

pBOOST2-wthIRF1

New DNA vaccine adjuvant of the pVAC plasmids expressing the wild-type IRF1 gene

Catalog # pbst2-wthirf1

For research use only

Version 20K16-MM

PRODUCT INFORMATION

Content:

- 20 µg of lyophilized pBOOST2-wthIRF1 plasmid expressing the human wild type IRF1 gene
- 1 ml of Zeocin™ (100 mg/ml)

Shipping and storage:

Products are shipped at room temperature. Lyophilized DNA is stable for 12 months when stored at -20°C. Resuspended DNA is stable for 12 months when stored at -20°C. Avoid repeated freeze-thaw cycles. 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

pBOOST2 plasmids were developed as genetic adjuvants for DNA vaccines to potentiate the immune response to a specific antigen. They feature different genes from the interferon regulatory factor family (IRF). IRFs are transcriptional activators for IFN- α , IFN- β and IFN-stimulated genes. In particular IRF-1, IRF-3 and IRF-7 act as direct transducers of virus-mediated signaling pathways activating IFN- α and IFN- β in infected cells. Recently, IRF-1, IRF-3 and IRF-7 were shown to be able to bias T cells towards type 1 or type 2 immune responses, leading to the activation of cytotoxic T cells and/or the production of antibodies. The method of plasmid DNA vaccine delivery is known to bias the immune response to a specific antigen towards a type 1 (T-cell) or type 2 (antibody) response¹. These biases can be further enhanced by the codelivery of IRFs to increase the efficacy of the vaccination^{2,3}.

PLASMID FEATURES

- **wthIRF1** (wild type human interferon regulatory factor 1)
IRF-1 primarily increases Th2 antibody responses². Following intramuscular or gene gun injections of DNA vaccines, IRF-1 can increase the titers of antibodies up to 10-fold.
- **hEF1 / HTLV prom** is a composite promoter comprising the Elongation Factor-1 α (EF-1 α) core promoter⁴ and the R segment and part of the U5 sequence (R-U5') of the Human T-Cell Leukemia Virus (HTLV) Type 1 Long Terminal Repeat⁵. 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.
- **SV40 pAn**: The Simian Virus 40 late polyadenylation signal enables efficient cleavage and polyadenylation reactions resulting in high levels of steady-state mRNA.
- **Ori** is a minimal *E. coli* origin of replication with the same activity as the longer Ori.

- **EM7** is a bacterial promoter that enables the constitutive expression of the antibiotic resistance gene in *E. coli*.

• **Sh-ΔCpG (Synthetic Zeocin® gene):** The *Sh ble* gene from *Streptomyces hindustanus* encodes a small protein that confers resistance to Zeocin™ by binding to the antibiotic. To reduce the amount of CpG motifs that may skew the raised antigen-specific immune response, pBOOST2 contains a CpG-free allele of the *Zeo^R* gene. All CpGs from the wild-type gene (50) were removed by synthesizing a new allele that contains no CpGs but encodes the exact same protein sequence.

References:

1. Robinson HL., 1999. DNA vaccines: basic mechanism and immune responses (Review). *Int J Mol Med*. 4(5):549-55.
2. Sasaki S. *et al.*, 2002. Regulation of DNA-raised immune responses by cotransfected interferon regulatory factors. *J Virol*. 76(13):6652-9.
3. Bransom JL. *et al.*, 2003. Super-activated interferon-regulatory factors can enhance plasmid immunization. *Vaccine*. 21(13-14):1363-70.
4. Kim, D.W. *et al.*, 1990. Use of the human elongation factor 1 alpha promoter as a versatile and efficient expression system. *Gene* 2: 217-223.
5. Takebe, Y. *et al.*, 1988. R 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*. 1: 466-472.

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.

TECHNICAL SUPPORT

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

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Intramuscular inoculation

Plasmid DNA solution

- Prepare the vaccine plasmid solution by resuspending 10 μ g of the vaccine plasmid DNA in 50 μ l saline solution.
- Prepare the pBOOST2 solution by mixing 10 μ g of pBOOST2-wthIRF1 and 90 μ g of the mock plasmid pBOOST2-null in 50 μ l saline solution for low dose, or 100 μ g of pBOOST2-wthIRF1 in 50 μ l saline solution for high dose.
- Combine both solutions to obtain a total of 110 μ g DNA in 100 μ l saline solution.

Note: The quantities are per mouse.

Intramuscular injections

- Inoculate 6 to 8-week old female BALB/c mice with 100 μ l plasmid DNA solution (described above) into the quadriceps at 0 and 4 weeks.
- Collect sera and analyze for antibodies at 8 weeks.

