein. Ferritin is ubiquitously expressed. Its synthesis is highly regulated by the iron status of the cell. The iron regulation is achieved at the translational level through the interaction between the iron-responsive element (IRE), located in the 5' untranslated region (5'UTR) of the ferritin mRNAs, and the iron regulatory protein⁴. To eliminate the iron regulation of the ferritin promoters, the 5'UTR of FerH and FerL have been replaced by the 5'UTR of the mouse and chimpanzee elongation factor 1 (EF1) genes, respectively.

MCS1 includes the following restriction sites:

Age I, Eco RV, Bam HI, Sal I and Avr II

- *Age I* is compatible with *Bsp E1* and *Sgr A1*
- *Eco RV* is compatible with any blunt-end restriction enzymes
- *Bam HI* is compatible with *Bgl II*, *Bst Y1* and *Bcl I*
- *Sal I* is compatible with *Ava I* and *Xho I*
- *Avr II* is compatible with *Xba I*, *Spe I* and *Nhe I*

MCS2 includes the following restriction sites:

Sgr A1, Bgl II, Xho I and Nhe I

- *Sgr A1* is compatible with *Bsp E1* and *Age I*
- *Bgl II* is compatible with *Bam HI*, *Bst Y1* and *Bcl I*
- *Xho I* is compatible with *Ava I* and *Sal I*
- *Nhe I* is compatible with *Xba I*, *Spe I* and *Avr II*

- **SV40 enhancer** which is comprised of a 72-base-pair repeat allows the enhancement of gene expression in a large host range. The enhancement varies from 2-fold in non-permissive cells to 20-fold in permissive cells. Furthermore, the SV40 enhancer is able to direct nuclear localization of plasmids⁵.

- **CMV enhancer:** The major immediate early enhancer of the human cytomegalovirus (HCMV), located between nucleotides -118 and -524, is composed of unique and repeated sequence motifs. The HCMV enhancer can substitute for the 72-bp repeats of SV40 and is severalfold more active than the SV40 enhancer⁶.

- **SV40 pAn:** the Simian Virus 40 late polyadenylation signal enables efficient cleavage and polyadenylation reactions resulting in high levels of steady-state mRNA. The efficiency of this signal was first described by Carswell *et al.*⁷

- **pMB1 Ori:** a minimal *E. coli* origin of replication to limit vector size, but with the same activity as the longer Ori.

- **FMDV IRES:** The internal ribosome entry site of the Foot and Mouth Disease Virus enables the translation of two open reading frames from one mRNA with high levels of expression⁸.

TECHNICAL SUPPORT

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- **EM7** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*.
- **hph gene** confers resistance to Hygromycin B both in *E. coli* and mammalian cells. In bacteria, *hph* is expressed from the constitutive *E. coli* EM7 promoter. In mammalian cells, *hph* is transcribed from the human FerH composite promoter as a polycistronic mRNA and translated via the FMDV IRES.
- **EF1 pAn** is a strong polyadenylation signal. InvivoGen uses a sequence starting after the stop codon of the EF1 cDNA and finishing after a bent structure rich in GT.

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.

Hygromycin B usage:

This antibiotic can be used for *E. coli* at 50-100 µg/ml in liquid or solid media and at 50-500 µg/ml to select Hygromycin-resistant mammalian cells.

References

1. Miyake K. et al., 2000. Innate recognition of lipopolysaccharide by Toll-like receptor 4/MD-2 and RP105/MD-1. *J Endotoxin Res.* 6(5):389-91.
2. Miyake K. et al., 1998. Mouse MD-1, a molecule that is physically associated with RP105 and positively regulates its expression. *J Immunol.* 161(3):1348-53.
3. Nagai Y. et al., 2002. Requirement for MD-1 in cell surface expression of RP105/CD180 and B-cell responsiveness to lipopolysaccharide. *Blood* 99(5):1699-705.
4. Eisenstein R.S. & Munro H.N., 1990. Translational regulation of ferritin synthesis by iron. *Enzyme* 44(1-4):42-58.
5. Dean D.A. et al., 1999. Sequence requirements for plasmid nuclear import. *Exp. Cell. Res.* 253:713-22.
6. Boshart M. et al., 1985. A very strong enhancer is located upstream of an immediate early gene of human cytomegalovirus. *Cell* 41(2):521-30.
7. 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.
8. Ramesh N. et al., 1996. High-titer bicistronic retroviral vectors employing foot-and-mouth disease virus internal ribosome entry site. *Nucleic Acids Res.* 24(14):2697-700.

