

Product datasheet for **MC229435**

Hipk2 (NM_001294144) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Hipk2 (NM_001294144) Mouse Untagged Clone
Tag: Tag Free
Symbol: Hipk2
Synonyms: 1110014O20Rik; B230339E18Rik; Stank
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC229435 representing NM_001294144
Red=Cloning site **Blue**=ORF **Orange**=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
GCC**CGGATCGCC**

ATGGCCCCGTGTACGAAGGTATGGCCTCACATGTGCAAGTTTTCTCCCCTCACACCCTCAATCAAGTG
CCTTCTGTAGTGTGAAGAACTAAAAGTAGAGCCAAGTCCAACCTGGGACATGACTGGGTACGGCTCCCA
CAGCAAAGTGTACAGCCAGAGCAAGAACATACCACCTTCTCAGCCAGCCTCCACAACCGTCAGCACCTCC
TTGCCAATCCCCAACCCAAGCCTACCTTACGAGCAGACCATCATCTTCCCAGGAAGTACTGGACACATAG
TTGTAACATCAGCAAGTAGCACTTCTGTACCCGGCAAGTCTCGGTGGACCACATAACCTAATGCGTCCG
AAGCACTGTGAGCCTCCTTGACACCTACCAAAAATGTGGACTCAAGCGCAAGAGTGAGGAGATTGAGAAC
ACAAGCAGCGTGCAGATCATTGAAGAGCATCCACCCATGATTCAGAATAATGCCAGCGGGGCCACTGTAG
CCTACTGCCACCACATCCACTGCCACCTCCAAAAACAGTGGCTCCAACAGTGAAGGGGACTATCAGCTGGT
GCAGCATGAAGTACTGTGCTCCATGACCAACACCTATGAGGTTTTAGAGTTCTTGGGGCGAGGGACATTT
GGGCAAGTGGTCAAGTGTGAAACGGGGCACCATGAAATTGTGGCCATTAAGATCCTCAAGAACCACC
CCTCCTATGCCCGGCAAGGCCAGATTGAAGTGAGCATCTTGGCCGGCTGAGCAGCGAGAGCGCTGATGA
CTACAATTTGTGCGGGCTATGAGTGCTTCCAGCACAAGAACCACACGTGCTTAGTCTTTGAGATGTTG
GAGCAGAACCCTTATGACTTTCTGAAACAGAACAAGTTTGTCCCTTGCCCTCAAGTATATTGCCCCAG
TCCTCCAGCAGGTAGCCACAGCCCTGATGAAACTCAAAAGCCTGGGTCTTATCCATGCTGACCTCAAACC
AGAAAACATCATGCTGGTGGATCCATCCAGACAACCGTACCGAGTGAAGGTCAATTGACTTTGGTTACGCT
AGTCATGTGTCCAAGCTGTCTGCTCTACGTACTTGAATCCAGATACTACCGGGCCCTGAGATCATCC
TTGGTTTGCCATTCTGTGAGGCAATTGACATGTGGTCCCTGGTTGTGTCATTGCTGAAGTGTCTCTGGG
CTGGCCATTATACCCAGGTGCTTCTGAGTATGATCAGATTCGGTATATTTCAAAAACACAGGGTTTACCA
GCTGAATATTTAAGTGCAGGGACAAAGACAAGTGGTTTTTCAACCGTGACACAGACTACCCGTATC
CTTTGTGAGGGCTAAAGACACCAGATGACCATGAAGCAGAGACGGGAATTAAGTCAAAGGAAGCAAGAAA
GTACATTTTCAACTGTTTGGATGATATGGCCAGGTGAACATGACAACAGATTTGGAAGGGAGTGACATG
TTAGTAGAGAAGGCAGACCCGACGGGAGTTCATTGACCTGTTAAAGAAGATGCTGACCATCGATGCTGATA



