

# **Product datasheet for MC210165**

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## Sh3kbp1 (NM\_001135728) Mouse Untagged Clone

### **Product data:**

**Product Type:** Expression Plasmids

Product Name: Sh3kbp1 (NM\_001135728) Mouse Untagged Clone

Tag: Tag Free Symbol: Sh3kbp1

Synonyms: 1200007H22Rik; 1700125L08Rik; 5830464D22Rik; Al447724; Cin85; IN85; Ruk; Seta

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

Cell Selection: Neomycin

Fully Sequenced ORF: >MC210165 representing NM\_001135728

Red=Cloning site Blue=ORF Orange=Stop codon

ATGGGAGAGAGACAATTGGGAAGAAGTTACCTCCAGCTACATCAACTCCAGACCCATCGAAGACGGAGA TGGACAGCAGGACAAAGACCAAGGATTACTGCAAAGTAATATTTCCATATGAGGCACAGAATGATGATGA ATTGACAATCAAGGAAGGAGATATAGTGACTCTCATCAATAAGGACTGCATCGACGTAGGCTGGTGGGAA GGAGAACTCAATGGCAGACGAGGCGTGTTTCCTGACAACTTCGTGAAGTTACTTCCATCGGACTTTGACA AGGAAGGGAATAGACCCAAGAAACCGCCACCTCCATCAGCTCCTGTCGTCAAACAAGGGGCAGGTACCAC AGAGAAAACATGAAATTAAAAAAATACCTCCTGAAAGACCAGAAACCCTTCCAAACAGAACAGAAGAG AAAGAAAGACCAGAGAGAGAGCCAAAACTGGATTTACAGAAGCCTTCAGTTCCTGCCATCCCACCAAAAA AGCCTCGGCCACCTAAGACCAATTCTCTGAACAGACCTGGTGCATTGCCCCCAAGAAGACCTGAGCGACC ATCCTGGATAAGGACCTCTCAGACCGCAGCAATGATATTGACTTAGAAGGTTTTGACTCTGTGATATCAT CTACTGAGAAACTCAGCCACCCAACCACAAGCAGACCAAAAGCTACAGGAAGACGGCCTCCATCCCAGTC ACTCACATCTTCTTCCCTTTCAAGCCCCGATATCTTTGACTCCCCAAGTCCCGAAGAAGATAAAGAGGAA CACATTTCGCTTGCGCACAGAGGAATAGACGTGTCAAAGAAGACTTCCAAGACTGTTACCATATCCCAAG TGTCTGATAACAAGACATCCCTGCCACCCAAGCCAGGGACCATGGCAGCTGCCAGCAGTGGGCCAGCTTC TCTCTCTCAGTGGCATCCTCACCCATGTCATCCTCTTTGGGAACAGCTGGACAGAGAGCCAGTTCTCCA TCTCTGTTCAGCACAGAAGGAAAGCCAAAGATGGAGCCAGCAGTGAGCAGCCAGGCTGCTATCGAGGAGC TTAAGATGCAAGTCCGTGAGCTGAGGACCATCATTGAGACCATGAAGGACCAGCAGAAACGTGAGATTAA GCAGTTACTGTCAGAATTGGATGAAGAGAAAAAGATCCGGCTCCGGTTGCAGATGGAAGTGAACGACATA AAGAAAGCTCTTCAATCAAAGTGA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATTACAAGGATGACGACGATAAGGTTTAA



#### Sh3kbp1 (NM\_001135728) Mouse Untagged Clone - MC210165

**Restriction Sites:** Sgfl-Mlul

**ACCN:** NM 001135728

**Insert Size:** 1284 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 001135728.2, NP 001129200.1

 RefSeq Size:
 4170 bp

 RefSeq ORF:
 1284 bp

 Locus ID:
 58194

 UniProt ID:
 Q8R550

 Cytogenetics:
 X F4

**Gene Summary:** Adapter protein involved in regulating diverse signal transduction pathways. Involved in the

regulation of endocytosis and lysosomal degradation of ligand-induced receptor tyrosine kinases, including EGFR and MET/hepatocyte growth factor receptor, through an association with CBL and endophilins. The association with CBL, and thus the receptor internalization, may be inhibited by an interaction with PDCD6IP and/or SPRY2. Involved in regulation of ligand-dependent endocytosis of the IgE receptor. Attenuates phosphatidylinositol 3-kinase activity by interaction with its regulatory subunit. May be involved in regulation of cell adhesion; promotes the interaction between TTK2B and PDCD6IP. May be involved in the regulation of cellular stress response via the MAPK pathways through its interaction with MAP3K4. Is involved in modulation of tumor necrosis factor mediated apoptosis. Plays a role in the regulation of cell morphology and cytoskeletal organization. Required in the control of cell shape and migration (By similarity).[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (3, also known as Rukm3) differs in the 5' UTR, lacks a portion of the 5' coding region, and initiates translation at an alternate start codon, compared to variant 1. The encoded isoform (3) has a distinct N-terminus and is shorter than isoform 1.