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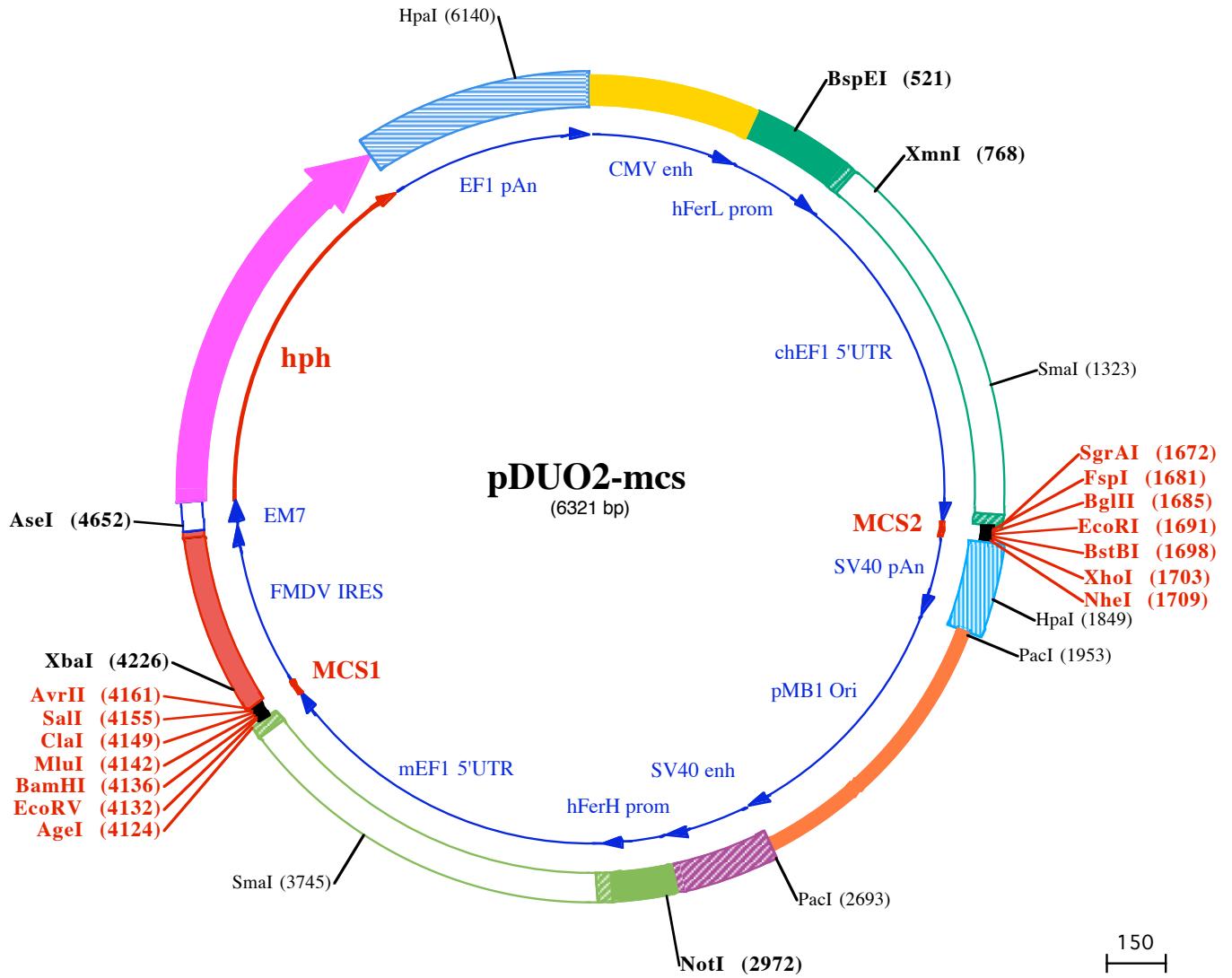
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1 CCTGCAGGGCTTACATAACTTACGGTAATGGCCCGCTGGCTGACCGCCCAACGACCCCCGCCATTGACGTCAATAATGACGTATGTTCCCAGTAGTAA
 101 CGCCAATAGGGACTTCATTGACGTCAATGGGTGGAGTTACGGTAACACTGCCACTGGCACTACATCAAGTGTATCATATGCCAAGTACGCC
 201 TATTGACGTCAATACGGTAATGGCCCGCTGGCATTATGCCACTACATGACCTTATGGACTTCCTACTTGGCAGTACATCTACGTATTAGTCATC
 301 GCTATTACCATGATGCGGTTGGCAGTACATCAATGGCGTGGATAGCGGTTGACTCACGGGATTCCAAGTCTCACCCATTGACGTCAATG
 401 GGAGTTGTTTGACTAGTCAGGGCCCAACCCCCCAAGCCCCATTCAACACACGCTGGCGTACAGGCGGTGACTTCCCTGTTGGCGGG
BspEI (521)
 501 GGGCTGAGACTCCTATGTCGGATTGGTCAGGCACGGCCTCGGCCCTGCCACCGCAGATTGGCGCTAGGCCCTCCGAGCGCCCTGCC
 601 TCCGAGGGCCGGCGCACATAAAAAGCCGCCCTAGCCACGTCCTCGCAGTTCGGCGTCCCGGGCTGTCTCAAGCTTGCAGAACACAG
XmnI (768)
 701 taagtccgtgtgggtcccgccctggctttacggttatggcccttgcgtgccttgaaatttcacccatgcccctggctgcagtagtgcattc
 801 ttgtatcccgagcttcgggtggaaatgggtggagatggcggccctgcgtgccttgcgttaaggagcccttcgcctcgcttgcgttgc
 901 ctggggccgcgcgtctaattctggcgcctgtctcgctgtttcgtaagtctctagccataaaattttgataaccagctgcacgc
 1001 cttttttctggcagatagttgttaatcgccggcaggatctgcacactgtgtatccgggtttggggccgcggccgcacggggccgtc
 1101 agccacatgttcggcgaggcgccctgcgagcgcggccaccgagaatcgacggggtagtctcaaactggccgcgtctggcctggc
 1201 gcccgtgtatccccccctggcgcaaggctggccggcaccagttgcgtgagcggaaatggccctccggccctgcgcaggagc
SmaI (1323)
 1301 tcaaaatggaggacgcggccggggagagcggccgggtgagtcacccacacaaggaaaggcccccattccatccgtcgcattgtactcc
 1401 cggagtaccggcgccgcaggcacccgtcattgtcgagctttggagtagtcgtcttaggtgggggggggggggggggggggggggg
 1501 ccacactgagttggggagactgaagagttggccagcttgcacttgcattgtatctcccttggaaatttgccttttgcatttt
FspI (1681) EcoRI (1691)
 1601 tcaagcctcagacagttcaagttttttttccatccatccatccatccatccatccatccatccatccatccatccatccatccatcc
SgrAI (1672) BglII (1685) BstBI (1698)
 1601 tcaagcctcagacagttcaagttttttttccatccatccatccatccatccatccatccatccatccatccatccatccatccatcc
