

# pUNO1-SpikeV6-dfur

Expression vector encoding the SARS-CoV-2 N.Y. variant (B.1.569 lineage) Spike (delta furin) gene

Catalog code: p1-spike-v6-df

<https://www.invivogen.com/ny-b1526-spike-expression-vectors>

For research use only

Version 21E11-ED

## PRODUCT INFORMATION

### Contents

- 20 µg of lyophilized pUNO1-SpikeV6-dfur (plasmid DNA)
- 2 x 1 ml of Blasticidin (10 mg/ml)

### Storage and Stability

- Product is shipped at room temperature.
- Store lyophilized DNA at -20°C.
- Resuspended DNA is stable for 1 year at -20°C.
- Store Blasticidin at 4°C or -20°C. The expiry date is specified on the product label.

### Quality control

- Plasmid construct is confirmed by restriction analysis and full-length open reading frame (ORF) sequencing.
- After purification by ion exchange chromatography, predominant supercoiled conformation is verified by electrophoresis.

## PLASMID FEATURES

### New York Variant SARS-CoV-2 Spike cassette

- EF-1 $\alpha$ /HTLV hybrid promoter is a composite promoter comprised of the Elongation Factor-1 $\alpha$  (EF-1 $\alpha$ ) core promoter<sup>1</sup> and the 5' untranslated region of the Human T-Cell Leukemia Virus (HTLV). EF-1 $\alpha$  utilizes a type 2 promoter that encodes a "house-keeping" gene. It is expressed at high levels in all cell cycles and lower levels during the G0 phase. Additionally, since the promoter is not tissue-specific it is highly expressed in all cell types. The R segment and part of the U5 sequence (R-U5') of the HTLV Type 1 Long Terminal Repeat<sup>2</sup> has been coupled to the EF-1 $\alpha$  promoter to enhance stability of DNA and RNA. This modification not only increases steady state transcription, but also significantly increases translation efficiency.

### • Codon-optimized Spike ORF

pUNO1-SpikeV6-dfur contains the Spike (S) coding sequence from the New York (N.Y.) variant (B.1.526 lineage). This variant is characterized by a number of mutations within the Spike coding sequence (see below)<sup>3</sup>. The furin cleavage site in pUNO1-SpikeV4-dfur has been inactivated (dfur) by the inclusion of two mutations (R683/5A). Furthermore, to improve expression of the S protein in cell lines, the gene is codon-optimized and the last 19 amino acids, which contain an ER-retention motif (KxHxx), have been removed<sup>4,5</sup>.

pUNO1-SpikeV6-dfur includes the following sequence features:

- **S1 domain:** L5F, T95I, D253G, D614G
- **RBD:** E484K
- **S1/S2 boundary:** R683A, R685A
- **S2 domain:** A701V

Spike (S) is a structural glycoprotein expressed on the surface of SARS-CoV-2. It mediates membrane fusion and viral entry into target cells upon binding to the host receptor ACE2 and the proteolytic activity of host proteases such as furin and TMPRSS2<sup>6</sup>.

For more information visit: <https://www.invivogen.com/sars2-spike>

- SV40 pAn is the Simian Virus 40 late polyadenylation (pAn) signal and it enables efficient cleavage and polyadenylation reactions resulting in high levels of steady-state mRNA<sup>7</sup>

### Antibiotic selection cassette

- hCMV (human cytomegalovirus) enhancer & promoter drive the expression of the blasticidin resistance gene (*bsr*) in mammalian cells.
- EM7 is a bacterial promoter that enables the constitutive expression of the blasticidin resistance gene (*bsr*) in *E. coli*.
- *bsr* (blasticidin resistance gene) encodes a deaminase from *Bacillus cereus* that confers resistance to the antibiotic blasticidin. The expression of the *bsr* gene is driven by the CMV promoter/enhancer and the bacterial EM7 promoter. Therefore, Blasticidin can be used to select stable clones in mammalian cells and *E. coli* transformants.
- Human β-Globin pAn is a strong polyadenylation (pAn) signal placed downstream of *bsr*. The use of β-globin pAn minimizes interference and possible recombination events with the SV40 pAn signal<sup>8</sup>.

### General features of pUNO1-SpikeV6-dfur

- pMB1 ori is a minimal *E. coli* origin of replication.

