

Recombinant Human AUH, His-tagged

Cat. No. AUH-27419TH Lot. No. (See product label)

SPECIFICATION

Product Overview	Recombinant full length Human AUH with an N-terminal His tag; 293 amino acids with tag, MWt 31.4 kDa.
Species	Human
Source	E.coli
ProteinLength	272 amino acids
Description	The methylglutaconyl-CoA hydratase, mitochondrial protein binds to the AU-rich element (ARE), a common element found in the 3' UTR of rapidly decaying mRNA such as c-fos, c-myc and granulocyte/ macrophage colony stimulating factor. ARE elements are involved in directing RNA to rapid degradation and deadenylation. AUH is also homologous to enol-CoA hydratase, an enzyme involved in fatty acid degradation, and has been shown to have intrinsic hydratase enzymatic activity. AUH is thus a bifunctional chimera between RNA binding and metabolic enzyme activity. A possible subcellular localization in the mitochondria has been demonstrated for the mouse homolog of this protein which shares 92% identity with the human protein. It has been suggested that AUH may have a novel role as a mitochondrial located AU-binding protein. Human AUH is expressed as a single mRNA species of 1.8 kb, and translated as a 40-kDa precursor protein which is subsequently processed to a 32-kDa mature form.
Conjugation	HIS

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Molecular Weight	31.400kDa inclusive of tags
Form	Liquid
Purity	>95% by SDS-PAGE
Storage buffer	Preservative: None Constituents: 20% Glycerol, 0.1M Sodium chloride, 20mM Tris HCl, 1mM DTT, pH 8.0
Storage	Shipped at 4°C. Upon delivery aliquot and store at -20°C or -80°C. Avoid repeated freeze / thaw cycles.
Sequences of amino acids	MGSSHHHHHSSGLVPRGSHMSSEMKTDELVRHLEENRGIVVLGINRAYGKN SLSKNLIKMLSKAVDALKSDKKVRTIIIRSEVPGIFCAGADLKERAKMSSSEVGPVSK IRAVINDIANLPVPTIAAIDGLALGGGLELALACDIRVAASSAKMGLVETKLAIIPGGGG TQRLPRAIGMSLAKELIFSARVLDGKEAKAVGLISHVLEQNQEGDAAYRKALDLAREF LPQGPVAMRVAKLAINQGMEVDLVTGLAIEEACYAQTPTKDRLEGLLAFKEKRPPR YKGE

GENE INFORMATION

Gene Name	AUH AU RNA binding protein/enoyl-CoA hydratase [Homo sapiens]
Official Symbol	AUH
Synonyms	AUH; AU RNA binding protein/enoyl-CoA hydratase; AU RNA binding protein/enoyl Coenzyme A hydratase , AU RNA binding protein/enoyl Coenzyme A hydratase; methylglutaconyl-CoA hydratase, mitochondrial;
Gene ID	549

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mRNA Refseq	NM_001698
Protein Refseq	NP_001689
MIM	600529
Uniprot ID	Q13825
Chromosome Location	9q22
Pathway	Branched-chain amino acid catabolism, organism-specific biosystem; Leucine degradation, leucine => acetoacetate + acetyl-CoA, organism-specific biosystem; Leucine degradation, leucine => acetoacetate + acetyl-CoA, conserved biosystem;
Function	RNA binding; enoyl-CoA hydratase activity; lyase activity; mRNA 3-UTR binding; methylglutaconyl-CoA hydratase activity;