

Product datasheet for **MC223591**

Otud4 (NM_001081164) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids
Product Name: Otud4 (NM_001081164) Mouse Untagged Clone
Tag: Tag Free
Symbol: Otud4
Synonyms: 4930431L18Rik; AI449692; D8Ert69e; mKIAA1046
Vector: pCMV6-Entry (PS100001)
E. coli Selection: Kanamycin (25 ug/mL)
Cell Selection: Neomycin
Fully Sequenced ORF: >MC223591 representing NM_001081164
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGGATCGC**C

ATGGAGGCAGCCGTCGGCGGCCGACGGCGTGGACCAGGGCGGCGTGGGGCCGCTGGAGGATGAGACGC
 CCATGGACGCCTATCTGCGCAAACCTGGGCTTGTATCGGAAATGGTCGCCAAGGACGGGTCGTGCTTGT
 CCGGGCTGTGGCCGAGCAGGTGTGCACCTCAGTCTCGGCATGTGGAGTTAGGATGGCCTGTATCCGC
 TACCTTCGAGAGAACAGAGAGAAATTTGAAGCGTTTATAGAAGGGTCATTTGAAGAATATTTAAACGTT
 TGGAAAATCCACAGGAATGGGTAGGACAAGTGGAAATAAGTGCCCTTTCACCTATGTACAGGAAAGATT
 TGTAATATATCAGGAGCCAAATGTTTCTCCTTCACATGTAACGAAAATAATTTCTGAGAAGGTGTTA
 CTGTGTTTTCAAATGGAAATCATTATGACATTGTCTATCCCATAAACATATAAGATAGTTCTGCTATGT
 GTCAGTCTCTCCTTATGAGTTGCTGTATGAGAAGGATTCAAAACCTGATGTTAGTAAATCATGATGGG
 ACTAGAAGCCTCTGAGGTGGCTGAGGAGAGCAACAGTGAGATATCAGACTCTGAGGACGACAGCTGCAAG
 AGTAAAAGTACTGCTGCTACTGATGTGAATGGATTTAAACCCTCAGGCAGTGAGAACCCTAAGAACAATG
 GAACTCAGTGCCTTCTTTGTCCAGAAAGGTTCTTAAGTCACTCAACCCAGCAGCTATAGAAATGT
 GGAGTATGAAATTTGGTTGAAGTCTAAACAAGCTCAACAAAAACGTGATTATCCATTGCTGCTGGCTTA
 CAGTATGAAGTTGGAGATAAATGCCACGTTAGATTGGATCATAATGGGAAATTATCTAATGCAGACATTC
 ATGGGGTTCACCTGAGAAATGGACTGGTTTTGTCTGAAGAAGTGGGAAAAAACATACACCGAAGAACCT
 CAAGCCACCTCCCCAGAAAGCTGGAACACGGTGTGAGGAAAGAAGATGAAAAACCTAATTCTGGGCAA
 AATTTCCATTAGATACAGATTACAGAGGGCCAAAGAATCTAAACAAGCCAAATCAAGCCCCATCTGCAC
 TACCTCCTCGACTCCAGCATCCTCATCGGGTGAAGACAGCATGCATTCTCCAGTCATTCTACAGGGTC
 CCAGTCTCAGAAATCCTCAGTGAGCATAAGAATCTAAGTAGGATGCCCTCACAGATCACAAGAAAACCT
 GACCGTGAAGGGCTGAGGACTTTGATCAGTGAGTCGTGAATCTTACTATTTTGGCCTCTCCCCAGAAG
 AACGCAGAGAGAAGCAAGCTATTGAAGAGTCTCGTTTACTGTATGAGATTCAGAACCAGGATGAACAGGC
 TTTCCCTGCCCTTTCTAGTTCATCAGTCAGTCAGTCACCTTCTCAGAATAGCAATGCGTGTGTCCCAAGG
 AAGTCTTCACATGCAAGGGACAGGAAAGGAAGCATGCGGAGAGCAGACGCAGAGGAACGAAAGGACAAAG