## APPLICATIONS

### Stable gene expression in mammalian cells.

pUNO1 plasmids are designed for both transient and stable transfection in mammalian cell lines by selection with Blasticidin. Furthermore, they facilitate high levels of expression of the gene of interest.

### Antibody screening by flow cytometry

pUNO1-SpikeV6-dfur has been specifically designed for mammalian cell expression of the SARS-CoV-2 S protein. Notably, due to the inactivated furin cleavage site, when this plasmid is expressed by a host cell (e.g. 293T cells) there is high surface expression of the full-length S protein<sup>4,9</sup>. Ideal for SARS-CoV-2 S-specific antibody screening by flow cytometry (*in-house data*).

## 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 water.
- Store the 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.

### • Blasticidin usage

Blasticidin should be used at 25-100 µg/ml in bacteria and 1-30 µg/ml in mammalian cells. Blasticidin is supplied as a 10 mg/ml colorless solution in HEPES buffer.

## TECHNICAL SUPPORT

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## REFERENCES

1. Kim D. et al., 1990. Use of the human elongation factor 1 $\alpha$  promoter as a versatile and efficient expression system. Gene 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.
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6. Hoffmann M. et al., 2020. SARS-CoV-2 cell entry depends on ACE2 and TMPRSS2 and is blocked by a clinically proven protease inhibitor. Cell. 181:1-16.
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8. Yu J. & Russell J., 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.
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## RELATED PRODUCTS

Product	Description	Cat. Code
Blasticidin ChemiComp GT116	Selection antibiotic Competent <i>E.coli</i>	ant-bl-1 gt116-11
<b>COVID-19 Product Range</b>		
HEK-Blue™ hACE2 Cells	Cell line	hkb-hace2
A549-hACE2-TMPRSS2 Cells	Cell Line	a549-hace2-tpsa
pUNO1-hACE2	Expression vector	puno1-hace2
pUNO1-hTMPRSS2a	Expression vector	puno1-htp2a
Anti-CoV2RBD-c1-hlgG1	Recombinant Antibody	cov2rbdc1-mab1

For a complete list of InvivoGen's COVID-19 related products visit:  
<https://www.invivogen.com/covid-19>

