

### Product datasheet for TP318271

#### OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

# NAGPA (NM\_016256) Human Recombinant Protein

**Product data:** 

**Product Type:** Recombinant Proteins

**Description:** Recombinant protein of human N-acetylglucosamine-1-phosphodiester alpha-N-

acetylglucosaminidase (NAGPA), 20 µg

Species: Human
Expression Host: HEK293T

Expression cDNA >RC218271 representing NM\_016256
Clone or AA Red=Cloning site Green=Tags(s)

Sequence:

MATSTGRWLLLRLALFGFLWEASGGLDSGASRDDDLLLPYPRARARLPRDCTRVRAGNREHESWPPPPAT PGAGGLAVRTFVSHFRDRAVAGHLTRAVEPLRTFSVLEPGGPGGCAARRRATVEETARAADCRVAQNGGF FRMNSGECLGNVVSDERRVSSSGGLQNAQFGIRRDGTLVTGYLSEEEVLDTENPFVQLLSGVVWLIRNGS IYINESQATECDETQETGSFSKFVNVISARTAIGHDRKGQLVLFHADGQTEQRGINLWEMAEFLLKQDVV NAINLDGGGSATFVLNGTLASYPSDHCQDNMWRCPRQVSTVVCVHEPRCQPPDCHGHGTCVDGYCQCTGH FWRGPGCDELDCGPSNCSQHGLCTETGCRCDAGWTGSNCSEECPLGWHGPGCQRPCKCEHHCPCDPKTGN

CSVSRVKQCLQPPEATLRAGELSFFTRTAWLALTLALAFLLLISIAANLSLLLSRAERNRRLHGDYAYHP

LQEMNGEPLAAEKEQPGGAHNPFKD

**TRTRPLEQKLISEEDLAANDILDYKDDDDKV** 

Tag: C-Myc/DDK

**Predicted MW:** 53.3 kDa

**Concentration:**  $>0.05 \mu g/\mu L$  as determined by microplate BCA method

**Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining

**Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3, 10% glycerol

**Preparation:** Recombinant protein was captured through anti-DDK affinity column followed by conventional

chromatography steps.

**Note:** For testing in cell culture applications, please filter before use. Note that you may experience some

loss of protein during the filtration process.

Storage: Store at -80°C.





#### NAGPA (NM\_016256) Human Recombinant Protein - TP318271

Stability: Stable for 12 months from the date of receipt of the product under proper storage and handling

conditions. Avoid repeated freeze-thaw cycles.

**RefSeq:** NP 057340

 Locus ID:
 51172

 UniProt ID:
 Q9UK23

 RefSeq Size:
 2219

 Cytogenetics:
 16p13.3

RefSeq ORF: 1545

Synonyms: APAA; UCE

**Summary:** Hydrolases are transported to lysosomes after binding to mannose 6-phosphate receptors in the

trans-Golgi network. This gene encodes the enzyme that catalyzes the second step in the formation of the mannose 6-phosphate recognition marker on lysosomal hydrolases. Commonly known as 'uncovering enzyme' or UCE, this enzyme removes N-acetyl-D-glucosamine (GlcNAc)

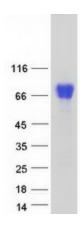
residues from GlcNAc-alpha-P-mannose moieties and thereby produces the recognition marker. The encoded preproprotein is proteolytically processed by furin to generate the mature enzyme, a homotetramer of two disulfide-linked homodimers. Mutations in this gene are associated with

developmental stuttering in human patients. [provided by RefSeq, Oct 2015]

**Protein Families:** Transmembrane

Protein Pathways: Lysosome

# **Product images:**



Coomassie blue staining of purified NAGPA protein (Cat# TP318271). The protein was produced from HEK293T cells transfected with NAGPA cDNA clone (Cat# [RC218271]) using MegaTran 2.0 (Cat# [TT210002]).