antibodies

Datasheet for ABIN2783232 anti-HADHB antibody (C-Term)

3 Images

2 Publications



Overview

Target:	HADHB
Target Details	
Purification:	Affinity Purified
Characteristics:	This is a rabbit polyclonal antibody against HADHB. It was validated on Western Blot and immunohistochemistry.
Predicted Reactivity:	Cow: 93%, Dog: 100%, Guinea Pig: 93%, Horse: 93%, Human: 100%, Mouse: 100%, Pig: 100%, Rabbit: 93%, Rat: 100%, Yeast: 83%, Zebrafish: 86%
Sequence:	LLLGPTYATP KVLEKAGLTM NDIDAFEFHE AFSGQILANF KAMDSDWFAE
Immunogen:	The immunogen is a synthetic peptide directed towards the C terminal region of human HADHE
Product Details	
Application:	Western Blotting (WB), Immunohistochemistry (IHC)
Conjugate:	This HADHB antibody is un-conjugated
Clonality:	Polyclonal
Host:	Rabbit
Reactivity:	Human, Mouse, Rat, Pig, Zebrafish (Danio rerio), Rabbit, Cow, Dog, Saccharomyces cerevisiae, Guinea Pig, Horse
Binding Specificity:	C-Term
Target:	HADHB
Quantity:	100 µL

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Target Details	
Alternative Name:	HADHB (HADHB Products)
Background:	HADHB is the beta subunit of the mitochondrial trifunctional protein, which catalyzes the last
	three steps of mitochondrial beta-oxidation of long chain fatty acids. The mitochondrial
	membrane-bound heterocomplex is composed of four alpha and four beta subunits, with the
	beta subunit catalyzing the 3-ketoacyl-CoA thiolase activity. Mutations in HADHB gene result in
	trifunctional protein deficiency. The protein can also bind RNA and decreases the stability of
	some mRNAs. This gene encodes the beta subunit of the mitochondrial trifunctional protein,
	which catalyzes the last three steps of mitochondrial beta-oxidation of long chain fatty acids.
	The mitochondrial membrane-bound heterocomplex is composed of four alpha and four beta
	subunits, with the beta subunit catalyzing the 3-ketoacyl-CoA thiolase activity. Mutations in this
	gene result in trifunctional protein deficiency. The encoded protein can also bind RNA and
	decreases the stability of some mRNAs. The genes of the alpha and beta subunits of the
	mitochondrial trifunctional protein are located adjacent to each other in the human genome in a
	head-to-head orientation. Alternatively spliced transcript variants have been found, however,
	their full-length nature is not known. Publication Note: This RefSeq record includes a subset of
	the publications that are available for this gene. Please see the Entrez Gene record to access
	additional publications.
	Alias Symbols: MGC87480, MSTP029, TP-BETA, ECHB, MTPB
	Protein Interaction Partner: HUWE1, TP53, TUBG1, SUMO2, SUMO3, UBC, PARK2, PPP6R1,
	ADRB2, CSNK2A2, PAN2, ATF2, vpu, TIMM8B, COX17, IQCB1, SUOX, CHCHD4, TUBB1, RPS8,
	OPA1, HSP90AB1, HADHA, DES, GRK5, FBX06, CDK2, LEO1, RTF1, TK1, SMN1, PSMA3, GRB7,
	RCC1, CDKN1A, ANXA7, Edc4, MYC, SQST
	Protein Size: 474
Molecular Weight:	47 kDa
Gene ID:	3032
NCBI Accession:	NM_000183, NP_000174
UniProt:	P55084
Pathways:	Monocarboxylic Acid Catabolic Process
Application Details	
Application Notes:	Optimal working dilutions should be determined experimentally by the investigator.

Restrictions: For Research Use only

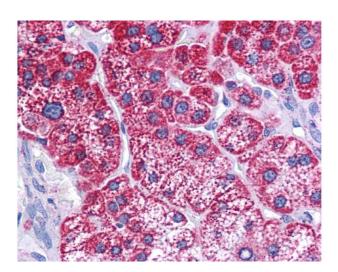
Comment:

Antigen size: 474 AA

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Liquid
Lot specific
Liquid. Purified antibody supplied in 1x PBS buffer with 0.09 % (w/v) sodium azide and 2 % sucrose.
Sodium azide
This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which should be handled by trained staff only.
Avoid repeated freeze-thaw cycles.
-20 °C
For short term use, store at 2-8°C up to 1 week. For long term storage, store at -20°C in small aliquots to prevent freeze-thaw cycles.
Craxton: "Evolutionary genomics of plant genes encoding N-terminal-TM-C2 domain proteins and the similar FAM62 genes and synaptotagmin genes of metazoans." in: BMC genomics , Vol. 8, pp. 259, (2007) (PubMed).

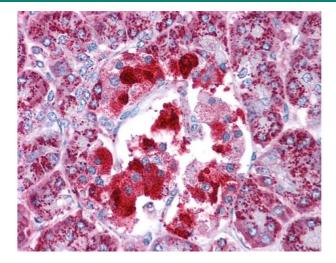
Images



Immunohistochemistry

Image 1.

Images



70 kDa___ 60 kDa___ 48 kDa___ 36 kDa___ 21 kDa___

Immunohistochemistry

Image 2.

Western Blotting

Image 3. WB Suggested Anti-HADHB Antibody Titration: 0.25ug/ml Positive Control: HepG2 cell lysate HADHB is supported by BioGPS gene expression data to be expressed in HepG2

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