### TECHNICAL SUPPORT

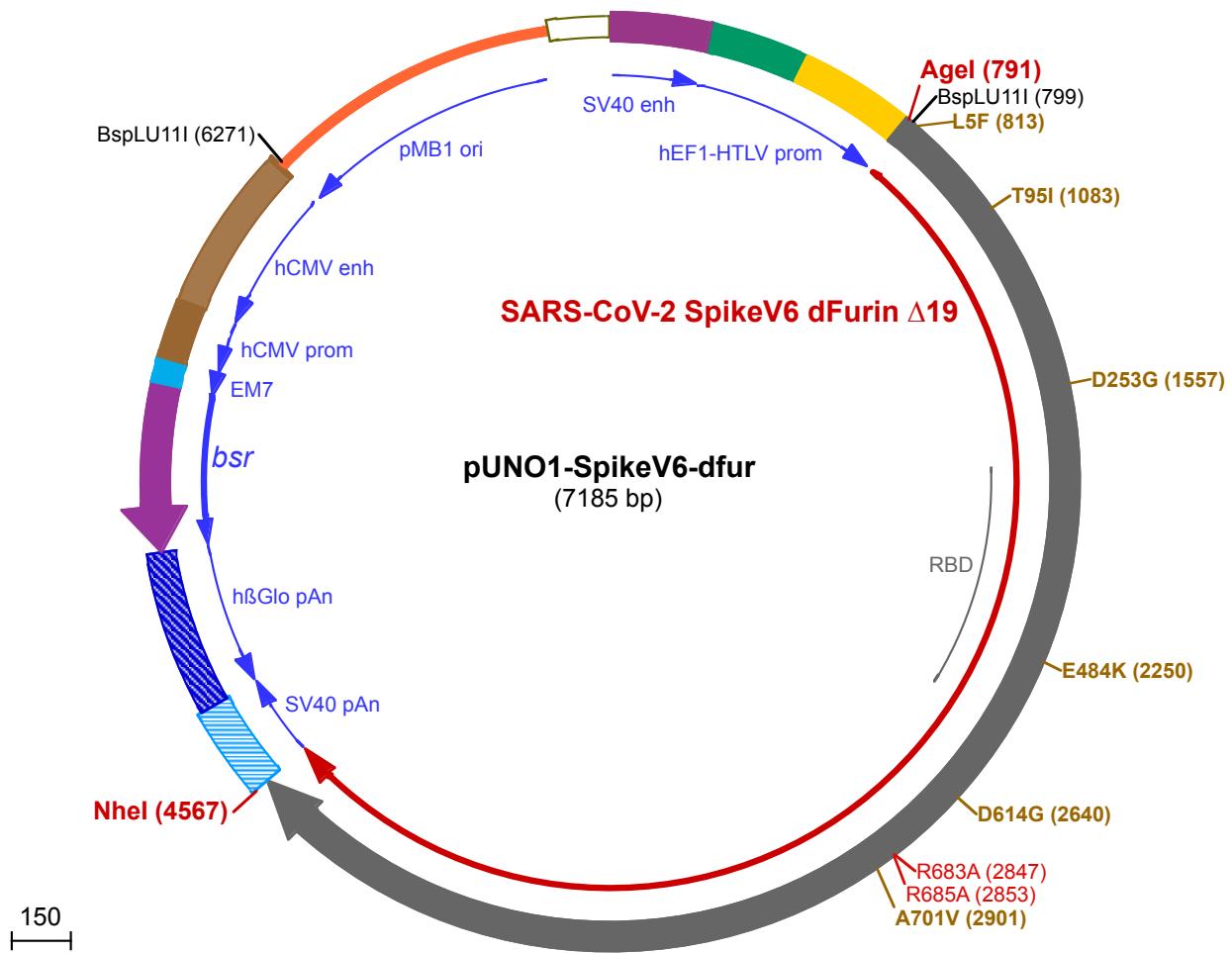
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1 GGACCTGCAGGCCGCTGAAATAACCTCTGAAAGAGGAACCTGGTAGGTACCTCTGAGGCCGAAAGAACAGCTGTGGAATGTGTCAGTTAGGGTGTG  
 101 GAAAGTCCCCAGGCTCCCAAGCAGGAGAAGTATGCAAAGCATGCATCTCAATTAGTCAGCAACCAGGTGTGAAAGTCCCAGGCTCCAGCAGGCAG  
 201 AAGTATGCAAAGCATGCATCTCAATTAGTCAGCAACCAGGTGTGAAAGTCCCAGGCTCCAGCAGGCAG  
 301 GAAGTTGGGGGAGGGTCGCAATTGAACGGGTGCCTAGAGAAGGTGGCGGGTAAACTGGAAAGTGATGTCGTACTGGCTCCGCTTTCCC  
 401 GAGGGTGGGGGAGAACCGTATATAAGTCAGTAGTCAGCTTTCGCAACGGTTGCGCCAGAACACAGCTGAAGCTCGAGGGCT  
 501 GCATCTCTCCTCACCGCCCCCGCCCTACCTGAGGCCCATCCACGCCGGTTGAGTCGCGTCTGCCGCTCCGCTGGTGCCTCTGAACTGC  
 601 GTCCGCCGTCTAGTAAGTTAAAGCTCAGGTCGAGACGGGCTTGTCCGGCCTCCCTGGAGCCTACCTAGACTCAGCCGCTCTCACGCTTGC

BspLU11I(799)

**AgeI (791)**

701 CTGACCCCTGCTTGTCAACTCTACGTCTTGTCTGCTGCG↓GTACAGATCCAAGCTGTGACCGGCGCTACCTGAGATCACCGTCAA

L5F (813)

801 CATGTTTGTGTTCTTIGTGTTGCTTCACTGGTCAGTCCCAATGCGTTAATCTCACCAACCGAACCTCCACCCGATATAACAAATTCTTCACC  
 1 M F V F F V U L L P L V S S Q C V N L T T R T Q L P P A Y T N S F T  
 901 AGAGGAGTGTACTATCCTGACAAAGTGTTCGGTCAAGTGTCTCCTCACTCTACTCAGGACCTTTCTGCTCTTCTAAGTGTACATGGTTCATG  
 34 R G V Y Y P D K V F R S S V L H S T Q D L F L P F F S N V T W F H

T95I (1083)

