Mouse ACY1 / Aminoacylase-1 Protein (His Tag)

Catalog Number: 50008-M08H



General Information

Gene Name Synonym:

1110014J22Rik; Acy-1

Protein Construction:

A DNA sequence encoding the mouse ACY1 (Q99JW2) (Met 1-Ser 408) was expressed with a C-terminal polyhistidine tag.

Source: Mouse

Expression Host: HEK293 Cells

QC Testing

Purity: > 95 % as determined by SDS-PAGE

Bio Activity:

Measured by its ability to cleave N-acetyl-L-Methione (Ac-Met) . The specific activity is >4,000 pmoles/min/ μg .

Endotoxin:

< 1.0 EU per µg of the protein as determined by the LAL method

Stability:

Samples are stable for up to twelve months from date of receipt at -70 $^{\circ}\mathrm{C}$

Predicted N terminal: Gln 18

Molecular Mass:

The recombinant mouse ACY1 consists of 402 amino acids and has a calculated molecular mass of 45 kDa as estimated by SDS-PAGE under reducing conditions.

Formulation:

Lyophilized from sterile PBS, pH 7.4

Normally 5 % - 8 % trehalose, mannitol and 0.01% Tween80 are added as protectants before lyophilization. Specific concentrations are included in the hardcopy of COA. Please contact us for any concerns or special requirements.

Usage Guide

Storage:

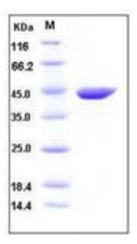
Store it under sterile conditions at -20° C to -80° C upon receiving. Recommend to aliquot the protein into smaller quantities for optimal storage.

Avoid repeated freeze-thaw cycles.

Reconstitution:

Detailed reconstitution instructions are sent along with the products.

SDS-PAGE:



Protein Description

Aminoacylase 1 (ACY1), a metalloenzyme that removes amide-linked ACY1 groups from amino acids and may play a role in regulating responses to oxidative stress. Both the C-terminal fragment found in the two-hybrid screen and full-length ACY1 co-immunoprecipitate with SphK1. Though both C-terminal and full-length proteins slightly reduce SphK1 activity measured in vitro, the C-terminal fragment inhibits while full-length ACY1 potentiates the effects of SphK1 on proliferation and apoptosis. It suggested that ACY1 physically interacts with SphK1 and may influence its physiological functions. As a homodimeric zinc-binding enzyme, Aminoacylase 1 catalyzes the hydrolysis of N alpha-acylated amino acids. Deficiency of Aminoacylase 1 due to mutations in the Aminoacylase 1 (ACY1) gene follows an autosomal-recessive trait of inheritance and is characterized by accumulation of N-acetyl amino acids in the urine.

References

1.Sommer A, et al. (2011) The molecular basis of aminoacylase 1 deficiency. Biochim Biophys Acta. 1812(6): 685-90. 2.Maceyka M, et al. (2004) Aminoacylase 1 is a sphingosine kinase 1-interacting protein. FEBS Lett. 568(1-3): 30-4. 3.Cook RM, et al. (1993) Human aminoacylase-1. Cloning, sequence, and expression analysis of a chromosome 3p21 gene inactivated in small cell lung cancer. J Biol Chem. 268(23): 17010-7.

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