

AASS Polyclonal Antibody

Catalog No :	YT0041
Reactivity :	Human;Rat;Mouse;
Applications :	WB;ELISA
Target :	AASS
Fields :	>>Lysine degradation;>>Metabolic pathways
Gene Name :	AASS
Protein Name :	Alpha-aminoadipic semialdehyde synthase mitochondrial
Human Gene Id :	10157
Human Swiss Prot	Q9UDR5
No :	
Mouse Swiss Prot	Q99K67
NO : Immunogen :	The antiserum was produced against synthesized peptide derived from human AASS. AA range:251-300
Specificity :	AASS Polyclonal Antibody detects endogenous levels of AASS protein.
Formulation :	Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.
Source :	Polyclonal, Rabbit,IgG
Dilution :	WB 1:500 - 1:2000. ELISA: 1:10000. Not yet tested in other applications.
Purification :	The antibody was affinity-purified from rabbit antiserum by affinity- chromatography using epitope-specific immunogen.
Concentration :	1 mg/ml
Storage Stability :	-15°C to -25°C/1 year(Do not lower than -25°C)



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Observed Band :	102kD	
Cell Pathway :	Lysine biosynthesis;Lysine degradation;	
Background :	This gene encodes a bifunctional enzyme that catalyzes the first two steps in the mammalian lysine degradation pathway. The N-terminal and the C-terminal portions of this enzyme contain lysine-ketoglutarate reductase and saccharopine dehydrogenase activity, respectively, resulting in the conversion of lysine to alpha-aminoadipic semialdehyde. Mutations in this gene are associated with familial hyperlysinemia. [provided by RefSeq, Jul 2008],	
Function :	catalytic activity:N(6)-(L-1,3-dicarboxypropyl)-L-lysine + NAD(+) + H(2)O = L- glutamate + 2-aminoadipate 6-semialdehyde + NADH.,catalytic activity:N(6)-(L-1,3-dicarboxypropyl)-L-lysine + NADP(+) + H(2)O = L-lysine + 2-oxoglutarate + NADPH.,disease:Defects in AASS are the cause of hyperlysinemia [MIM:238700]. Hyperlysinemia is an autosomal recessive condition characterized by hyperlysinemia lysinuria and variable saccharopinuria.,function:Bifunctional enzyme that catalyzes the first two steps in lysine degradation. The N-terminal and the C-terminal contain lysine-ketoglutarate reductase and saccharopine dehydrogenase activity, respectively.,induction:Induced by starvation.,pathway:Amino-acid degradation; L- lysine degradation via saccharopine pathway; glutaryl-CoA from L-lysine: step 1/6.,pathway:Amino-acid degradation; L-lysine degradation via saccharopine pathway; glutaryl-CoA from L-lys	
Subcellular Location :	Mitochondrion .	
Expression :	Expressed in all 16 tissues examined with highest expression in the liver.	
Sort :	1564	
No4 :	1	

Products Images









Western blot analysis of the lysates from COLO205 cells using AASS antibody.