

# pUNO1-hSTING-H232

Expression vector containing H232 isoform human STING (R232H) open reading frame

Catalog code: puno1-hsting-h232

<https://www.invivogen.com/hsting-h232>

## For research use only

Version 19K10-MM

## PRODUCT INFORMATION

### Contents

- 20 µg of lyophilized plasmid DNA
- 2 x 1 ml blasticidin at 10 mg/ml

### Storage and Stability

- Product is shipped at room temperature.
- Lyophilized DNA should be stored at -20°C.
- Resuspended DNA should be stored at -20°C and is stable at least for 1 year.
- Store blasticidin 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 full-length open reading frame (ORF) sequencing.
- Plasmid DNA was purified by ion exchange chromatography.

## GENERAL PRODUCT USE

- **Subclone gene into another vector.** Two unique restriction sites flank the gene, allowing convenient excision. The 5' site is BspEI which is compatible with AgeI, XmaI, NgoMIV and SgrAI. The 3' site is NheI which is compatible with XbaI, SpeI, and AvrII.
- **Stable gene expression in mammalian cells.** pUNO1 plasmids can be used directly in transfection experiments both *in vitro* and *in vivo*. pUNO1 plasmids contain the blasticidin-resistance gene (*bsr*) driven by the CMV promoter/enhancer in tandem with the bacterial EM7 promoter. This allows the amplification of the plasmid in *E. coli*, as well as the selection of stable clones in mammalian cells using the same selective antibiotic. pUNO1 allows high levels of expression and secretion of the gene product.

## 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 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 at 10 mg/ml in HEPES buffer.

## PLASMID FEATURES

- **Bsr (blasticidin resistance gene):** The *bsr* gene from *Bacillus cereus* encodes a deaminase that confers resistance to the antibiotic blasticidin. The *bsr* gene is driven by the CMV promoter/enhancer in tandem with the bacterial EM7 promoter. Therefore, blasticidin can be used to select stable mammalian cells transfectedants and *E. coli* transformants.
- **CMV promoter & enhancer** drives the expression of the blasticidin resistance in mammalian cells.

### • Human STING-R232H

**ORF size:** 1140 bp

**Cloning fragment size:** 1181 bp

STING (stimulator of interferon genes; also known as TMEM173, MITA, MPYS, and ERIS) is essential for the IFN response to microbial or self-DNA, and acts as a direct sensor of cyclic dinucleotides (CDNs). CDNs are important messengers in bacteria, affecting numerous responses of the prokaryotic cell, but also in mammalian cells, acting as agonists of the innate immune response. Several non-synonymous variants of STING have been described in the human population. R232H has been identified as a natural variant allele of STING occurring in ~14% of the human population<sup>1</sup>. H232 contains a single amino acid substitution R232H. The R232H isoform has a diminished response to bacterial and metazoan CDNs when compared to the wild-type allele<sup>1-2</sup>. R232H has been the most commonly used human STING allele in published structural studies.

• **EF-1 $\alpha$ /HTLV hybrid promoter** is a composite promoter comprised of the Elongation Factor-1 $\alpha$  (EF-1 $\alpha$ ) core promoter<sup>3</sup> and the 5' untranslated region of the Human T-Cell Leukemia Virus (HTLV). EF-1 $\alpha$  utilizes a type 2 promoter that encodes for a «house keeping» gene. It is expressed at high levels in all cell cycles and lower levels during G0 phase. The promoter is also non-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>4</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 possibly through mRNA stabilization.

• **SV40 pAn:** The Simian Virus 40 late polyadenylation signal enables efficient cleavage and polyadenylation reactions, resulting in high levels of steady-state mRNA<sup>5</sup>.

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

• **Human beta-Globin polyA** is a strong polyadenylation (pAn) signal placed downstream of *bsr*. The use of beta-globin pAn minimizes interference<sup>6</sup> and possible recombination events with the SV40 polyadenylation signal.

