

## Product datasheet for MC223911

### Hipk2 (NM\_010433) Mouse Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Hipk2 (NM_010433) 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:	>MC223911 representing NM_010433 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCCGCGATCGCC

ATGGCCCCGTGTACGAAGGTATGGCCTCACATGTGCAAGTTTTCTCCCCTCACACCCTCAATCAAGTG  
CCTTCTGTAGTGTGAAGAACTAAAAGTAGAGCCAAGTCCAACCTGGGACATGACTGGGTACGGCTCCCA  
CAGCAAAGTGTACAGCCAGAGCAAGAACATACCACCTTCTCAGCCAGCCTCCACAACCGTCAGCACCTCC  
TTGCCAATCCCCAACCCAAGCCTACCTTACGAGCAGACCATCATCTTCCCAGGAAGTACTGGACACATAG  
TTGTAACATCAGCAAGTAGCACTTCTGTACCCGGCAAGTCTCGGTGGACCACATAACCTAATGCGTCCG  
AAGCACTGTGAGCCTCCTTGACACCTACCAAAAATGTGGACTCAAGCGCAAGAGTGAGGAGATTGAGAAC  
ACAAGCAGCGTGCAGATCATTGAAGAGCATCCACCCATGATTCAGAATAATGCCAGCGGGGCCACTGTAG  
CCTACTGCCACCACATCCACTGCCACCTCCAAAAACAGTGGCTCCAACAGTGAAGGGGACTATCAGCTGGT  
GCAGCATGAAGTACTGTGCTCCATGACCAACACCTATGAGGTTTTAGAGTTCTTGGGGCGAGGGACATTT  
GGGCAAGTGGTCAAGTGTGAAACGGGGCACCATGAAATTGTGGCCATTAAGATCCTCAAGAACCACC  
CCTCCTATGCCCGCAAGGCCAGATTGAAGTGAGCATCTTGGCCGGCTGAGCAGCGAGAGCGCTGATGA  
CTACAATTTGTGCGGGCTATGAGTGCTTCCAGCACAAGAACCACACGTGCTTAGTCTTTGAGATGTTG  
GAGCAGAACCCTTATGACTTTCTGAAACAGAACAAGTTTGTCCCTTGCCCTCAAGTATATTGCCCCAG  
TCCTCCAGCAGGTAGCCACAGCCCTGATGAAACTCAAAAGCCTGGGTCTTATCCATGCTGACCTCAAACC  
AGAAAACATCATGCTGGTGGATCCATCCAGACAACCGTACCGAGTGAAGGTCAATTGACTTTGGTTACGCT  
AGTCATGTGTCCAAAGCTGTCTGCTCTACGTACTTGAATCCAGATACTACCGGGCCCTGAGATCATCC  
TTGGTTTGCCATTCTGTGAGGCAATTGACATGTGGTCCCTGGTTGTGTCATTGCTGAAGTGTCTCTGGG  
CTGGCCATTATACCCAGGTGCTTCTGAGTATGATCAGATTCGGTATATTTCAAAAACACAGGGTTTACCA  
GCTGAATATTTAAGTGCAGGACAAAGACAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAGTAAAG  
CCTTGTGGAGGCTAAAGACACCAGATGACCATGAAGCAGAGACGGGAATTAAGTCAAAAGGAAGCAAGAAA  
GTACATTTTCAACTGTTTGGATGATATGGCCAGGTGAACATGACAACAGATTTGGAAGGGAGTGACATG  
TTAGTAGAGAAGGCAGACCCGACGGGAGTTCATTGACCTGTTAAAGAAGATGCTGACCATCGATGCTGATA



