pDUO2-mCD14/TLR2

A plasmid coexpressing the murine CD14 and TLR2 genes

Catalog code: pduo2-mcd14tlr2

https://www.invivogen.com/pduo-cd14-tlr2

For research use only

Version 19I23-MM

PRODUCT INFORMATION

Contents

- 20 µg of pDUO2-mCD14/TLR2 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.

To date, ten toll-like receptors have been reported in humans (TLR1 to TLR10) and only nine in mice (TLR1 to TLR9). 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 pDUO2 comprise the coding sequence (without introns) from the ATG to the Stop codon.

PLASMID FEATURES

• **Murine CD14** (1098 bp)**/Murine TLR2** (2352 bp)
TLR2 is involved in the recognition of multiple products of Gram-positive bacteria, mycobacteria and yeast. The first studies reported that TLR2 mediated LPS response but TLR2 has since been shown to confer responsiveness to the lipopeptides present in LPS preparations. However, it seems that some types of LPS can activate TLR2[1]. TLR2 is known to heterodimerize with other TLRs, a property believed to extend the range of PAMPs that TLR2 can recognize. TLR2 cooperates with TLR6 in the response to peptidoglycan[2] and diacylated mycoplasmal lipopeptide, and associtates with TLR1 to recognize triacylated lipopetides. Furthermore, pathogen recognition by TLR2 is strongly enhanced by CD14.

• **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 protein[3]. 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.

• **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 plasmids[4].

• **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 enhancer[5].

• **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.*[6]

• **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 expression[7].

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 (hygromycin resistance gene):** confers resistance to Hygromycin B both in *E. coli* and mammalin 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 H_2O. Store resuspended plasmid at -20°C.

Plasmid amplification and cloning

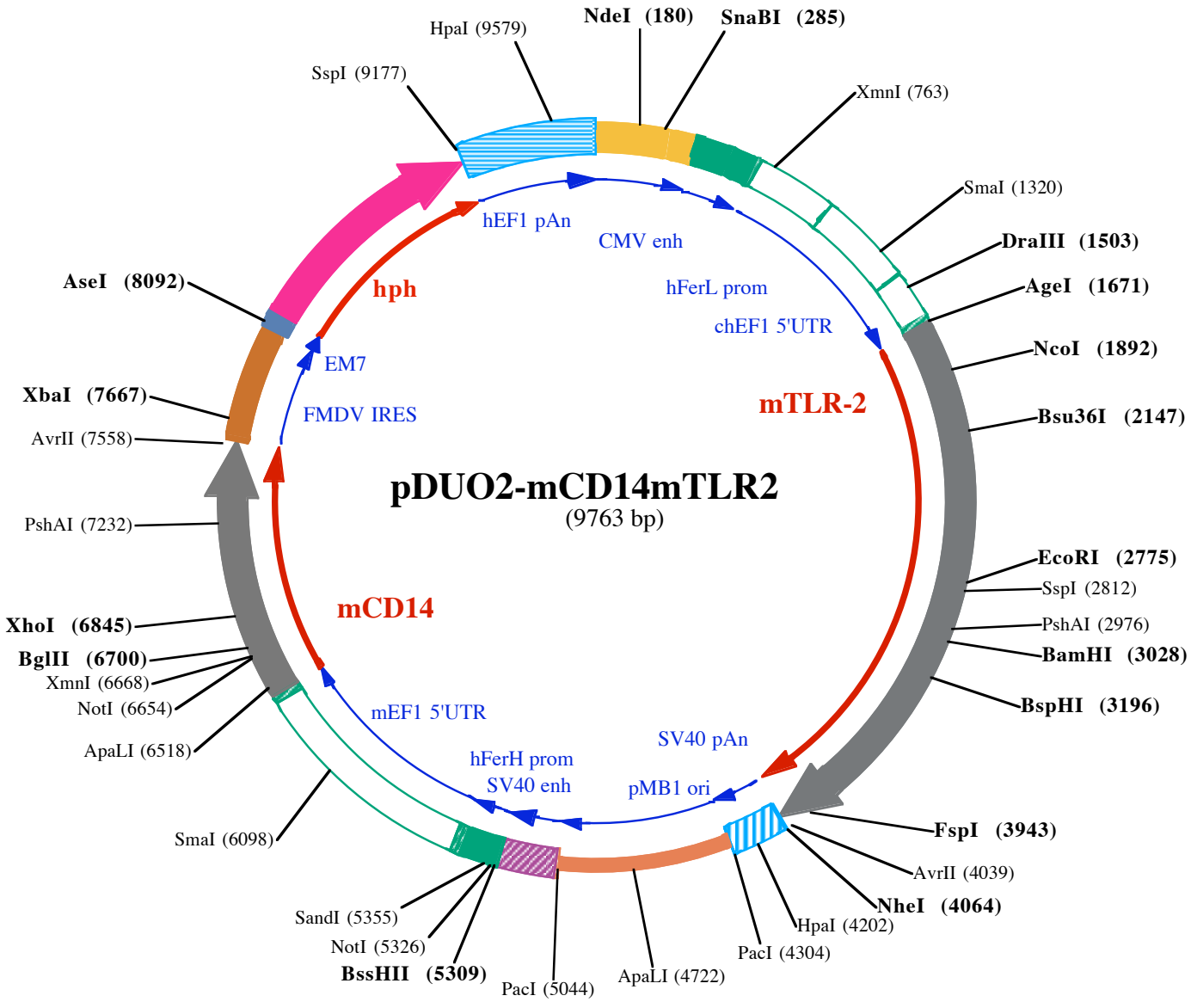
Plasmid amplification and cloning can be performed in *E. coli* GT116 or other commonly used laboratory *E. coli* strains, such as DH5α.