1. Yi G. et al., 2013. Single nucleotide polymorphisms of human STING can affect Innate immune response to cyclic dinucleotides. PLoS One 8(10):e77846. 2. Diner E. et al., 2013. The innate immune gene sensor cGAS produces a noncanonical cyclic dinucleotide that activates human STING. Cell Rep 3(5):1355-61. 3. 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. 4. 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. 5. Carswell S. & Alwine J., 1989. Efficiency of utilization of the simian virus 40 late polyadenylation site: effects of upstream sequences. Mol Cell Biol. 9(10):4248-58. 6. 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.

## RELATED PRODUCTS

Product	Description	Cat. Code
Blasticidin ChemiComp GT116	Selection antibiotic Competent <i>E. coli</i>	ant-bl-1 gt116-11

### TECHNICAL SUPPORT

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

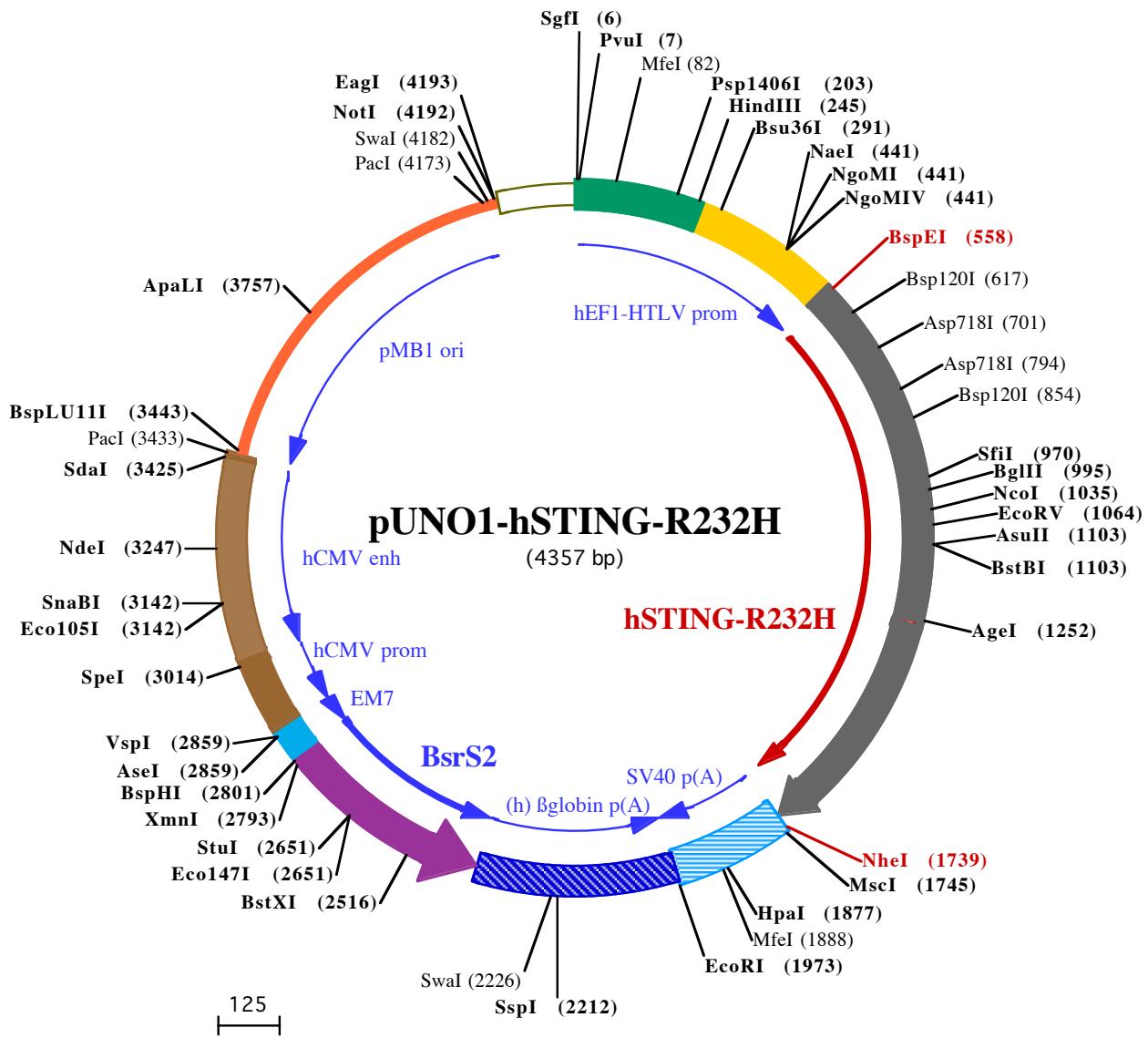
InvivoGen USA (International): +1 (858) 457-5873