AGAGAGTCACTCCCATTGAAACTCTGAACCACCCCTTTGTACCATGACACACCTGCTTGACTTCCCCCA
 TAGTGCCCATGTCAAATCTTGTTTCCAAAACATGGAGATTTGCAAGCGCCGGGTGAATATGTATGACACA
 GTGAACCAGAGCAAAACACCTTTTCATCACTCACGTGGCCCCCAGCACATCCACCAACTTGACCATGACCT
 TTAACAACCAGCTGACCACTGTCCACAACCAGCCCTCAGCGGCATCCATGGCCCGGTGGCCCCGGGAG
 CATGCCCTGCAGACGGGAACAGCCAAATTTGTGCCGACCTGATCCCTCCAGCAAGCTCTCATCGTG
 TGTCCCCTGGCTTCCAAGGCCGTGACGGCCTCTCCCTCCAAGCAGCTGGCTACTCAGTGGCGATGGAAA
 ATGCTGTCCCCATCGTCACCCAGGCGCCAGGAGCTCAGCCTCTTCCAGATCCAACCAAGGCTGCTTCA
 GGCTGGCCAGGTGGGGCCCAACAGATCCTACTTCTCCAGCATGGCAGCAGCTGACTGGCGTGGCCACC
 CACACATCTGTACAGCATGCAGCTGTGATTCTGAGACCATGGCAGGCACTCAGCAACTGGCTGACTGGA
 GGAACACGCATGCTCATGGAAGTCATTACAATCCCATCATGCAGCAGCCTGCACTCTTGACTGGTATGT
 GACCTTCCAGCTGCCAGCCCTTAAATGTGGGTGTAGCCCATGTGATGAGGCAACAACCAACCAGCACC
 ACCTCCTCCAGGAAGAGTAAGCAGCACCAGTCTCTGTGAGAAATGTCTCCACCTGTGAGGTGACCTCTT
 CACAGGCTATCAGCTCCCCTCAGCGATCCAAGCGTGTCAAGGAGAACACTCCCCACGGTGCATGGT
 ACACAGCAGCCAGCTTGCAGCACCTCAGTCACCTGTGGGTGGGCGACGTGGCCTCCAGCACCACCGG
 GAGCGACAGCGCAGACGATTGTATCCCCGACACCCGAGCCCCACAGTCAGTGTATCACCATCAGCA
 GTGACACCGATGAAGAAGAGGAGCAGAAGCAGCCCCCACCAGCAGATCTCCAAGCAAAGAAAAATGT
 CATCAGCTGTGTACCGTCCACGACTCTCCCTACTCTGACTCCTCCAGCAACACCAGCCCTACTCGGTG
 CAGCAGCGCACAGGGCACAACGGCACCAACACCTTGGACACCAAGGGGGCCCTAGAGAATCACTGCACTG
 GCAACCCCGCACCATCATCGTACCCCACTGAAGACCCAGGCCAGTGAAGTGTGGTAGAATGTGACAG
 CCTAGGGCCAGCGATCAGTGCCAGTCACCATTCGTCTCTTCAAGTCCAAGTCTCCAGCACCCTGACC
 TCCACCACTGGACTCTTCAGGGAGCTTTCAGGAGCCATCGCTACCGTCCAGCAGCGCCAGGCCCC
 ACTTCCAGCAGCAGCAGCCCTCAATCTCAGCCAGGCCAGCAGCACATGGCTGCGGACCCGACCCGGGAG
 TCACCGTCGACAGCAGGCTACATCACTCCGACCATGGCGCAAGCTCCGTACACCTTCCCGCAACAGC
 CCCAGCCATGGCACTGTTCAACCCACCTGGCTGCGGCTGCCACCTCCCACCCAGCCTCACCTCTACA
 CCTATACAGCGCCACAGCCCTAGGCTCCACCGCACTGTAGCCACCTGGTGGCATCCCAGGCTCAGC
 ACGCCACACCGTGCAGCACACTGCCTATCCGGCCAGCATCGTCCACCAGGTCCCTGTGAGCATGGGGCC
 CGGGTCTGCCCTCGCCACCATCCACCCAGTCAGTATCCAGCCAGTTTGCCACCCAGACCTATATCA
 GCGCCTCGCCAGCCTCCACCGTCTACACTGGATACCCACTGAGTCTGCCAAGGTCAACCAATACCTTA
 CATATAA

ACGCGTACGCGGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001294144
- Insert Size:** 3507 bp
- OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
- OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal tag.
- Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

Reconstitution Method:

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM_001294144.1](#), [NP_001281073.1](#)

RefSeq Size: 4175 bp

RefSeq ORF: 3507 bp

Locus ID: 15258

UniProt ID: [Q9QZR5](#)

Cytogenetics: 6 B1

Gene Summary:

Serine/threonine-protein kinase involved in transcription regulation, p53/TP53-mediated cellular apoptosis and regulation of the cell cycle. Acts as a corepressor of several transcription factors, including SMAD1 and POU4F1/Brn3a and probably NK homeodomain transcription factors. Phosphorylates PDX1, ATF1, PML, p53/TP53, CREB1, CTBP1, CBX4, RUNX1, EP300, CTNNB1, HMGA1 and ZBTB4. Inhibits cell growth and promotes apoptosis through the activation of p53/TP53 both at the transcription level and at the protein level (by phosphorylation and indirect acetylation). The phosphorylation of p53/TP53 may be mediated by a p53/TP53-HIPK2-AXIN1 complex. Involved in the response to hypoxia by acting as a transcriptional co-suppressor of HIF1A. Mediates transcriptional activation of TP73. In response to TGF β , cooperates with DAXX to activate JNK. Negative regulator through phosphorylation and subsequent proteasomal degradation of CTNNB1 and the antiapoptotic factor CTBP1. In the Wnt/beta-catenin signaling pathway acts as an intermediate kinase between MAP3K7/TAK1 and NLK to promote the proteasomal degradation of MYB. Phosphorylates CBX4 upon DNA damage and promotes its E3 SUMO-protein ligase activity. Activates CREB1 and ATF1 transcription factors by phosphorylation in response to genotoxic stress. In response to DNA damage, stabilizes PML by phosphorylation. PML, HIPK2 and FBXO3 may act synergically to activate p53/TP53-dependent transactivation. Promotes angiogenesis, and is involved in erythroid differentiation, especially during fetal liver erythropoiesis. Phosphorylation of RUNX1 and EP300 stimulates EP300 transcription regulation activity. Triggers ZBTB4 protein degradation in response to DNA damage. Modulates HMGA1 DNA-binding affinity. In response to high glucose, triggers phosphorylation-mediated subnuclear localization shifting of PDX1. Involved in the regulation of eye size, lens formation and retinal lamination during late embryogenesis. [UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (4) uses two alternate in-frame splice sites in the coding region compared to variant 2. The encoded isoform (4) is shorter than isoform 2. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on alignments.