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. Netea MG. et al. 2002. Does the shape of lipid A determine the interaction of LPS with Toll-like receptors? Trends Immunol, 23(3):135-9.
2. Ozinsky A. et al. (2000). The repertoire for pattern recognition of pathogens by the innate immune system is defined by cooperation between Toll-like receptors. PNAS 97(25):13766-71.
3. Eisenstein RS. and Munro HN. 1990. Translational regulation of ferritin synthesis by iron. Enzyme 44(1-4):42-58.
4. Dean DA. *et al.* 1999. Sequence requirements for plasmid nuclear import. Exp. Cell. Res. 253:713-22.
5. Boshart M. *et al.* 1985. A very strong enhancer is located upstream of an immediate early gene of human cytomegalovirus. Cell 141(2):521-30.
6. Carswell S., and Alwine JC. 1989. Efficiency of utilization of the simian virus 40 late polyadenylation site: effects of upstream sequences. Mol. Cell Biol. 10: 4248-4258.
7. 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.


pDUO2-mCD14mTLR2
(9763 bp)
NdeI (180)
SnaBI (285)
XmnI (763)
SmaI (1320)
DraIII (1503)
AgeI (1671)
NcoI (1892)
Bsu36I (2147)
EcoRI (2775)
SspI (2812)
PshAI (2976)
BamHI (3028)
BspHI (3196)
FspI (3943)
AvrII (4039)
NheI (4064)
HpaI (4202)
PacI (4304)
ApaLI (4722)
PacI (5044)
BssHII (5309)
NotI (5326)