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E-mail: info@invivogen.com





**PvuI** (7)  
**SgfI** (6)  
 1 GGATCTCGATCGTCCGGTCCCCGTCACTGGGAGAGCGCACATGCCACAGTCCCCGAGAAGTTGGGGGAGGGGTCGGCAATTGAACGGTGCTA  
 101 GAGAAGGGTGGCAGGGGTAACCTGGAAAGTGTGCGTGTACTGGCTCGCTTTCCGAGGGTGGGGAGAACGTATAAGTCAGTAGTCGGC  
 202 Psp1406I (203) HindIII (245) Bsu36I (291)  
 203 TGAACGTTTCGCAACGGGTTGCCAGAACACAGCTGAAGCTCGAGGGGCTCGCATCTCCTCACGCCGCCCTACCTGAGGCCGC  
 303 CATCCACGCCGGTTGAGTCGCTCTGCCGCCCTCCGCTGTGGTGCCTCTGAACCTCGCTCCACGCTTGCCTGACCTGCTCAACTCTACGTCTTGTTCGTTCTG  
 404 CCTTGTCCGGCGCTCCCTGGAGCCTACCTAGACTCAGCCGGCTCCACGCTTGCCTGACCTGCTGCTCAACTCTACGTCTTGTTCGTTCTG  
 505 BspEI (558)  
 505 TTCTCGCCGTTACAGATCCAAGCTGTGACCGGGCGCTACCTGAGATCACCGGCTCCGGACAGCATGCCACTCCAGCCTGCATCCATCCCCTGTC  
 606 Bsp120I (617) Asp718I (701)  
 606 CCAGGGGTACGGGCCAGAACGGCAGCCTGGTCTGACTGAGTCCTGCTGGTACCCCTGGGGCTAGGAGAGCCACAGAGCACACTCTGGTAC  
 13▶ P R G H G A Q K A A L V L L S A C L V T L W G L G E P P E H T L R Y  
 707 Asp718I (794)  
 707 CTGGTCCTCACCTAGCCTCCCTGAGCTGGGACTGCTGTTAACGGGCTGAGCCTGGCTGGAGGCTGCCACATCCACTCAGGTACCGGGCA  
 47▶ L V L H L A S L Q L G L L L N G V C S L A E E L R H I H S R Y R G  
 807 Bsp120I (854)  
 807 GCTACTGGAGGACTGTGGGGCTGCCTGGCTGCCCTCCGGCTGGGCTCTGGGCTGTGCTCCATATTCTACTACTCCCTCCAAATCGGGTC  
 80▶ S Y W R T V R A C L G C P L R R G A L L L S I Y F Y Y S L P N A V  
 908 BgIII (995)  
 908 908 GGCCGCCCTTCACTGGATGCTGGCTCTGGGCTCTCGCAGGCACTGAACATCTCTGGGCTCAAGGGCCTGGCCCAGCTGAGATCTCGAGT  
 114▶ G P P F T W M L A L L G L S Q A L N I L L G L K G L A P A E I S A V  
 1009 BstBI (1103)  
 1009 NeoI (1035) EcoRV (1064) AsuII (1103)  
 147▶ C E K G N F N V A H G L A W S Y Y I G Y L R L I L P E L Q A R I R  
 1110 CTTACAATCAGATTACAACAACCTGCTACGGGTGAGTGAGGCAAGCCTGAGATCTGGGACTGTGGGGTGCCTGATAACCTGAGTATG  
 181▶ T Y N Q H Y N N L L R G A V S Q R L Y I L L P L D C G V P D N L S M  
 1211 AgeI (1252)  
 1211 GCTACCCCCAACATTGCTTCTGGATAACTGGCCAGCAGACGGTACCATGCTGGCATCAAGGATGGTTACAGAACAGCATCTAGAGCTCT  
 215▶ A D P N I R F L D K L P Q T G D H A G I K D R V Y S N S I Y E L L  
 1312 1312 GGAGAACGGCAGCGGGCGGGCACCTGTCCTGGAGTACGCCACCCCCCTGAGACTTGTGTCATGTCACAATACAGTCAGCTGGCTTAGCCGG  
 248▶ E N G Q R A G T C V L E Y A T P L Q T L F A M S Q Y S Q A G F S R  