Note: For more information see the article by Sasaki S. et al.¹

TECHNICAL SUPPORT

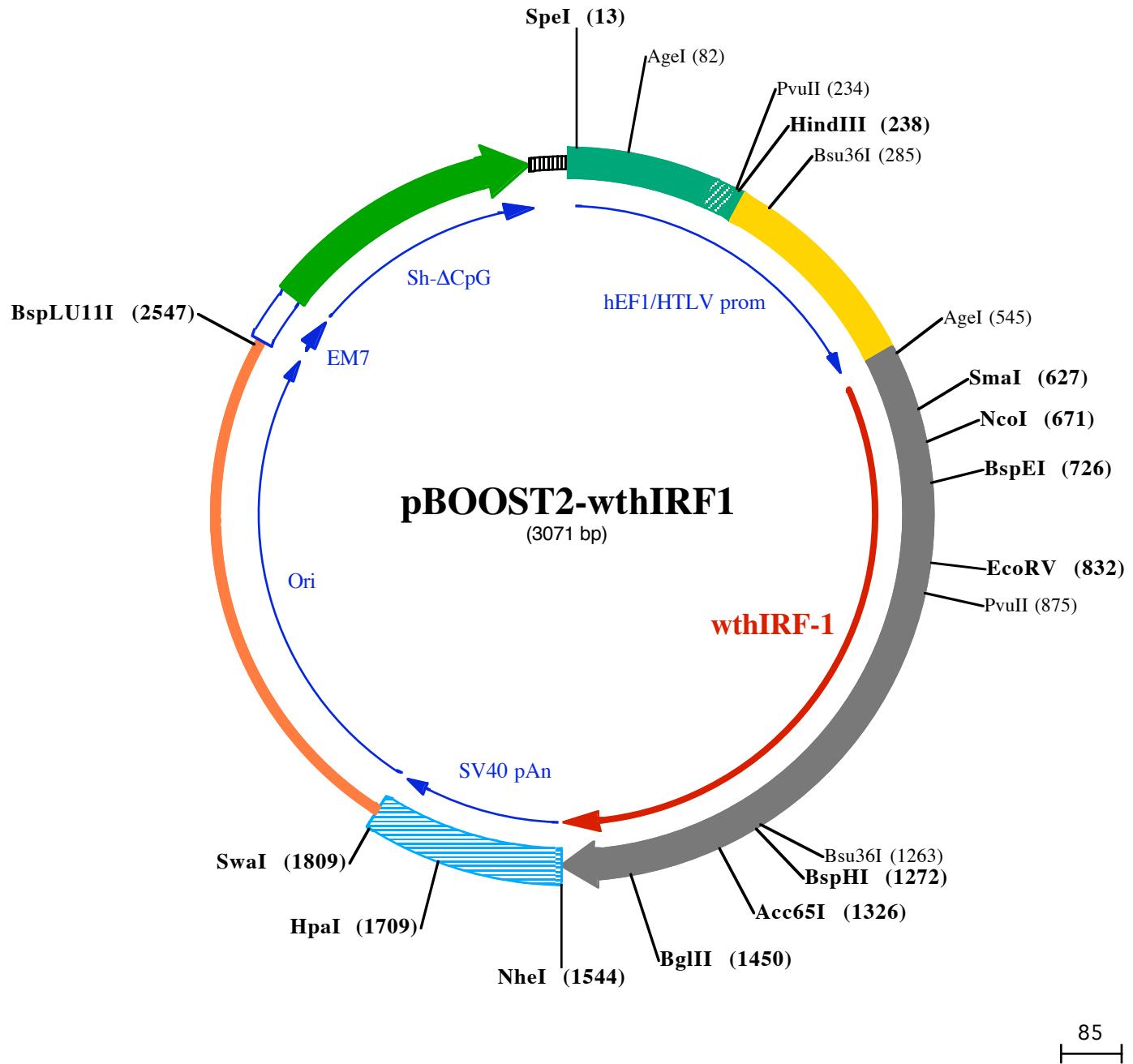
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SpeI (13)

1 CCTGCAGGCCACTAGTCAGTGGCAGAGCGCACATGCCACAGTCCCAGAGAAGTTGGGGAGGGTCGCAATTGAACCGGTCTAGAGAAGGT

101 GGCAGGGTAACTGGAAAGTGATTCGTACTGGCTCCGCTTTCCGAGGGTGGGAGAACGTATAAGTCAGTAGTCGGTGAACGT

HindIII (238)

PvuII (234)

201 TCTTTTCGCAACGGTTGCCAGAACACAGCTGAAGCTCGAGGGCTGCATCTCCTCACGGCCGCCCTACCTGAGGCCATCCA

301 CGCCGGTTAGTCGCTCTGCCCTCCGCTGTTGCTGAACCTGCCTCCGCTAGGTAAGTTAAAGCTCAGGTCGAGACGGGCTT

401 GTCCGGCGCTCCCTGGAGCTACCTAGACTCAGCCGCTCCACGCTTGCTGACCTGCTCAACTCTACGTTGCTTCTGTTCT

AgeI (545)

