

GOT2 monoclonal antibody (M09), clone 4H8

Catalog # H00002806-M09 Size 100 ug

Applications



Western Blot (Cell lysate)

GOT2 monoclonal antibody (M09), clone 4H8. Western Blot analysis of GOT2 expression in HepG2(Cat # L019V1).



Sandwich ELISA (Recombinant protein)

Detection limit for recombinant GST tagged GOT2 is 0.3 ng/ml as a capture antibody.



Western Blot detection against Immunogen (36.74 KDa).

Specification

Product Description

Mouse monoclonal antibody raised against a partial recombinant GOT2.

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Immunogen	GOT2 (NP_002071, 331 a.a. ~ 430 a.a) partial recombinant protein with GST tag. MW of the GST ta g alone is 26 KDa.
Sequence	LNTPDLRKQWLQEVKGMADRIIGMRTQLVSNLKKEGSTHNWQHITDQIGMFCFTGLKPEQVERLIK EFSIYMTKDGRISVAGVTSSNVGYLAHAIHQVTK
Host	Mouse
Reactivity	Human
Interspecies Antigen Sequence	Mouse (93); Rat (93)
Isotype	lgG2b Kappa
Quality Control Testing	Antibody Reactive Against Recombinant Protein. Western Blot detection against Immunogen (36.74 KDa).
Storage Buffer	In 1x PBS, pH 7.4
Storage Instruction	Store at -20°C or lower. Aliquot to avoid repeated freezing and thawing.

Applications

- Western Blot (Cell lysate)
 GOT2 monoclonal antibody (M09), clone 4H8. Western Blot analysis of GOT2 expression in HepG2(Cat # L019V1).
 <u>Protocol Download</u>
- Western Blot (Recombinant protein)

Protocol Download

- Sandwich ELISA (Recombinant protein)
 Detection limit for recombinant GST tagged GOT2 is 0.3 ng/ml as a capture antibody.
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- ELISA

Gene Info — GOT2	
Entrez GenelD	2806

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Product Information

GeneBank Accession#	<u>NM_002080</u>
Protein Accession#	<u>NP_002071</u>
Gene Name	GOT2
Gene Alias	FLJ40994, KAT4, KATIV, mitAAT
Gene Description	glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransferase 2)
Omim ID	<u>138150</u>
Gene Ontology	<u>Hyperlink</u>
Gene Summary	Glutamic-oxaloacetic transaminase is a pyridoxal phosphate-dependent enzyme which exists in c ytoplasmic and inner-membrane mitochondrial forms, GOT1 and GOT2, respectively. GOT plays a role in amino acid metabolism and the urea and tricarboxylic acid cycles. The two enzymes are homodimeric and show close homology. [provided by RefSeq
Other Designations	aspartate aminotransferase 2 kynurenine aminotransferase IV

Pathway

- <u>Alanine</u>
- <u>Arginine and proline metabolism</u>
- Biosynthesis of alkaloids derived from ornithine
- Biosynthesis of phenylpropanoids
- Biosynthesis of plant hormones
- Carbon fixation in photosynthetic organisms
- Cysteine and methionine metabolism
- Isoquinoline alkaloid biosynthesis
- Metabolic pathways
- Novobiocin biosynthesis
- Phenylalanine
- Phenylalanine metabolism
- Tyrosine metabolism