 1413 1413 AGGATAGGCTTGAGCAGGCAAACCTCTTGCCGACACTGAGGACATCTGGAGATGCCCTGAGTCAGAACACTGCCCTATTGCCCTACAG  
 282▶ E D R L E Q A K L F C T R L E D I L A D A P E S Q N N C R L I A Y Q  
 1514 1514 GAACCTGCAAGATGACAGCAGCTTCTGCTGCCAGGAGTTCTCGGACCTGGCAGGAGAAAAGGAAGAGTTACTGTGGGAGCTGAAGACCTC  
 316▶ E P A D D S S F S L S Q E V L R H L R Q E E K E E V T V G S L K T S  
 1615 1615 AGCGTGCCTCAGTACCTCACGATGCCAACAGAGCTGAGCTCATAGTGAATGAAAAGCCCTCCCTCCGACGGATTCTCTGAGACCCAG  
 349▶ A V P S T S T M S Q E P E L L I S G M E K P L P L R T D F S •  
 1716 MscI (1745)  
 1716 NheI (1739)  
 1817 HpaI (1877) MfeI (1888)  
 1918 EcoRI (1973)  
 1918 AGGGGAGGTGGAGGTTTTAAAGCAAGTAAACCTCTACAAATGGTATGGAATTCTAAACATAGCAAAACTTAACTCCAAATCAAG  
 2019 CCTCTACTGAATCTTCTGAGGGATGAATAAGGCATAGGCATAGGGCTGTCATGTCATTAGCTGTTGCAGCTCACCTTTCATGGAA  
 2119 SspI (2212)  
 2119 GTTTAAGATATAGTGTATTTCCAAGGTTGAACAGCTCTTCTTATGTTAAATGCACTGACCTCCACATCCCTTTAGTAAATATT  
 2220 SwI (2226)  
 2220 AGAAATAATTAAACATCATTGCAATGAAATAATGTTTATTAGGCAGAACATCCAGATGCTCAAGGCCCTCATATAATCCCCAGTTAGTTAGTT  
 2321 GGACTTAGGAACAAAGGAACCTTAATAGAAATTGGACAGCAAGAACGGAGCTCTAGCTTGTAGTTCTGGTACTTGAGGGGATGAGTTCTCAAT  
 1411 • N R T Y K L P I L E E I  
 2422 BstXI (2516)  
 2422 GGTGGTTTGACCAGCTGCCATTCTCAATGAGCACAAAGCAGTCAGGAGCATAGTCAGAGATGAGCTCTGCACATGCCACAGGGCTGACCA  
 128▶ T K V L K G N M E I L V F C D P A Y D S I L E R C M G C P S V V R  
 2523 TGATGGATCTGCAACCTCATCAGAGTAGGGGCTGAGCAGCCACAATGGTCAAAGTCTCTGCCCCGGTGCACAGCAGACCAATGGCAATGGCT  
 94▶ I S R D V E D S Y P H R V A V I T D F D K Q G N S V A S G I A I A  
 2624 StuI (2651)  
 2624 Eco147I (2651)  
 2624 TCAGCACAGACAGTGCACCTGCCAATGTTAGGCCCTCAATGGAGCAGCAGAGATGATCTCCCAAGTCTGGTCTGATGGCCGCCGACATGGGCTTGT  
 60▶ E A C V T V R G I Y A E I H V A S I I E G T K T R I A A A G V H H K N  
 2725 BspHI (2801)  
 2725 XmnI (2793)  
 2725 GTCCTCATAGAGCATGGTATCTCTCAGTGGCACCTCCACAGCTCCAGATCTGCTGAGAGATGTTGAAGGTCTTCATGATGGCCCTCTATAGTGA  
 27▶ D E Y L M T I K E T A V E V L E L D Q Q S I N F T K M ←  
 2826 VspI (2859)  
 2826 AseI (2859)  
 2826 TCGTATTATACTATGCCGATATACTATGCCGATATTGTCAAACAGCGTGGATGGCGTCTCCAGCTTATCTGACGGTCACTAAACGAGCTG  
 2926 SpeI (3014)  
 2926 TTATATAGACCTCCCACCGTACACGCCCTACCGCCATTGCGTAATGGGGCGAGTTGTTACGACATTGAAAGTCCGTTGATTTACTAGTCAAAA  
 3026 CAAACCTCCATTGACGTCAATGGGGAGACTTGGAAATCCCCGTGAGTCAAACCGCTATCCACGCCATTGATGACTGCCAAACCGCATCATG