1001 CAATCCATGTGCTGGGACAAACGGCACCAACGGCTTCGACAACCCGTATTGCCATTCAATGATGGGTGTACTTGCCTCATGAGAAATCCAACAT  
 67 A I H V S G T N G T K R F D N P V L P F N D G V Y F A S I E K S N I  
 1101 CATTGAGGATGGATTTTGGACTACTCTGGACTCAAAGACACAGAGCCTGCTGATCGTTAACACGCCACAAACGTTGTCATCAAAGTGTGCGAATT  
 100 I R G W I F G T T L D S K T Q S L L I V N N A T N V V I K V C E F  
 1201 CAGTTTGCAATGATCCCTTCTGGAGTGTACTATCAGAAATAACAAGTCTGGATGGAGAGCGAACCTGGTCTACAGCAGCGAACAAACTGCA  
 134 Q F C N D P F L G V Y Y H K N N K S W M E S E F R V Y S S A N N C  
 1301 CCTTCGAGTACGTGAGTCACCCCTTCTGATGGACCTGGAAGGGAAACAGGGAAACTTCAAGAACCTGAGAGAGTTGTCTTTAAGAACATCGACGGCTA  
 167 T F E Y V S Q P F L M D L E G K Q G N F K N L R E F V F K N I D G Y  
 1401 TTTAAGATCTATAGTAAGCATACGCCATCACCTGGTAAGGGATCTCCCCAGGGCTTCAAGCCCTGGAACCTTGGTTGACTTGCTATTGGTATC  
 200 F K I Y S K H T P I N L V R D L P Q G F S A L E P L V D L P I G I

D253G (1557)

1501 AATATCACCAGATTTCAGACCCCTTGGCATTGCATGGTCTTATCTTACTCCAGGTGTTCTCTCCGGTGGACTGCCGCCGCTGCCTACTATG  
 234 N I T R F Q T L L A L H R S Y L T P G G S S S G W T A G A A A Y Y  
 1601 TCGGCTATCTGCAACCAAGAACGTTCTGCTCAAGTACAACGAAAACGGCACTATTACGGATGCTGTTGATTGCCCTGGACCCCTGTGAGACTAA  
 267 V G Y L Q P R T F L L K Y N E N G T I T D A V D C A L D P L S E T K  
 1701 ATGCACCCCTCAAGAGCTTACCGTGTGAGAAGGGATTACCAACAGTAATTCCGGTCAACCCACCGAACAGTGTGCGTTCCAAATATCACC  
 300 C T L K S F T V E K G I Y Q T S N F R V Q P T E S I V R F P N I T

D253G (1557)

1801 AATCTGTGCTCCCTTGGCGAAGTGTCAATGCTACAAGGTTGCTCTGTGTACGCATGGAATAGGAAACGCATCTCAAATTGTGTCGTTGATTACTCCG  
 334 N L C P F G E V F N A T R F A S V Y A W N R K R I S N C V A D Y S

E484K (2250)

1901 TGCTGTACAATTCCGCTTTCTCAACCTTCAAGTGTATGGCGTTCACCTACCAAACTTAACGACCTGTGCTTCAACTATGTTGATGCCGACTCTT  
 367 V L Y N S A S F S T F K C Y G V S P T K L N D L C F T N V Y A D S F

2001 TGTGATACGAGGCGATGAAGTGTGAGACAGATTGACCCAGGGCAGACGGCAAATTGCCACTACAACACTACAAGCTTCCAGATGACTTTACCGGATGTGTT  
 400 V I R G D E V R Q I A P G Q T G K I A D Y N Y K L P D D F T G C V

2101 ATTGCATGAACTCAAACAACTGGATTCCAAGGTGGTGGCAACTATAACTACCTGTATAGACTGTTAGGAAATCCAACCTGAAACCATCGAGCGAG  
 434 I A W N S N N L D S K V G G N Y N Y L Y R L F R K S N L K P F E R

D614G (2640)

2201 ATATAAGCACAGAAATCTACCAAGGCTGGAAAGTACGCCCTGCAACGGCGGAAAGGGTCAACTGCTACTTCCATTGAGGTTACGGATTCCAGCCTAC  
 467 D I S T E I Y Q A G S T P C N G V K G F N C Y F P L Q S Y G F Q P T

