

Product datasheet for **SC307751**

DNAH12 (NM_198564) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	DNAH12 (NM_198564) Human Untagged Clone
Tag:	Tag Free
Symbol:	DNAH12
Synonyms:	DHC3; DLP3; DLP12; DNAH7L; DNAH12L; DNAHC3; DNAHC12; DNHD2; HDHC3; HL-19; HL19
Vector:	<u>pCMV6 series</u>
Fully Sequenced ORF:	>NCBI ORF sequence for NM_198564, the custom clone sequence may differ by one or more nucleotides ATGTCAGATGCAAACAAAGCTGCCATTGCAGCAGAAAAGGAAGCTCTGAAGTTA CCCCCATTGTCCATCTCCAGAAAACATAGGCGTTGATACACCAACACAAAGTAAGCTG CTAAAATACAGAAGATCCAAGGAGCAGCAGCAGAAAATTAATCAGTTAGTAATTGATGGA GCCAAAAGAAATTTAGACAGAACACTGGGTAAAAGAACACCTCTATTACCACCACCTGAT TATCCTCAAACATGACCAGTGAATGAAAAAAAAAGGATTCAACTATATTTATATGAAG CAATGTGTAGAAAGTAGTCCTTTAGTACCTATTCAGCAGGAATGGCTGGATCACATGTTA AGGCTGATACCTGAGTCTTTAAAGGAAGGAAAGAAAGAGAAGAACTTCTTGAAAGTCTC ATAAATGAGGTGTCAAGTGACTTTGAAAACAGCATGAAGAGATATTTGGTGCAGAGCGTT CTTGTAACACCAGTTAAATCGCTTGAAGATGAAGGAGGTCCTTTACCTGAATCTCCT GTAGGCCTAGATTATTCTAATCCTTGGCATTCTAGCTATGTGCAGGCAAGAAATCAAATA TTCTCTAATTTGCACATTATTCATCCAACATGAAAATGTTACTGGACCTTGGTTATACA ACATTTGCTGATACAGTTTTGTTGGACTTCACAGGAATTAGAGCTAAAGGTCCAATTGAC TGTGAATCACTGAAAACCTGATCTATCAATACAACTAGAAACGCAGAAGAGAAGATAATG AATACATGGTATCCAAAGTTATAAATCTCTTTACCAAGAAGGAGGCACTAGAAGGTGTT AAACCTGAAAAATTGGATGCATTTTATAGCTGTGTTCCACACTTATGTCAAATCAGCTA AAGGATCTATTAAGGAGAACTGTAGAAGATTTGTAATACTTTGACCCAAAAGATCAA CAAAGGCTGCCAATATTTAAGATAGAATTGACATTTGATGACGACAAAATGGAATTTTAT CCTACCTTTCAAGATTTGGAAGATAATGTCTTGAGTTTGGTGAACGAATAGCCGAAGCT CTGCAGAAATGTCCAAACAATCCCCTCTTGGCTATCAGGAACCTCAACACCAGTAAATCTT GACACAGAACTTCTGAACACGTGTTACTACTGGGCTGTTGATACACTGAAGGCAGCAGTA CATCGGAACCTAGAAGGTGCAAGAAAGCATTATGAGACATATGTTGAAAAATATAATTGG CTCCTTGATGGGACTGCAGTTGAGAATATAGAGACTTTTCAGACAGAAGATCATACTTTT GATGAATATACAGAGGAGCTGGATTGCTGGGTGGTATGGGAAGTGTATTTTTTAA
Restriction Sites:	Please inquire
ACCN:	NM_198564



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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:	<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:	<u>NM_198564.1</u> , <u>NP_940966.1</u>
RefSeq Size:	2014 bp
RefSeq ORF:	1374 bp
Locus ID:	201625
UniProt ID:	<u>Q6ZR08</u>
Cytogenetics:	3p14.3
Gene Summary:	<p>Force generating protein of respiratory cilia. Produces force towards the minus ends of microtubules. Dynein has ATPase activity; the force-producing power stroke is thought to occur on release of ADP. Involved in sperm motility; implicated in sperm flagellar assembly (By similarity).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (2) lacks multiple 3' exons but has an alternate 3' exon, as compared to variant 3. The resulting isoform (2) is much shorter and has a different C-terminus, as compared to isoform 3. 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.</p>