**SnaBI** (3142)  
**Eco105I** (3142)

3126 **GTAATAGCGATGACTAATACGTAGATGACTGCCAAGTAGGAAAGTCCCATAAGTCATGTACTGGGCATAATGCCAGGCAGGCCATTACCGTCATTGAC**

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**NdeI** (3247)

3227 **GTCAATAGGGGGCGTACTTGGCATATGATACACTTGATGACTGCCAAGTGGCAGTTACCGTAATACTCCACCCATTGACGTCAATGGAAGTCCCTA**

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**SdaI** (3425)

3328 **TTGGCGTTACTATGGAACATACGTCAATTGACGTCAATGGCGGGGTCGGTGGCGGTAGCCAGGCAGGCCATTACCGTAAGTTATGTAACGCC**

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PacI (3433)    **BspLU11I** (3443)

3428 **TGCAG G TT AA TT** AAGAACATGTGAGCAAAGGCCAGCAAAGGCCAGGAACCGTAAAAGGCCGCGTGTGGCGTTTTCCATAGGCTCCGCCCCCT  
3527 **GACGAGCATCACAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGGACTATAAAGATACCAGGCCTTCCCCCTGAAAGCTCCCTCGCGCTC**

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3628 **TCCTGTTCCGACCCCTGCGCTTACCGATACTGTCCGCTTCCCTCGGAAAGCGTGGCGCTTCTAGCTCACGCTGTAGGTATCTCAGTCGG**

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**ApaLI** (3757)

3729 **TGTAGGTCGTCGCTCCAAGCTGGGCTGTGACGAACCCCCCGTTAGCCCCGACCGCTGCCCTATCGGTAACTATGCTTTGAGTCCAACCCGTA**

---

3830 **AGACACGACTTATGCCACTGGCAGCAGCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCCTGCTACAGAGTTCTGAAGTGGTGGCTAACTA**

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3931 **CGGCTACACTAGAAGAACAGTATTGGTATCTGCCTCTGCTGAAGCCAGTTACCTCGAAAAAGAGTTGGTAGCTCTTGATCCGCAAACAAACCCG**

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4032 **CTGGTAGCGGTGGTTTTTGTGCAAGCAGATTACGCGCAGAAAAAAAGGATCTCAAGAAGATCCTTGATCTTCTACGGGTCTGACGCTCAG**

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**EagI** (4193)

PacI (4173)    Swal (4182)    **NotI** (4192)

4133 **TGGAACGAAAACTCACGTTAAGGGATTTGGTATGGCTAGTTAAATTAAACATTAAATC AGCGGCCAATAAAATATCTTATTTCATTACATCTGT**

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4233 **TGTTGGTTTTTGTGAATCGTAACATACGCTCTCCATAAAACAAAACGAAACAAACAAACTAGCAAAATAGGCTGCCCCAGTGCAAGTGCAG**

4334 **GTCAGAACATTCTATCGAA**