NheI (1709)
XhoI (1703)
 1701 AACTCGAGGCTAGCTGGCCAGACATGATAAGATAACATTGATGAGTTGGACAAACCACAACTAGAATGCACTGAAAAAAATGCTTATTGTGAAATTG
HpaI (1849)
 1801 TGATGCTATTGCTTATTGTAACCATTATAAGCTGAATAACACAAGTTAACACAAATTGCAATTTCATTTATGTTCAAGGTCAGGGGAGGTGTTG
PacI (1953)
 1901 GAGGTTTTAAAGCAAGTAAACCTCTACAAATGTTGATGGAAATGTTAAATTACTAGCCATGACCAAAATCCCTAACGTGAGTTCTTCACTG
 2001 AGCGTCAGACCCCGTAGAAAGATCAAAGGATCTTCTGAGATCTTTCTGCGCTGAATCTGCTGCTGCAACACAAAAACCCCGTACCGCG
 2101 GTGGTTGTTGCCGGATCAAGAGCTACCAACTCTTCTGAGGTAACCTGGCTCAGCAGAGCGCAGATACAAATACTGTTCTAGTGTAGCG
 2201 AGTTAGGCCACCACTCAAGAACACTGTCAGCACCGCCTACATACCTCGCTGCTGAATCCTGTTACAGTGGCTGTCAGTGGCAGTC
 2301 TACCGGGTTGACTCAAGACGATAGTTACCGATAAGGCAGCGGCTGGCTGAACGGGGGTTCTGACACAGCCAGCTGGAGCGAACGAC
 2401 ACCGAACCTGAGATACTACAGCGTAGCTATGAGAAAGGCCACCGTCCGAAGGGGAAAGGGGAGCAGGTACCGTAAGCGGAGGGTGG
 2501 GAGAGCCACGAGGGAGCTCCAGGGGAAACGCCGGTATCTTATAGTCCTGCGGTTGCCACCTCTGACTTGAGCGTGA
PacI (2693)
 2601 GTCAGGGGGCGGAGCCTATGGAAAAACGCCAGCACGCGCCCTTTACGGTCTGGCTTTGCTGGCTTGTACATGTTTAATTAAACCTG
 2701 CAGGGCCTGAAATAACCTCTGAAAGAGGAATTGTTAGGTACCTCTGAGGCTGAAAGAACCCAGCTGTGGAATGTTGTCAGTTAGGGTGT
 2801 CCCAGGCTCCCAGCAGGAGAAGTATGCAAAAGCATGCAATTAGTCAGCAACCAGGTGTTGAAAGTCCCCAGGCTCCAGCAGGAGTATG
NotI (2972)
 2901 CAAAGCATCTCAATTAGTCAGCAACCATAGTCCACTAGTCCGCCAGAGCGCGCGAGGGCCCTCCAGCGCCGCCCTCCCCACAGCAGGGCG
 3001 GGTCCCGGCCACCGAAGGAGCGGGCTGGGGGGCGCTGATTGGCGGGGGCGCTGACGCCAGCGGTATAAGAGACCACAGCGACCC
 3101 GCAGGGCCAGACGTTCTCGCCGAAGCTTGCCTGAGACGAGgtgaggg
 3201 ggccggcccccgtgtcgccggggattagtcgtcgagcatcccggtcgatgtcgccggccggccggccggccggccggccggccggcc
 3301 gtagccctcgccctgtgtccggcttgcgtggatgcgtgtccgc
 3401 ttgcctctgtccgtgtccggatgcgtggatgcgtggatgcgtggatgcgtggatgcgtggatgcgtggatgcgtggatgcgtgg
 3501 aatggggggacaggagtggcgctggggcccccgttgcgtggatgcgtggatgcgtggatgcgtggatgcgtggatgcgtggatgc
 3601 caggctggggtagcgtgcgtggccatgtggcccccagcaccggcgttgcgtggatgcgtggatgcgtggatgcgtggatgcgtgg

