

Product datasheet for SC304609

YOD1 (NM 018566) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: YOD1 (NM_018566) Human Untagged Clone

Tag: Tag Free Symbol: YOD1

Synonyms: DUBA8; OTUD2; PRO0907

Mammalian Cell N

Selection:

Neomycin

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

Fully Sequenced ORF: >SC304609 representing NM_018566.

Blue=Insert sequence Red=Cloning site Green=Tag(s)

GATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGTTTGGCCCCGCTAAAGGTCGCCATTTTGGAGTCCACCCGGCGCCCTGGTTTCCCCGGCGGCGTCTCC CAACAGGCTGCCGGGACCAAAGCTGGCCCGCGGGTGCCTGGCCTGTGGGCAGCCGGACCGACACGATG TGGCGGCTCCGCTGCAAGGCCAAGGACGGCACCCATGTTTTGCAGGGGCTGTCCAGCCGGACCCGGGTG CGGGAACTCCAGGGCCAAATTGCCGCCATCACCGGGATCGCCCCCGGCGGTCAGCGAATCCTCGTCGGA TACCCTCCGAGTGCCTGGATCTCAGCAATGGGGATACCATTCTGGAAGACTTGCCCATCCAATCTGGT GACATGCTGATCATTGAAGAAGACCAAACCAGGCCCAGAAGTTCACCTGCATTTACTAAACGTGGTGCT TCTAGTTACGTCAGGGAAACTTTGCCTGTGCTTACCAGAACCGTGGTCCCAGCAGACAACTCTTGCCTC TTTACTAGTGTGTACTATGTCGTCGAAGGAGGAGTCTTGAATCCAGCTTGTGCCCCTGAGATGAGACGC CTCATAGCACAATTGTAGCAAGCGATCCAGACTTCTATAGTGAGGCAATACTGGGAAAAACAAATCAA GAGTACTGTGACTGGATCAAAAGGGATGACACTTGGGGAGGAGCAATAGAGATATCGATTTTGTCCAAG GGATATACCAAAAGGGTTCTGCTTATTTATGATGGCATCCACTATGATCCACTTCAGCGTAACTTCCCT GATCCAGATACACCTCCTCTGACCATTTTCTCCTCTAATGATGATATTGTTCTTGTACAAGCACTGGAA TTAGCAGATGAAGCTAGAAGAAGGAGACAGTTTACTGATGTCAACCGCTTCACCCTGAGATGCATGGTA TGTCAGAAAGGATTAACTGGACAAGCAGAAGCAAGGGAACATGCCAAGGAGACAGGCCATACCAACTTT **GGAGAAGTGTGA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGAT

TACAAGGATGACGACGATAAGGTTTAAACGGCCGGC

Restriction Sites: Sgfl-Mlul



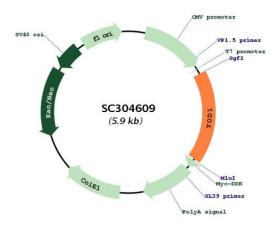
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Plasmid Map:



ACCN: NM_018566 **Insert Size:** 1047 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).

OTI Annotation: This TrueClone is provided through our Custom Cloning Process that includes sub-cloning

into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.

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 018566.3</u>

 RefSeq Size:
 6265 bp

 RefSeq ORF:
 1047 bp

 Locus ID:
 55432



YOD1 (NM_018566) Human Untagged Clone - SC304609

UniProt ID: Q5VVQ6

Cytogenetics: 1q32.1

Protein Pathways: Biosynthesis of unsaturated fatty acids, Limonene and pinene degradation

MW: 38.3 kDa

Gene Summary: Protein ubiquitination controls many intracellular processes, including cell cycle progression,

transcriptional activation, and signal transduction. This dynamic process, involving ubiquitin

conjugating enzymes and deubiquitinating enzymes, adds and removes ubiquitin. Deubiquitinating enzymes are cysteine proteases that specifically cleave ubiquitin from ubiquitin-conjugated protein substrates. The protein encoded by this gene belongs to a DUB subfamily characterized by an ovarian tumor (OTU) domain. Alternative splicing results in

multiple transcript variants. [provided by RefSeq, Jan 2013]

Transcript Variant: This variant (1) 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.