

Recombinant Human EHHADH 293 Cell Lysate

Cat. No. EHHADH-6686HCL **Lot. No.** (See product label)

SPECIFICATION

Product Overview

Species	Human
Source	HEK293
Description	Antigen standard for enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A dehydrogenase (EHHADH), transcript variant 1 is a lysate prepared from HEK293T cells transiently transfected with a TrueORF gene-carrying pCMV plasmid and then lysed in RIPA Buffer. Protein concentration was determined using a colorimetric assay. The antigen control carries a C-terminal Myc/DDK tag for detection.
Components	This product includes 3 vials: 1 vial of gene-specific cell lysate, 1 vial of control vector cell lysate, and 1 vial of loading buffer. Each lysate vial contains 0.1 mg lysate in 0.1 ml (1 mg/ml) of RIPA Buffer (50 mM Tris-HCl pH7.5, 250 mM NaCl, 5 mM EDTA, 50 mM NaF, 1% NP40). The loading buffer vial contains 0.5 ml 2X SDS Loading Buffer (125 mM Tris-Cl, pH6.8, 10% glycerol, 4% SDS, 0.002% Bromophenol blue, 5% beta-mercaptoethanol).
Size	0.1 mg
Storage Instruction	Store at -80°C. Minimize freeze-thaw cycles. After addition of 2X SDS Loading Buffer, the lysates can be stored at -20°C. Product is guaranteed 6 months from the date of shipment.

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Applications

ELISA, WB, IP. WB: Mix equal volume of lysates with 2X SDS Loading Buffer. Boil the mixture for 10 min before loading (for membrane protein lysates, incubate the mixture at room temperature for 30 min). Load 5 ug lysate per lane.

GENE INFORMATION

Gene Name [EHHADH enoyl-CoA, hydratase/3-hydroxyacyl CoA dehydrogenase \[Homo sapiens \]](#)

Official Symbol EHHADH

Synonyms

EHHADH; enoyl-CoA, hydratase/3-hydroxyacyl CoA dehydrogenase; ECHD, enoyl Coenzyme A, hydratase/3 hydroxyacyl Coenzyme A dehydrogenase; peroxisomal bifunctional enzyme; PBE; 3,2-trans-enoyl-CoA isomerase; peroxisomal enoyl-CoA hydratase; L-3-hydroxyacyl-CoA dehydrogenase; L-bifunctional protein, peroxisomal; enoyl-Coenzyme A, hydratase/3-hydroxyacyl Coenzyme A dehydrogenase; LBP; ECHD; LBFP; PBE; L-PBE; MGC120586;

Gene ID [1962](#)

mRNA Refseq [NM_001166415](#)

Protein Refseq [NP_001159887](#)

MIM [607037](#)

UniProt ID [Q08426](#)

Chromosome Location 3q26.3-q28

Pathway Butanoate metabolism, organism-specific biosystem; Butanoate metabolism,

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conserved biosystem; Fatty acid metabolism, organism-specific biosystem; Fatty acid metabolism, conserved biosystem; Lysine degradation, organism-specific biosystem; Lysine degradation, conserved biosystem; Lysine degradation, lysine =>

Function

3-hydroxyacyl-CoA dehydrogenase activity; coenzyme binding; dodecenoyl-CoA delta-isomerase activity; enoyl-CoA hydratase activity; isomerase activity; lyase activity; nucleotide binding; oxidoreductase activity;

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