

Product datasheet for **MC221199**

Ythdc1 (NM_177680) Mouse Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	Ythdc1 (NM_177680) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Ythdc1
Synonyms:	A730098D12Rik; C80342; mKIAA1966
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)



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Fully Sequenced ORF: >MC221199 representing NM_177680
 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC
 GCC**CGCATCGCC**

ATGGCGGCCGACAGCCGGGAGGAGAAAGATGGGGAACCTAATGTTTTGGATGATATTTTGACTGAAGTAC
 CAGAACAGGATGATGAATTGTATAATCCAGAGAGTGAACAAGATAAAAAATGAGAAAAAGGATCAAAAAG
 AAAAAAGTGAAGAAATGGAGTCTACTGACACCAAGCGACAAAAACCTTCCATCCATTCAAGACAATTGATT
 TCTAAGCCACTAAGCTCATCTGTAAGCAATAATAAAGAATAGTTAGTACAAAAGGAAAGTCGGTTACAG
 AATATAAAAAATGAGGAATATCAAAGATCTGAAAGGAACAAGCGTCTAGATGCTGATCGAAAAATTCGCT
 GTCAAGCAGTTCCCTAGAGAACCTTACAAGAGTCAACCAGAAAAAACTGTCTACGGAAAAAGGATTCT
 GAAAGAAGGGCCAAGTCTCTACACCAGATGGTCTGAGAGAATTGGGCTGAAGTTGATAGACGTGCAA
 GCAGATCCAGCCAGTCTCAAAGGAAGAGGTGAACTCTGAAGACTATGGCTCTGACCACGAGACAGGAAG
 CAGTGGTCTTCTGAACAGGGCAACAACACTGAGAATGAGGAGGAAGGAGGGGAGGAAGATGTAGAGGAA
 GATGAAGAGGTAGATGAAGATGCAGAAGATGATGAGGAAGTGGATGAAGATGCAGAGGAGGAGGAGGAAG
 AGGATGAAGAGGAAGAAGAGGATGAGGACGAGGATGAAGAGGAAGAAGAATATGAACAGGATGAGAGAGA
 TCAGAAGGAAGAAGGAAATGATTATGACACCCGAAGTGAAGGCGAGTATTCTGGTCCGAGTCTGTTTCC
 TTCACAGATGGATCTGTGATCTGGGTGAGGAACAGATGGATCAGATGAGAAAAAGAAGGAAAGGAAAGA
 GAGCTCGAGGCATATCACCCATTGTCTTTGATAGAAGTGGCAGTTCTGCATCAGAGTCATATGCAGGTTC
 AGAAAAAGAAGCATGAGAAATTATCATCTCCGTTCTGCTGTCCGAAAAGATCAAACAGTAACTCAA
 TATGTCCTTCAGGATGCAAGATTTTCTCATAAAGAGTAATAACCATGAGAATGTGTCTCTTGCCAAAG
 CAAAGGGTGTATGGTCCACATTGCCTGTAATGAGAAGAAATTAATCTTGCAATTTAGATCTGCAAGGAG
 TGTTATATTAATATTTTCTGTGACAGAAAAGTGGAAAAATTTCAAGGTTTTGCCAGATTGTCTTCAGAATCA
 CACCATGGCGGATCTCCTATACACTGGGTGCTTCCAGCAGGAATGAGTGCTAAAAATGCTTGAGGTGTTT
 TAAAAATAGACTGGATTTGCAGGCGTGAATTACCTTTACTAAATCAGCTCATCTACCAATCCCTGGAA
 TGAACATAAGCCAGTAAAGATTGGACGTGATGGACAGGAAATGAACTTGAATGTGGAACCTCAGCTTTGT
 CTTCTGTTTCCCCTGATGAAAGTATTGACTTGTATCAGCTCATTCAAAAATGCGTCACAAGAGAAGAA
 TGCATTCTCAGCCTCGATCAAGAGGACGTCATCCCGTCGAGAACCAGTCCGGGATGTGGGAAGGCGTCCG
 ACCAGAAGATTATGATATTCATAACAGCAGAAAAGAAACCAAGGATTGACTATCCCCCTGAGTTTCACCAG
 AGACCAGGATGTAAGGATCCCCGATACCAGGAAGTGGACAGTTTTACAAATCTTATCCCAACAGAC
 GATTTTCAGGAGTTCGCCGAGATGTGTTTTAAATGGGTCTACAATGATTATGTGAGGGAAATTCATAA
 CATGGGACCACCGCTCCTTGCAAGGAATGCCTCCTTACCCGGGAATAGAACAACCTCCACACCATCCT
 TACTACCAGCACCATGCCCCACCTCCTCAAGCCCACCCCTTACTCAGGACACCATCCGGTACCACATG
 AAGCAAGATACAGAGATAAACGAGTACATGACTATGATATGAGGGTCTGATGATTTCTTCGCCGACGCA
 AGCTGTGGTCAGTGGTCCGAGAAGTAGACCTCGAGAAAGAGATCGGGAGCGAGAGCGAGACCCGCCAGAG
 GATAACAGAAGAGATAGAGAGCGAGACAGAGGTCTGATCGAGAAAAGAGAGAGAGAAAGAATATGTGATC
 GGGACAGAGACCGAGGGGAGAGAGGTCTTATCGAAGAT**TAA**

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT
 ACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: SgfI-MluI

ACCN: NM_177680

Insert Size: 2211 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:	<ol style="list-style-type: none">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_177680.3 , NP_808348.2
RefSeq Size:	3034 bp
RefSeq ORF:	2211 bp
Locus ID:	231386
UniProt ID:	E9Q5K9
Cytogenetics:	5 E1
Gene Summary:	<p>Regulator of alternative splicing that specifically recognizes and binds N6-methyladenosine (m6A)-containing RNAs (By similarity). M6A is a modification present at internal sites of mRNAs and some non-coding RNAs and plays a role in the efficiency of mRNA splicing, processing and stability (By similarity). Acts as a key regulator of exon-inclusion or exon-skipping during alternative splicing via interaction with mRNA splicing factors SRSF3 and SRSF10 (By similarity). Specifically binds m6A-containing mRNAs and promotes recruitment of SRSF3 to its mRNA-binding elements adjacent to m6A sites, leading to exon-inclusion during alternative splicing (By similarity). In contrast, interaction with SRSF3 prevents interaction with SRSF10, a splicing factor that promotes exon skipping: this prevents SRSF10 from binding to its mRNA-binding sites close to m6A-containing regions, leading to inhibit exon skipping during alternative splicing (By similarity). May also regulate alternative splice site selection (By similarity). Also involved in nuclear export of m6A-containing mRNAs via interaction with SRSF3: interaction with SRSF3 facilitates m6A-containing mRNA-binding to both SRSF3 and NXF1, promoting mRNA nuclear export (By similarity). Also recognizes and binds m6A on other RNA molecules (By similarity). Involved in random X inactivation mediated by Xist RNA: recognizes and binds m6A-containing Xist and promotes transcription repression activity of Xist (By similarity). Involved in S-adenosyl-L-methionine homeostasis by regulating expression of MAT2A transcripts, probably by binding m6A-containing MAT2A mRNAs (PubMed:29262316).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (1) lacks an alternate in-frame exon and uses an alternate in-frame splice junction compared to variant 4. The resulting isoform (1) has the same N- and C-termini but is shorter compared to isoform 4. 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.</p>