SmaI (3745)

3701 atcccgccggcaccagttgcgtgcgtggaaagatggcgctccggccctgttgcaggagctaaaaatggaggacgcggcagccggtgagccggc
 3801 gggtagtcacccacacaaggaaaggccgtgtccctaccggctgtcttcgtgaccctgttgcattggccatagtccatcggcc
 3901 ttgagcacgcgtatgcggcgccccggggatgtatggcggtggatgttgcacattggggggactgtcggccatggcc
 4001 gtcattttggaaatttgtcccttgcgtttggcgactaatttcggcttgcgttcaaggatctttaaacccttttagGTGTTGTG

EcoRV (4132) MluI (4142) SalI (4155)
 AgeI (4124) BamHI (4136) ClaI (4149) AvrII (4161)

4101 AAAACCACCGCTAATTCAAAGCAACCGGTATCCGATCCACCGTATCGATTGCGACCCTAGGAGCAGGTTCCCAATGACACAAAACGTGCAACT

XbaI (4226)

4201 TGAAACTCCGCTGGTCTTCAGGGTAACCTTGACTGCCTGGTCCACGCTCGATCCACTGGCAGTGTAGTAACAGCACTGTT
 4301 GCTTCGTAGCGGAGCATGACGCCGTGGAACTCCCTGGTAAACAAGGACCCACGGGCCAAAGCCACGCCACCGGCCGTATGTGCAACC
 4401 CCAGCACGGCGACTTACTGCACCCACTTAAAGTGACATTGAAACTGGTACCCACACTGGTGACGGCTAAGGATGCCCTCAGGTACCCGAG
 4501 GTAACACCGGACACTGGATCTGAGAAGGGACTGGGCTCTATAAAAGCGCTGGTTAAAAGCTTCTATGCCGAATAGGTGACCGGAGGTCGGC

