

Product datasheet for TP527349

Hipk2 (NM_001136065) Mouse Recombinant Protein

Product data:

OriGene Technologies, Inc.

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Product Type:	Recombinant Proteins
Description:	Purified recombinant protein of Mouse homeodomain interacting protein kinase 2 (Hipk2), with C-terminal MYC/DDK tag, expressed in HEK293T cells, 20ug
Species:	Mouse
Expression Host:	HEK293T
Expression cDNA Clone or AA Sequence:	>MR227349 representing NM_001136065 <mark>Red</mark> =Cloning site Green=Tags(s)
	MASHVQVFSPHTLQSSAFCSVKKLKVEPSSNWDMTGYGSHSKVYSQSKNIPPSQPASTTVSTSLPIPNPS LPYEQTIIFPGSTGHIVVTSASSTSVTGQVLGGPHNLMRRSTVSLLDTYQKCGLKRKSEEIENTSSVQII EEHPPMIQNNASGATVATATTSTATSKNSGSNSEGDYQLVQHEVLCSMTNTYEVLEFLGRGTFGQVVKCW KRGTNEIVAIKILKNHPSYARQGQIEVSILARLSTESADDYNFVRAYECFQHKNHTCLVFEMLEQNLYDF LKQNKFSPLPLKYIRPVLQQVATALMKLKSLGLIHADLKPENIMLVDPSRQPYRVKVIDFGSASHVSKAV CSTYLQSRYYRAPEIILGLPFCEAIDMWSLGCVIAELFLGWPLYPGASEYDQIRYISQTQGLPAEYLLSA GTKTTRFFNRDTDSPYPLWRLKTPDDHEAETGIKSKEARKYIFNCLDDMAQVNMTTDLEGSDMLVEKADR REFIDLLKKMLTIDADKRVTPIETLNHPFVTMTHLLDFPHSAHVKSCFQNMEICKRRVNMYDTVNQSKTP FITHVAPSTSTNLTMTFNNQLTTVHNQPSAASMAAVAPRSMPLQTGTAQICARPDPFQQALIVCPPGFQG LQASPSKHAGYSVRMENAVPIVTQAPGAQPLQIQPGLLAQAWPGGAQQILLPPAWQQLTGVATHTSVQHA AVIPETMAGTQQLADWRNTHAHGSHYNPIMQQPALLTGHVTLPAAQPLNVGVAHVMRQQPTSTTSSRKSK QHQSSVRNVSTCEVTSSQAISSPQRSKRVKENTPPRCAMVHSSPACSTSVTCGWGDVASSTTRERQRQTI VIPDTPSPTVSVITISSDTDEEEEQKHAPTSTVSKQRKNVISCVTVHDSPYSDSSSNTSPYSVQQRTGHN GTNTLDTKGGLENHCTGNPRTIIVPPLKTQASEVLVECDSLGPAISASHHSSSFKSKSSSTVTSTSGHSS GSSSGAIAYRQQRPGPHFQQQQPLNLSQAQQHMAADRTGSHRRQQAYITPTMAQAPYTFPHNSPSHGTVH PHLAAAAHLPTQPHLYTYAPTALGSTGTVAHLVASQGSARHTVQHTAYPASIVHQVPVSMGPRVLPSPT IHPSQYPAQFAHQTYISASPASTVYTGYPLSPAKVNQYPYI
Tag:	C-MYC/DDK
Predicted MW:	127.3 kDa
Concentration:	>0.05 ug/ul as determined by microplate BCA method
Durity:	> 80% as determined by SDS PAGE and Coomassia blue staining
runty.	- 50% as determined by 5D3-FAGE and Coomassie blue staining



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	Hipk2 (NM_001136065) Mouse Recombinant Protein – TP527349
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol
Note:	For testing in cell culture applications, please filter before use. Note that you may experience some loss of protein during the filtration process.
Storage:	Store at -80°C after receiving vials.
Stability:	Stable for 12 months from the date of receipt of the product under proper storage and handling conditions. Avoid repeated freeze-thaw cycles.
RefSeq:	<u>NP 001129537</u>
Locus ID:	15258
UniProt ID:	<u>Q9QZR5</u> , <u>A0A0R4J204</u>
RefSeq Size:	3972
Cytogenetics:	6 B1
RefSeq ORF:	3486
Synonyms:	1110014O20Rik; B230339E18Rik; Stank
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 TGFB, 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]

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