501 GCGCGTTACAGATCCAAGCTGACCGCGCTACCTGAGATCACGGTAGGAGGCCAGCATGCCCATCACTGGATGCGCATGAGACCTGGCTAGA
1► Met Pro Ile Thr Arg Met Arg Met Arg Pro Trp Leu Gl

SmaI (627)

BsU36I (285)

601 GATGCAGATTAATTCAAACAAATCCGGGCTATCTGATTAATAAGAGGAGATGATCTTCAGATCCATGAAAGCATGCTGCCAAGCATGGCTGG
13► uMet Gl nIle Asn Ser Asn Gl nIle Pro Gl yLeu lIe Trp lIe Asn Lys Gl uGl uMet lIe Phe Gl nIle Pro Trp Lys His Al aAl aLys His Gl yTrp

BspEI (726)

701 GACATACAAGGATGCCCTGTTGGAGCTGGCCATTACACAGGCCATACAAAGCAGGGAAAAGGAGGCCAGATCCAAGCAGTGGAGGCCA
47► Asp lIe Asn Lys Asp Al aCys Leu Phe Arg Ser Trp Al aIe Hi sThr Gl yArg Tyr Lys Al aGl yGl uLys Gl uPro Asp Pro Lys Thr Trp Lys Al aA

EcoRV (832)

PvuII (875)

801 ACTTCGCTGTGCCATGAACTCCCTGCCAGATATCGAGGAGGTAAAGACCAGAGCAGGAACAAGGGCAGCTCAGCTGTGCGAGTGTACCGGATGCTCC
80► snPhe Arg Cys Al aMet Asn Ser Leu Pro Asp lIe Gl uGl uVal Lys Asp Gl nSer Arg Asn Lys Gl ySer Ser Al aVal lArg Val Tyr Arg Met Leu Pr
901 ACCTCTACCAAGAACAGAGAAAAGAAAGCTGAAGTCCAGCCGAGATGCTAAAGGCAAGAGGAAGTCATGTTGGGATTCAGCCCTGAT
113► oPro Leu Thr Lys Asn Gl nArg Lys Gl yArg Ser Ser Arg Asp Al aLys Ser Lys Al aLys Arg Lys Ser Cys Gl yAsp Ser Ser Pro Asp
1001 ACCTCTCTGATGGACTCAGCAGCTCCACTCTGCTGATGACCAACAGCAGCTACACAGTTCCAGGCTACATGCAAGACTTGGAGGTGGAGCAGGCCCTGA
147► Thr Phe Ser Asp Gl yLeu Ser Ser Thr Leu Pro Asp Asp His Ser Ser Tyr Thr Val lPro Gl yTyr Met Gl nAsp Leu Gl uVal lGl uGl nAl aLeu T
1101 CTCCAGCACTGTCGCCATGTGCTGTCAGCAGCACTCCCCGACTGGCACATCCAGTGGAGTTGTGCCGGACAGCACCAAGTGTATCTGTACAACCTCCA
180► hr Pro Al aLeu Ser Pro Cys Al aVal Ser Ser Thr Leu Pro Asp Trp His lIe Pro Val Gl uVal Val lPro Asp Ser Thr Ser Asp Leu Tyr Asn Phe Gl

Bsu36I (1263) **BspHI (1272)**

1201 GGTGTACCCATGCCCTCACCTCTGAAGCTAACACAGATGAGGATGAGGAAGGGAAATTACCTGAGGACATCATGAAGCTTGGAGCAGTCGGAGTGG
213► nVal Ser Pro Met Pro Ser Thr Ser Gl uAl aThr Thr Asp Gl uAsp Gl uGl yLys Leu Pro Gl uAsp lIe Met Lys Leu Leu Gl uGl nSer Gl uTrp

Acc65I (1326)

1301 CAGCCAACAAACGTTGGATGGGAAGGGTACCTACTCAATGAACCTGGAGTCCAGCCCACCTCTGCTATGGAGACTTAGCTGTAAGGAGGCCAGAAA
247► Gl nPro Thr Asn Val Asp Gl yLys Gl yTyr Leu Leu Asn Gl uPro Gl yVal Gl nPro Thr Ser Val lTyr Gl yAsp Phe Ser Cys Lys Gl uGl uPro Gl uI

BglII (1450)

