

Product datasheet for PH312434

OriGene Technologies, Inc.

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ASAH1 (NM_004315) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

Description: ASAH1 MS Standard C13 and N15-labeled recombinant protein (NP_004306)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC212434

or AA Sequence:

.

Predicted MW: 46.3 kDa

Protein Sequence: >RC212434 representing NM_004315

Red=Cloning site Green=Tags(s)

MNCCIGLGEKARGSHRASYPSLSALFTEASILGFGSFAVKAQWTEDCRKSTYPPSGPTYRGAVPWYTINL DLPPYKRWHELMLDKAPMLKVIVNSLKNMINTFVPSGKVMQVVDEKLPGLLGNFPGPFEEEMKGIAAVTD IPLGEIISFNIFYELFTICTSIVAEDKKGHLIHGRNMDFGVFLGWNINNDTWVITEQLKPLTVNLDFQRN NKTVFKASSFAGYVGMLTGFKPGLFSLTLNERFSINGGYLGILEWILGKKDAMWIGFLTRTVLENSTSYE EAKNLLTKTKILAPAYFILGGNQSGEGCVITRDRKESLDVYELDAKQGRWYVVQTNYDRWKHPFFLDDRR

TPAKMCLNRTSQENISFETMYDVLSTKPVLNKLTVYTTLIDVTKGQFETYLRDCPDPCIGW

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-Myc/DDK

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

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

Labeling Method: Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine

Buffer: 25 mM Tris-HCl, 100 mM glycine, pH 7.3

Storage: Store at -80°C. Avoid repeated freeze-thaw cycles.

Stability: Stable for 3 months from receipt of products under proper storage and handling conditions.

RefSeq: NP 004306

RefSeq Size: 2503 RefSeq ORF: 1233

Synonyms: AC; ACDase; ASAH; PHP; PHP32; SMAPME

Locus ID: 427



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UniProt ID: <u>Q13510</u>, <u>Q53H01</u>, <u>A8K0B6</u>

Cytogenetics: 8p22

Summary: This gene encodes a member of the acid ceramidase family of proteins. Alternative splicing

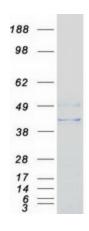
results in multiple transcript variants, at least one of which encodes a preproprotein that is proteolytically processed. Processing of this preproprotein generates alpha and beta subunits that heterodimerize to form the mature lysosomal enzyme, which catalyzes the degradation of ceramide into sphingosine and free fatty acid. This enzyme is overexpressed in multiple human cancers and may play a role in cancer progression. Mutations in this gene are associated with the lysosomal storage disorder, Farber lipogranulomatosis, and a neuromuscular disorder, spinal muscular atrophy with progressive myoclonic epilepsy.

[provided by RefSeq, Oct 2015]

Protein Families: Druggable Genome

Protein Pathways: Lysosome, Metabolic pathways, Sphingolipid metabolism

Product images:



Coomassie blue staining of purified ASAH1 protein (Cat# [TP312434]). The protein was produced from HEK293T cells transfected with ASAH1 cDNA clone (Cat# [RC212434]) using MegaTran 2.0 (Cat# [TT210002]).