AGAGAGTCACTCCCATTGAAACTCTGAACCACCCCTTTGTCACCATGACACACCTGCTTGACTTCCCCCA  
 TAGTGCCCATGTCAAATCTTGTTTCAAACCATGGAGATTTGCAAGCGCCGGGTGAATATGTATGACACA  
 GTGAACCAGAGCAAAACACCTTTTCATCACTCACGTGGCCCCAGCACATCCACCAACTTGACCATGACCT  
 TTAACAACCAGCTGACCACTGTCCACAACCAGGCTCCCACCACCTCCAGTGCCACTCTTTCTTAGCCAA  
 TCCCGAAGTCTCCATACTAACTACCAATCTGCACTCTACCAGCCCTCAGCGGCATCCATGGCCGCGGTG  
 GCCCGCGGAGCATGCCCTGCAGACGGGAACAGCCAAATTTGTGCCGACCTGATCCCTCCAGCAAG  
 CTCTCATCGTGTGTCCCCTGGCTTCCAAGGCCTGCAGGCCTCCTCCCAAGCACGCTGGCTACTCAGT  
 GCGGATGGAAAATGCTGTCCCCATCGTACCCAGGCGCCAGGAGCTCAGCCTCTTCAGATCCAACCAGGC  
 CTGCTTGCTCAGCAGGCCTGGCCAGGTGGGGCCCAACAGATCCTACTTCTCCAGCATGGCAGCAGTGA  
 CTGGCGTGGCCACCCACACATCTGTACAGCATGCAGCTGTGATTCTGAGACCATGGCAGGCACTCAGCA  
 ACTGGCTGACTGGAGAACACGCATGCTCATGGAAGTCATTACAATCCCATCATGCAGCAGCCTGCACTC  
 TTGACTGGTCATGTGACCCTTCCAGCTGCCAGCCCTTAAATGTGGGTGTAGCCCATGTGATGAGGCAAC  
 AACCAACCAGCACCCTCCTCCAGGAAGAGTAAGCAGCACCAGTCATCTGTGAGAAATGTCTCCACCTG  
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 CGGTGCGCCATGGTACACAGCAGCCAGCTTGCAGCACCTCAGTCACCTGTGGGTGGGGCAGCTGGCCT  
 CCAGCACCCCGGGAGCGACAGCGGCAGACGATTGTATCCCCGACACCCCGAGCCCCACAGTCAGTGT  
 CATCACCATCAGCAGTGACACCGATGAAGAAGAGGAGCAGAAGCAGCCCCACCAGCACAGTCTCCAAG  
 CAAAGAAAAATGTCATCAGCTGTGTACCCTCCAGCTCTCCCTACTCTGACTCCTCCAGCAACACCA  
 GCCCTACTCGGTGCAGCAGCGCACAGGGCACAACGGCACCAACACCTTGGACACCAAGGGGGGCTAGA  
 GAATCACTGCACTGGCAACCCCGCACCATCATCGTACCCCACTGAAGACCCAGGCCAGTGAAGTGTG  
 GTAGAATGTGACAGCCTAGGGCCAGCGATCAGTGCCAGTACCATTCTGCTCCTCTCAAGTCCAAGTCT  
 CCAGCACCGTGACCTCCACCAGTGGACACTTTCAGGGAGCTCTCAGGAGCCATCGCCTACCGTCAGCA  
 GCGGCCAGGCCCCACTTCCAGCAGCAGCAGCCCTCAATCTCAGCCAGGCCAGCAGCATGGCTGCG  
 GACCGCACCGGGAGTACCGTGCAGCAGGCGCTACATCACTCCGACCATGGCGCAAGTCCGTACACCT  
 TCCCGCACAAACAGCCCCAGCCATGGCACTGTTCACCCCACTGGCTGCGGCTGCCACCTCCCCACCA  
 GCCTCACCTCTACACCTATACAGCGCCACAGCCCTAGGCTCCACCGGCACTGTAGCCCACTGGTGGCA  
 TCCCAAGGCTCAGCAGCCACACCGTGCAGCAGTGCCTATCCGGCCAGCATCGTCCACCAGGTCCTG  
 TGAGCATGGGGCCCCGGTCTGCCCTCGCCACCATCCACCCAGTCAGTATCCAGCCAGTTTGGCCA  
 CCAGACCTATATCAGCGCTCGCCAGCCTCCACCGTCTACTGGATACCCACTGAGTCTGCCAAGGTC  
 AACCAATACCCTTACATA

ACGCGTACGCGGCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT  
 ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:**

Sgfl-Mlul

**ACCN:**

NM\_010433

**Insert Size:**

3591 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).

**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\\_010433.2](#), [NP\\_034563.2](#)

**RefSeq Size:** 4259 bp

**RefSeq ORF:** 3591 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 $\beta$ , 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 (2) represents the longest transcript and encodes the longest isoform (2). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.