2301 AAACGGGGTGGTTACCAACCTATGTGCGTAGCTGAGTTGAGCTCTCCATGCCAGCCACAGTCTGTGGCCCAAGAAAAGCACCAATCTG  
 500 N G V G Y Q P Y R V V V L S F E L L H A P A T V C G P K K S T N L

2401 GTGAAGAACAAATCGTGAACCTTAACCTTAACGGACTCACAGGAACGGCGTATTGACGGAGAGTAACAAGAACGTTCTGCCATTCCAGCAGTCGGTC  
 534 V K N K C V N F N F N G L T G T G V L T E S N K K F L P F Q Q F G

2501 GCGATATTGCCACACTACCGACGCTGTCCGAGATCCCAGACATTGGAGATTCTGTATCACCCCTGTAGTTCCGGAGGTGAGCGTGTGATTACGCC  
 567 R D I A D T T D A V R D P Q T L E I L D I T P C S F G G V S V I T P

2601 CGGAACCAATACCGCAATCAGGTGCCGTCTGTATCAGGGTGAATTGACCGAGGTACCTGTCGCCATCACGCTGACCAACTACCCACATGG  
 600 G T N T S N Q V A V L Y Q G V N C T E V P V A I H A D Q L T P T W

2701 CGAGTATATTCCACCGGCTCAACGTCTTCAGACACGTGCTGGATGTCATCGGTGCAGAACACGTTAATAATAGCTACGAGTGTGATATCCCCATCG  
 634► R V Y S T G S N V F Q T R A G C L I G A E H V N N S Y E C D I P I  
 R685A (2853)  
 R683A (2847)  
 2801 GTGCTGGAATATGCGCCTTATCAAACCTAAACCAACTCTCCTAGGGCGGCAGCTAGTGTAGCATCCAAAGTATCATTGCCACACAATGAGCCTCGG  
 667► G A G I C A S Y Q T Q T N S P R A A A S V A S Q S I I A Y T M S L G  
**A701V (2901)**  
 2901 TGTAGAGAATTCTGTCGCCTACAGCAACAACCTCATTGCTATCCCTACTAACTTCACAATCAGTGTGACAACGTGAAATTCTGCCGTATCTGACCAAA  
 700► V E N S V A Y S N N S I A I P T N F T I S V T T E I L P V S M T K  
 3001 ACAAGCGTTGACTGCACCATGTCATCTGTCGGGATTCTACCGAATGTCAGCAATCTCCCTGCACAGGATCATTGCACTCAGCTGAATCGTGCC  
 734► T S V D C T M Y I C G D S T E C S N L L L Q Y G S F C T Q L N R A  
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 834► I K Q Y G D C L G D I A A R D L I C A Q K F N G L T V L P P L L T  
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 3501 GCAGATGGCCTACCGCTTAAACGGCATCGGTGACACAAAACGTTCTGTATGAAAACCAGAAACTCATGCCAACCGAGTTCAACAGTGTATCGTAAG  
 900► Q M A Y R F N G I G V T Q N V L Y E N Q K L I A N Q F N S A I G K  
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 1100► H W F V T Q R N F Y E P Q I I T T D N T F V S G N C D V V I G I V  
 4201 AATAACTGTTACGATCCTTGAGCCAGAGCTGGACTCCTCAAGGAGGAGCTTGACAAATATTAAAGAATCACACATCACCTGACGTCGACCTCG  
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 1200► Q E L G K Y E Q Y I K W P W Y I W L G F I A G L I A I V M V T I M  
**NheI (4567)**  
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 4601 GAGTTGGACAAACCAACTAGAATGCACTGAAAAAATGCTTATTGTGAAATTGTGATGCTATTGCTTATTGTAACCATTAAAGCTGCAATA  
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 141► • N R T Y K L P I L E E I T T K V L K G N M E I L V F C D P  
 5301 AGCATAGTCAGAGATGAGCTCTGTCACATGCCACAGGGCTGACCCCTGATGGATCTGCCACCTCATCAGAGTAGGGGTGCTGACGCCAACATG  
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 ←

5901 AGTCAAACCGCTATCCACGCCATTGATGACTGCCAAAACCGCATCATGGTAATAGCGATGACTAATACGTAGATGTACTGCCAAGTAGGAAAGTC  
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BspLU11I (6271)