AseI (4652)

4601 ACCTTCCTTGCAATTACTGACCCATGAATACAATGACTGTTGACAATTATCATCGGCATAGTATATCGGCATAGTATAACGACTCACTATAG
 4701 GAGGCCACCATGAAGAACCTGAACTGACAGACTCTCTGGAGAAGTTCTATTGAAAATTGATTCTGTTCTGATCTCATGCAGCTGCTGAA
 4801 GGTGAAGAAAGCAGAGCCTTCTTCTTGATGTTGAGGAAGAGGTTATGTTCTGAGGGTCATTCTGCTGATGTTTACAAGACAGATATTT
 4901 31 Gl yGl uGl uSer ArgAl aPheSer PheAspVal Gl yGl yArgGl yTyrVal LeuArgVal AsnSer CysAl aAspGl yPheTyrLysAspArgTyrVal T
 4901 64 yrArgHi sPheAl aSer Al aAl aLeuProI leProGl uVal LeuAspI leGl yGl uPheSer Gl uSer LeuThr TyrCys I leSer ArgArgAl aGl nGl
 5001 97 yVal Thr LeuGl nAspLeuProGl uThr Gl uLeuProAl aVal LeuGl nProValAl aGl uAl aMetAspAl al I eAl aAl aAspLeuSer Gl nThr
 5101 131 TCTGGATTGGCTTGGCTTGGCCCAAGGATTGGCAGTACACCACTGGAGGATTCTGGCCATTGCTGATCTCATGCTATCAGTGAGCAGAAG
 5201 164 Ser Gl yPheGl yProPheGl yProGl nGl yI leGl yGl yTyrThr Thr TrpArgAspPheI leCysAl aI I eAl aAspProHi sVal I TyrHi sTrpGl nT
 5201 197 CTGTGATGGTACACAGTTCTGCTCTGCTGAGCACTGATGCTGCAACCTGCTGAGCAATGGATGCCATTGCGAGCTGATCTGAGCAGAACC
 5301 231 aAspPheGl ySer AsnAsnVal LeuThr AspAsnGl yArgGl yThrAl aVal I leAspTrpSer Gl uAl aMetPheGl yAspSer Gl nTyrGl uVal Al a
 5401 264 AsnI lePhePheTrpArgProTrpLeuAl aCysMetGl uGl nGl nThr ArgTyrPheGl uArgArgHi sProGl uLeuAl aGl ySer ProArgLeuArgA
 5501 297 1aTyrMetLeuArgI IleGl yLeuAspGl LeuTyrGl nSer LeuValAspGl yAsnAspAl aAl aTrpAl aGl nGl yArgCysAspAl aI leVa
 5601 331 1ArgSer Gl yAl aGl yThr Val Gl yArgThr Gl nI leAl aArgArgSer Al aAl aVal TrpThrAspGl yCysValGl uVal LeuAl aAspSer Gl yAsn
 5701 364 AGGAGACCTCCACAAGACCCAGAGCCAAGGAATGAATATTAGCTAGATTATGCCATAACCTGCCACCCACTCTTAATCAGTGGTGAAGGAAACGGCT
 5801 397 CAGAACTGTTGTTCAATTGGCCATTAAAGTTAGTAGTAAAGACTGGTAATGATAACAATGCATGTAACACCTCAGAAGGAAAGGAAATGTT
 5901 431 TGTGGACCCTTGGTTCTTGGCTGTCAGTTTAAGTTAGTTAGTTAAATCAGTACTTTAATGAAACACTGGACAAAAATTG
 6001 464 CACAGAATTGGAGACCCATTAAAAAGTTAAATGAGAACCTGTTGTCCTGGCAACACCGAGACATTAGGTGAAGACATCTAATTCTGGTT

HpaI (6140)

6101 TACGAATCTGAAACTCTTGAATGTAATTCTGAGTTAACACTTCTGGTGGAGAATAGGTTGTTCCCCACATAATTGGAAGGGAGGAAT
 6201 ATCATTAAAGCTATGGAGGGTTGTTGATTACAACACTGGAGAGAAATGCAGCATGTTGCTGATTGCTGTACTAAACAGGCCAAACTGAGTC
 6301 CTTGGTTGCATAGAAAGCT