SandI (5355)
SmaI (6098)
ApaLI (6518)
NotI (6654)
XmnI (6668)
BglII (6700)
XhoI (6845)
PshAI (7232)
AvrII (7558)
XbaI (7667)
AseI (8092)
SspI (9177)
HpaI (9579)
hEF1 pAn
CMV enh
hFerL prom
chEF1 5'UTR
mTLR-2
SV40 pAn
pMB1 ori
hFerH prom
SV40 enh
mEF1 5'UTR
mCD14
EM7
FMDV IRES
hph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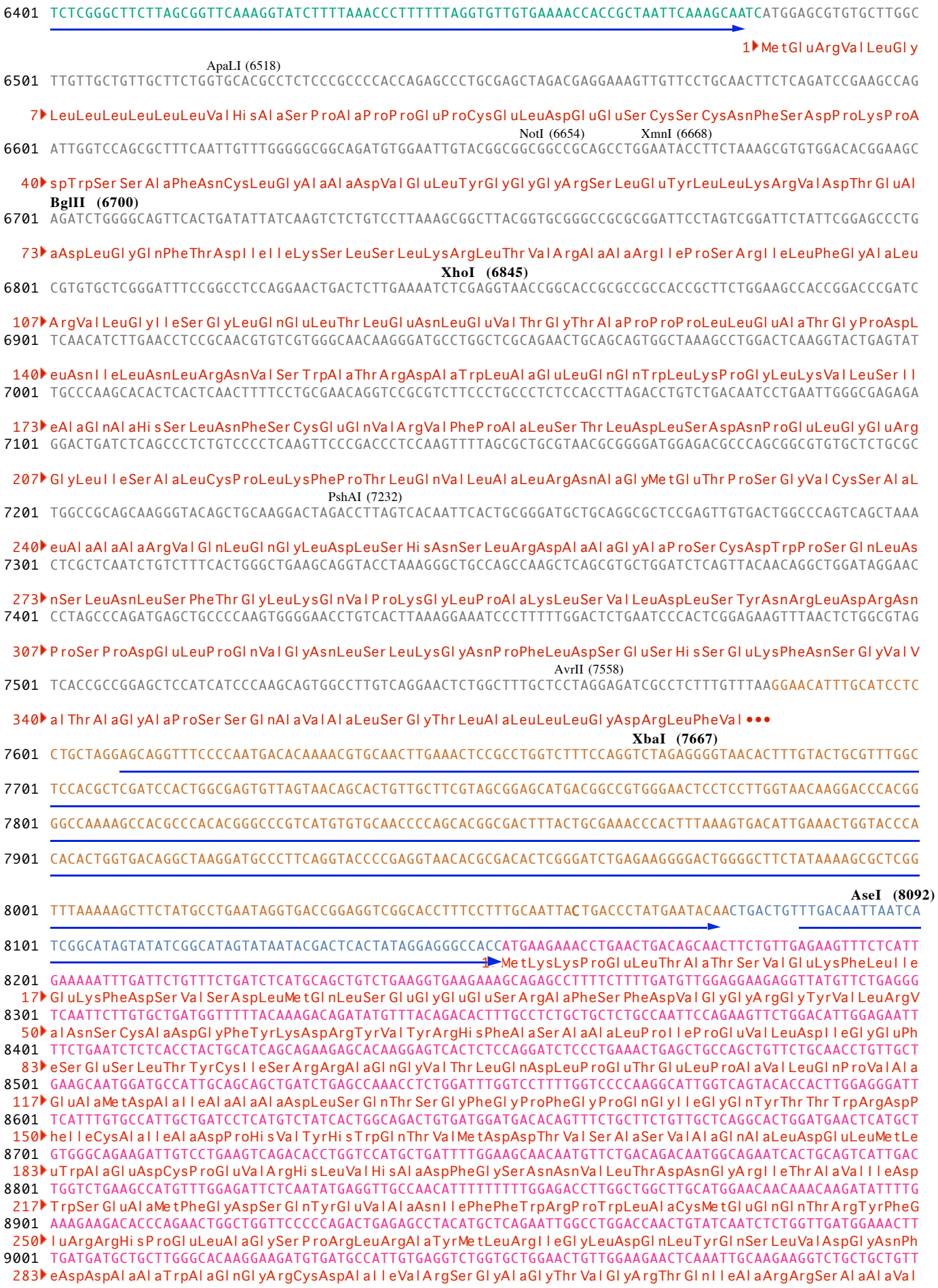
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AvrII (7558)
XbaI (7667)
AseI (8092)

SspI (9177)

9101 TGGACTGATGGATGTGTTGAAGTTCTGGCTGACTCTGGAAACAGGAGACCCTCCACAAGACCCAGAGCCAAGGAATGAATATTAGCTAGATTATCCCTAA

317 TrpThrAspGlyCysValGluValLeuAlaAspSerGlyAsnArgArgProSerThrArgProArgAlaLysGlu•••

9201 TACCTGCCACCCCACTCTTAATCAGTGGTGGAAGAACGGTCTCAGAACTGTTTGTTTCAATTGGCCATTTAAGTTTAGTAGTAAAAGACTGGTTAATGAT

9301 AACAATGCATCGTAAAACCTTCAGAAGGAAAGGAGAATGTTTTGTGGACCACTTTGGTTTTCTTTTTTGCGTGTGGCAGTTTTAAGTTATTAGTTTTTAA

9401 AATCAGTACTTTTTAATGGAAACAACTTGACCAAAAATTTGTCACAGAATTTTGAGACCCATTAAAAAAGTTAAATGAGAAACCTGTGTGTTCCTTTGGT

HpaI (9579)

9501 CAACACCGAGACATTTAGGTGAAAGACATCTAATTCTGGTTTTACGAATCTGGAAACTTCTTGAAAATGTAATTCTTGAGTTAACACTTCTGGGTGGAGA

9601 ATAGGGTTGTTTTCCCCCCACATAATTGGAAGGGGAAGGAATATCATTTAAAGCTATGGGAGGGTTGCTTTGATTACAACACTGGAGAGAAATGCAGCAT

9701 GTTGCTGATTGCCTGTCACTAAAACAGGCCAAAAACTGAGTCCTTGGGTTGCATAGAAAGCTG