1401 TTGACAGCCAGGGGGATATTGGCTGAGTCTACAGCTGCTTACAGAGATCTGAAGAACATGGATGCCACCTGGCTGGACAGCCTGCTGACCCAGT
280► lEasp Ser Pro Gl yGl yAsp lIe Gl yLeu Ser Leu Gl nArg Val Phe Thr Asp Leu Lys Asn Met Asp Al aThr Trp Leu Asp Ser Leu Leu Thr Pro Va

NheI (1544)

1501 CCGGTTGCCCTCCATCCAGGCCATTCCCTGTCAGCGTAGCTGACCTGAGCTCGACATGATAAGATAACATTGATGAGTTGGACAAACCACAACATGAG
313► lArg Leu Pro Ser lIe Gl nAl aIle Pro Cys Al aPro ***

1601 AGTAAAAAAATGCTTATTGTGAAATTGTGATGCTATTGTTATTGTGAAATTGTGATGCTATTGCTTATTGTAACCATTATAAGCTGAAT

HpaI (1709)

1701 AAACAAGTTAACAAACAATTGCAATTCTATTGTTAGGTTCAAGGGAGGTGGAGGTTTTAAAGCAAGTAAACCTTACAAATGTGTA
→

SwaI (1809)

1801 GATCCATTAAATGTTAATTAACTAGCCATGCCAAATCCCTAACGTGAGTTCTTCACTGAGCCTGAGACCCGTAGAAAAGATCAAAGGATCT

1901 TCTTGAGATCTTTCTGCGTAATCTGCTGCTGCAAACAAAAACCCACCGCTACCAGCGTGGTTTGGCGATCAAGAGCTACCAACTC

2001 TTTTCCGAAGGTAACTGGCTTCAGCAGAGCGCAGATACCAAATACTGTTCTAGTGTAGCCGTAGTTAGGCCACCACTCAAGAACTCTGACCG

2101 GCCTACATACCTGCTCTGCTAATCTGTTACCGAGCTGCTGCCAGTGGCTGCTGATAAGTCGTGCTTACCGGGTGGACTCAAGACGATAGTTACCGGAT

2201 AAGGCGCAGCGTGGCTGAACGGGGGTTGTCACAGCCCAGCTGGAGCGAACCTACCGAACCTACAGCGTGAAGCTGAGATAACCTACAGCGTGAAGCTATGAG

2301 AAAGGCCACGCTCCGAAGGGAGAACAGGCGACAGGTATCCGTAAGCGCAGGGTCGAACAGGAGAGCGCACAGGGAGCTCAGGGAAACGC

2401 CTGGTATCTTATAGCTCTGCGGTTGCCACCTCTGACTTGAGCGTCGATTTGTGATGCTCAGGGGGCGGAGCCTATGGAAAACGCCAGC

BspLU11I (2547)

2501 AACCGGCCCTTTACGGTCTGGCTTTGCTCACATGTTCTAATTAAATTTCAAAAGTAGTTGACATTACATCGGCATAG
→

2601 TATATCGGCATAGTATAACGACTCAATAAGGGCCATCATGGCCAAGTTGACCGATGCTGCTCCAGTGCACAGCCAGGGATGTGGCTGGAGCT
1► Met Al aLys Leu Thr Ser Al aVal lPro Val Leu Thr Al aArg Asp Val Al aGl yAl aV
2701 GTTGAAGTCTGGACTGACAGGTTGGGTTCTCAGAGATTGAGGATGACTTGTGAGGATGACTTGTGAGGATGAGCTGAGTGTGAGGAGTGGCTCCACCAACTT
20► Val Gl uPhe Trp Thr Asp Arg Leu Gl yPhe Ser Arg Asp Phe Val Gl uAsp Asp Phe Al aGl yVal Val lArg Asp Asp Val Thr Leu Phell Ie Ser Al aV
2801 TCCAGGACCAAGGTGGCTGACAACACCTGGCTGGGTGAGAGGACTGGATGAGCTGAGTGTGAGGAGTGGCTCCACCAACTT
53► al Gl nAsp Gl nVal Val lPro Asp Asn Thr Leu Al aTrp Val Trp Val lArg Gl yLeu Asp Gl uLeu Tyr Al aGl uTrp Ser Gl uVal Val Ser Thr Asn Ph
2901 CAGGGATGCCAGTGGCCCTGCCATGACAGAGATTGGAGAGCAGGCCCTGGGGAGAGAGTGGCTGAGAGACCCAGCAGCACTGTGCACTTGTG
86► eArg Asp Al aSer Gl yPro Al aMet Thr Gl uIle Gl yGl uGl nPro Trp Gl yArg Gl uPhe Al aLeu Arg Asp Pro Al aGl yAsn Cys Val His Phe Val
3001 GCAGAGGAGCAGGACTGAGATAAGAATTGAAACAAAAACCCGCCCGGGTTTTGTTAAATTAA
120► Al aGl uGl nAsp ***