

Product datasheet for MC213186

Tsku (NM_001168541) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Tsku (NM_001168541) Mouse Untagged Clone

Tag: Tag Free
Symbol: Tsku

Synonyms: 9530051K01Rik; E2ig4; Lrrc54; Tsk

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

Cell Selection: Neomycin

Fully Sequenced ORF: >MC213186 representing NM_001168541

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ATGCTGTGCTCTCTGTTCCTGCTGCTGCCGTGGCCAGACTGCAGACGACTCGGCCGTGTTTCCCTG GCTGCCAGTGTGAAGAAGAGACATTCGGCCTCTTTGACAGTTTCAGCCTGATCCGTGTGGACTGCAGCAG CCTGGGCCCCACATTGTGCCTGTGCCCATCCCTTTGGATACAGCCCACCTGGACCTGTCTTCCAACCGG ACAACCTGCTCACCAGCATCATGCCCTCTGCCTTCTCCCGACTGCGCTACCTGGAGTCACTTGACCTCAG CCACAATGGCCTGGCAGCCCTGCCGGCAGAGATTTTCACCAGCTCCCCCTTGAGTGACATCAATCTGAGC TCAGAGCCTGAACCTGTCCTGGAACCGATTCCGAGCTGTGCCCGATCTCCGAGACCTACCCCTGCGTTAC CTGAGCCTGGATGGGAACCCTCTGGCTACCATCAACCCAGATGCCTTCATGGGGCTGGCAGGCCTCACCC ACCTGTCACTGGCCAGCCTGCAGGGCATCCTCCATCTACCACCCCACGGCTTCCGGGAGCTCCCAGGCCT TCAGGTCCTGGACTTGTCAGGCAACCCCAAGCTCAAGTGGGCAGGAGCTGAGGTGTTTTCAGGCCTGGGT CTGCTTTACAAAGTGTCAGCGTAGGCCAGGATGTGCAGTGCCGGCGCCTGGTGCGGGAGGGCGCCTACCA CCGGCAGCCTGGCTCCAGCCCTAAGGTAGTCCTACACTGTGGAGACACCCAGGAATCTGCTGCCAGGGGC **CCAGACATTTTGTGA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul

ACCN: NM_001168541



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Insert Size: 1065 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 001168541.1, NP 001162013.1

RefSeq Size: 2760 bp RefSeq ORF: 1065 bp Locus ID: 244152 UniProt ID: Q8CBR6

Cytogenetics: 7 E1



Gene Summary:

Contributes to various developmental events and other processes such as wound healing and cholesterol homeostasis through its interactions with multiple signaling pathways (PubMed:21856951, PubMed:22995554, PubMed:25159578, PubMed:31391339). Wnt signaling inhibitor which competes with WNT2B for binding to Wnt receptor FZD4 and represses WNT2B-dependent development of the peripheral eye (PubMed:21856951). Plays a role in regulating the hair cycle by controlling TGFB1 signaling (PubMed:22995554). Required for the development of the anterior commissure in the brain by inhibiting neurite outgrowth (PubMed:21055390, PubMed:23206892). Essential for terminal differentiation of hippocampal neural stem cells (PubMed:31983064). Plays a role in regulating bone elongation and bone mass by modulating growth plate chondrocyte function and overall body size (PubMed:30271858). Required for development of the inner ear through its involvement in stereocilia formation in inner hair cells (PubMed:32127020). Facilitates wound healing by inhibiting secretion of TGFB1 from macrophages which prevents myofibroblast differentiation, maintaining inflammatory cell quiescence (PubMed:25159578). Plays a role in cholesterol homeostasis by reducing circulating high-density lipoprotein cholesterol, lowering cholesterol efflux capacity and decreasing cholesterol-to-bile acid conversion in the liver (PubMed:31391339). In one study, shown to negatively regulate sympathetic innervation in brown fat, leading to reduced energy expenditure (PubMed:31535079). In another study, shown not to affect brown fat thermogenic capacity, body weight gain or glucose homeostasis (PubMed:31767170).[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (1) represents the longest transcript. All four variants encode the same protein. 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.