

Recombinant Full Length Human NAA25 Protein, GST-tagged

Cat. No. NAA25-2023HF **Lot. No.** (See product label)

SPECIFICATION

Product Overview	Human C12orf30 full-length ORF (NP_079229.2, 1 a.a. - 972 a.a.) recombinant protein with GST-tag at N-terminal.
Species	Human
Source	In Vitro Cell Free System
ProteinLength	972 amino acids
Description	MDM20 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]).
Form	50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.
Molecular Mass	138.7 kDa
AA Sequence	MATRGHVQDPNDRRLRPIYDYLDNGNKNMAIQQADKLLKKHKDLHCAKVLKAIGLQ RTGKQEEAFTLAQEVAALEPTDDNSLQALTILYREMRPELVTKLYEAAVKKVPNSE EYHSHLFMAYARVGEYKKMQQAGMALYKIVPKNPYYFWSVMSLIMQSISAQDENLS KTMFLPLAERMVEKMKVEDKIEAEAEVELYYMILERLGKYQEALDVIRGKLGEKLTSE IQSREKCMAMYKKLSRWPECNALSRRLLKNSDDWQFYLTIFYDSVFRLEEAWSP PAEGEHSLEGEVHYSAEKAVKFIEDRITEESKSSRHLRGPHLAKLELIRRLRSQGCN DEYKLGDPPEELMFQYFKKFGDKPCCFTDLKVFDLLPATQCTKFINQLLGVVPLSTP TEDKLALPADIRALQQHLCVVQLTRLLGLYHTMDKNQKLSVVRELMLRYQHGLEFGK

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TCLKTELQFSDYYCLLAVHALIDVWRETGDETTWQALTLEEGLTHSPSNAQFKLL
 LVRIYCMLGAFEPVVDLYSSSLDAKHQHDITIGYLLTRYAESLGQYAAASQSCNFAIRF
 FHSNQKDTSEYIIQAYKYGAFEKIPEFIAFRNRLNNSLHFAQVRTERMLLDLLLEANIS
 TSLAESIKSMNLRPEEDDIPWEDLRDNRDLNVFFSWDPKDRDVSEEHKKLSLEEETL
 WLRIRSLTLRLISGLPSLNHPVEPKNSEKTAENGVSRRIDILRLLLQQLEATLETGKRFI
 EKDIQYPFLGPVPTRMGGFFNSGCSQCQISSFYLVNDIYELDTSGLEDTMEIQUERIEN
 SFKSLLDQLKDVFSKCKGDLLEVKGDNLKTHTPTLLENLVFFVETISVILWVSSYCESV
 LRPYKLNQKKKKKKKETSIIIMPPVFTSFQDYVTGLQTLISNVVDHIKGLETHLIALKLE
 ELILEDTSLSPEERKFSKTVQGVQSSYLHSLLEMGELLKKRLETTKKLKI

Storage Store at -80 centigrade. Aliquot to avoid repeated freezing and thawing.

Storage Buffer 50 mM Tris-HCl, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.

GENE INFORMATION

Gene Name NAA25 N(alpha)-acetyltransferase 25, NatB auxiliary subunit [Homo sapiens]

Official Symbol NAA25

Synonyms NAA25; N(alpha)-acetyltransferase 25, NatB auxiliary subunit; C12orf30, chromosome 12 open reading frame 30; N-alpha-acetyltransferase 25, NatB auxiliary subunit; FLJ13089; natB complex subunit MDM20; mitochondrial distribution and morphology 20; N-terminal acetyltransferase B complex subunit NAA25; NAP1; MDM20; C12orf30; DKFZp667K2112

Gene ID 80018

mRNA Refseq NM_024953

Protein Refseq NP_079229

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MIM	612755
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UniProt ID	Q14CX7
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