Escherichia coli acnB Protein

Catalog Number: 13102-ENAE



General Information

Gene Name Synonym:

yacl; yacJ

Protein Construction:

A DNA sequence encoding the E.Coli (strain K12) acnB (P36683) (Met 1-Val 865) was expressed and purified.

Source: E. coli

Expression Host: E. coli

QC Testing

Purity: > 97 % as determined by SDS-PAGE

Endotoxin:

Please contact us for more information.

Stability:

Samples are stable for up to twelve months from date of receipt at -70 °C

Predicted N terminal: Met 1

Molecular Mass:

The recombinant E.Coli acnB consisting of 865 amino acids and has a calculated molecular mass of 93.5 kDa. It migrates as an approximately 95 kDa band in SDS-PAGE under reducing conditions.

Formulation:

Lyophilized from sterile 40mM Tris, 1mM DTT, pH 8.2

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:

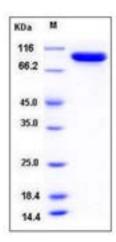
Store it under sterile conditions at -20 $^\circ\!\mathrm{C}$ to -80 $^\circ\!\mathrm{C}$ upon receiving. Recommend to aliquot the protein into smaller quantities for optimal storage.

Avoid repeated freeze-thaw cycles.

Reconstitution:

Detailed reconstitution instructions are sent along with the products.

SDS-PAGE:



Protein Description

Escherichia coli contains two major aconitases (Acns), AcnA and AcnB. They are distantly related monomeric Fe-S proteins that contain different arrangements of four structural domains. acnA is specifically subject to SoxRS-mediated activation, whereas acnB encodes the major aconitase that is synthesized earlier in the growth cycle than AcnA. It is concluded that AcnB is the major citric acid cycle enzyme. Aconitate hydratase 2 (acnB) catalyzes the isomerization of citrate to isocitrate via cis-aconitate as well as the dehydration of 2-methylisocitrate to cis-2-methylaconitate, thus it functions as the major citric-acid-cycle enzyme during exponential growth. Escherichia coli acnB serves as either an enzymic catalyst or a mRNA-binding post-transcriptional regulator, depending on the status of its iron sulfur cluster. AcnB represents a large, distinct group of Gramnegative bacterial aconitases that have an altered domain organization relative to mitochondrial aconitase and other aconitases.

References

1.Bradbury AJ. et al., 1996, 142 (Pt 2): 389-400. 2.Cunningham L. et al., 1997, Microbiology. 143 (Pt 12): 3795-805. 3.Jordan PA. et al., 1999, Biochem J. 344 Pt 3: 739-46.

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