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ACTCTCTACGTGGCCATACTCATGTGGATAAAAAACCGGAGCCAAGCACACTGGAGATCAGCGATGATAA
 ATGTACAAGAGTTTCATCACCATCTAAGTCAAAGAAAGAGTGCCCATCTCCTGTAGAACAAAAGCCAGCA
 GAACATATACCTTTGTCAAATCCAGCTCCCCTTCTAGTTTCTCCAGAAGTACATCTACTCCTGCGGTGC
 CTTCTTTACCAGCCACTGTGCCAGCCTGGCCAAGTGAACCTACAACCTTCGGACCAACAGGTGTCCCTGC
 TCAGATTCCCATTTTGTCAAGTACACAGACCCTGGACCTGATGCTGCCGTGTACAAGCGCATTTAACA
 CCTTCTCCGGTTCCTGTGTCAATTCAGGCAGTTAACAGCCCTTGATGCCTTTCCTCAGACAATGAGCC
 TCTATCAAGACCCCTCTATCCTGGGTTTCCTTGTAGTGAAGGGAGATCGAGCCATTGCACCACCTTA
 TTCACTGTGTGACACCGGGGAGGACCTGCCTAAAGATAAGAATATTCTTCGATTCTTCTCAATCTCGGT
 GTAAAGGCATATAGTTGTCTATGTGGGCCACATTCTTACCTATATCCTCTGCACCAGGCCTATATGG
 CAGCCTGCAGGATGTACCCAAAGGTCCCTGTTCCCGTGTATCCTCAGAATACTTGGTTCCAAGAAGCCCC
 TCCTGCTCAGAGTGAAAGTGACTGTCTTGACCCGATGCCACTACTCTCTGCACCCCGAGGCCAGTGT
 AATGGTCAGATGCCACAGGCAGAGATGGGACCGCCTGCATTTGCATCACCTCTGGTTATCCCTCCATCTC
 AGGTGTCTGAAGTGCATGGACAATTGTCTTACCAACCTGAACTGGAGTCTGAGAACCAGGGCAGCTTCT
 TCATGCTGAATATGAAGAGTCACCTAGTGGCAAGAACATGTACCCACAACAGTCTTTTGGCCTAACCCA
 TTTTATAGTCTGTTCCATTGCACCTCCTTTCTCCCTCATGTTTGGTATGGGTATCCTTTTCAGGGAT
 TCGTAGAAAAATCCTGTAATGAGGCAAAATATTGCTCCTGCCCTGATGATAAAGGAGAATTGGATTGGCC
 TTTGGAGAATCTAGATCTGTCTAAAGAATGCGATTCTGTCTCAGCAGTAGATGAGTTCCAGACGCCAGA
 GTTGAAGGTGCACATTCTCTGTCTGCAGCGAGTGTGAGCAGCAAGCACGAAGGCCGAGTGGAGCAGTCGT
 CCCAGACCCGGAAGGCAGACATAGACTTGGCTTCAAGTCTTCTGCAGTGAAGGAAAGGGTCATCCTCC
 CACTCAGATTCTAAACAGAGAAAAGAGAACCTGGGTCTGCTGAACCTGAGCCTAAGAGGACCATTCAAAGT
 CTGAAAGAAAACCAGAGAAAAGTAAAAGATCCCAAGACTGCTGCTGATGTGGTCAGCCCTGGGGCAATT
 CTGTGGATAGATTGCAAAGACCAAAAAGAAGAGAGTTTCAAGAATGAGAATGAAGTATCTAATATTTTGG
 AAGTGGCAGATCCAAGCAGTTTTATAATCAAACCTTACGGAAGCAGGAAGTACAAAAGTATTGGGGCTCT
 TCTGGTCGAGGTGGCTATCAACACGTGAGAGCGAGGATCCTGGAAAGGGCAGCCAAATCGAAGCCGGG
 ATGAAGGTTATCAGTACCATCGACATGTTAGAGGACGCCCATACAGGGGAGATAGGAGGAGATCAGGGAT
 GGGAGATGGCCACAGGGGACAACACTTAA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-MluI
- ACCN:** NM_001081164
- Insert Size:** 3321 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_001081164.1](#), [NP_001074633.1](#)

RefSeq Size: 7324 bp

RefSeq ORF: 3321 bp

Locus ID: 73945

UniProt ID: [B2RRE7](#)

Cytogenetics: 8 37.74 cM

Gene Summary: Deubiquitinase which hydrolyzes the isopeptide bond between the ubiquitin C-terminus and the lysine epsilon-amino group of the target protein. May negatively regulate inflammatory and pathogen recognition signaling in innate immune response. Upon phosphorylation at Ser-202 and Ser-204 residues, via IL-1 receptor and Toll-like receptor signaling pathway, specifically deubiquitinates 'Lys-63'-polyubiquitinated MYD88 adapter protein triggering down-regulation of NF-kappa-B-dependent transcription of inflammatory mediators (PubMed:29395066). Independently of the catalytic activity, acts as a scaffold for alternative deubiquitinases to assemble specific deubiquitinase-substrate complexes. Associates with USP7 and USP9X deubiquitinases to stabilize alkylation repair enzyme ALKBH3, thereby promoting the repair of alkylated DNA lesions (By similarity).[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (1) represents the longer transcript and encodes the longer isoform (1). 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.