

Product datasheet for MC207553

Zfand6 (NM_022985) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Zfand6 (NM_022985) Mouse Untagged Clone

Tag: Tag Free Symbol: Zfand6

Synonyms: 3110005P07Rik; Awp1; Za20d3

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

Cell Selection: Neomycin

Fully Sequenced ORF: >MC207553 representing NM_022985

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGAGATCTGCC

GCCGCGATCGCC

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul ACCN: NM_022985

Insert Size: 672 bp



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Zfand6 (NM_022985) Mouse Untagged Clone - MC207553

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: <u>NM 022985.6</u>, <u>NP 075361.2</u>

RefSeq Size: 1658 bp
RefSeq ORF: 672 bp
Locus ID: 65098
UniProt ID: Q9DCH6

Cytogenetics: 7 D3

Gene Summary: Involved in regulation of TNF-alpha induced NF-kappa-B activation and apoptosis. Involved in

modulation of 'Lys-48'-linked polyubiquitination status of TRAF2 and decreases association of TRAF2 with RIPK1 (By similarity). Required for PTS1 target sequence-dependent protein

import into peroxisomes and PEX5 stability; may cooperate with PEX6. In vitro involved in

PEX5 export from the cytosol to peroxisomes.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (6) differs in the 5' UTR compared to variant 1. Variants 1-6 all encode the same isoform (a). 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.