

Product datasheet for RC230366L4V

OriGene Technologies, Inc.

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EHHADH (NM_001166415) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: EHHADH (NM_001166415) Human Tagged ORF Clone Lentiviral Particle

Symbol: EHHADH

Synonyms: ECHD; FRTS3; L-PBE; LBFP; LBP; MFE1; PBFE

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_001166415

ORF Size: 2172 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC230366).

Sequence:
OTI Disclaimer:

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeq: <u>NM 001166415.1</u>, <u>NP 001159887.1</u>

 RefSeq Size:
 3993 bp

 RefSeq ORF:
 1884 bp

 Locus ID:
 1962

 UniProt ID:
 Q08426

 Cytogenetics:
 3q27.2

Protein Pathways: beta-Alanine metabolism, Butanoate metabolism, Fatty acid metabolism, Limonene and

pinene degradation, Lysine degradation, Metabolic pathways, PPAR signaling pathway, Propanoate metabolism, Tryptophan metabolism, Valine, leucine and isoleucine degradation





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MW: 79.5 kDa

Gene Summary: The

The protein encoded by this gene is a bifunctional enzyme and is one of the four enzymes of the peroxisomal beta-oxidation pathway. The N-terminal region of the encoded protein contains enoyl-CoA hydratase activity while the C-terminal region contains 3-hydroxyacyl-CoA dehydrogenase activity. Defects in this gene are a cause of peroxisomal disorders such as Zellweger syndrome. Two transcript variants encoding different isoforms have been found for this gene. [provided by RefSeq, Oct 2009]