

Product datasheet for SC307298

OriGene Technologies, Inc.

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N acetyl transferase 5 (NAA20) (NM_181528) Human Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: N acetyl transferase 5 (NAA20) (NM_181528) Human Untagged Clone

Tag: Tag Free Symbol: NAA20

Synonyms: dJ1002M8.1; NAT3; NAT3P; NAT5P

Vector: <u>pCMV6 series</u>

Fully Sequenced ORF: >NCBI ORF sequence for NM_181528, the custom clone sequence may differ by one or more

nucleotides

GAAAGATATGAGGAAAGCACTTTCCAGGGATACTGA

Restriction Sites: Please inquire **ACCN:** NM 181528

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 181528.1</u>, <u>NP 852669.1</u>

RefSeq Size: 1214 bp
RefSeq ORF: 336 bp
Locus ID: 51126
UniProt ID: A6NHA3
Cytogenetics: 20p11.23

Protein Pathways: Glycerophospholipid metabolism, Limonene and pinene degradation, Phenylalanine

metabolism, Tyrosine metabolism

Gene Summary: NAT5 is a component of N-acetyltransferase complex B (NatB). Human NatB performs

cotranslational N(alpha)-terminal acetylation of methionine residues when they are followed by asparagine (Starheim et al., 2008 [PubMed 18570629]).[supplied by OMIM, Apr 2009] Transcript Variant: This variant (3) lacks an alternate exon compared to variant 1, which causes a frameshift. The exon combination for the transcript is supported by full-length ESTs (BG548527.1 and CD641402.1). The resulting isoform (c) is shorter and has a distinct C-

terminus compared to isoform a.