

GAPDH Monoclonal Antibody(2B8), FITC Conjugated

CatalogNo: YM2053

Key Features

Host Species Mouse 	Reactivity • Human,Rat,Mouse,Mk,Dg,Ch,Hamster,Rabbit,Pig,sheep,Insect,Yeast	Applications • WB,IF,IHC
Isotype • IgG1	Conjugate • FITC	

Recommended Dilution Ratios

Optimal working dilutions should be determined experimentally by the investigator Suggested starting dilutions are:IF 1:250-1:2000 Flow Cyt 1:250-1:2000

Storage

Storage*	Stable for one year at -15°C to -25°C from date of shipment. For maximum recovery of product, centrifuge the original vial after thawing and prior to removing the cap. Aliquot to avoid repeated freezing and thawing. Store in dark.
Formulation	1mg/ml

Basic Information

Clonality	Monoclonal
Clone Number	2B8

Immunogen Information

Specificity GAPDH Monoclonal Antibody(2B8) FITC conjugated specially designed for your WB or IHC analysis.

Target Information

Gene name	GAPDH			
Protein Name	Glyceraldehyde-3-phosphate dehydrogenase			
	Organism	Gene ID	UniProt ID	
	Human	<u>2597;</u>	<u>P04406;</u>	
Cellular Localization	Cytoplasm, cytosol . Nucleus . Cytopla cytoskeleton . Translocates to the nuc SIAH1, which contains a nuclear local Perinuclear regions (PubMed:1282926	cleus following S-nitrosylation zation signal (By similarity). P	and interaction with	
Tissue specificity	Astrocytoma, Brain, Cajal-Retzius cell, G	Colon adenocarcinoma,Epitheli	iu	
Function	Catalytic activity:D-glyceraldehyde 3-phosphate + phosphate + NAD(+) = 3-phospho-D- glyceroyl phosphate + NADH.,Function:Independent of its glycolytic activity it is also involved in membrane trafficking in the early secretory pathway.,online information:Glyceraldehyde 3-phosphate dehydrogenase entry,pathway:Carbohydrate degradation; glycolysis; pyruvate from D-glyceraldehyde 3-phosphate: step 1.,pathway:Carbohydrate degradation; glycolysis; pyruvate from D-glyceraldehyde 3- phosphate: step 1/5.,PTM:Reversible S-nitrosylation of Cys-152 inhibits enzymatic activity and increases endogenous ADP-ribosylation, which inhibits the enzyme in a non-reversible manner. The latter modification is more likely to be a pathophysiological event associated with inhibition of gluconeogenesis.,sequence Caution:Differs quite extensively.,similarity:Belongs to the glyceraldehyde-3-phosphate dehydrogenase family.,subcellular location:Postnuclear and Perinuclear regions.,subunit:Homotetramer.,subunit:Homotetramer. Binds PRKCI.,			

Validation Data

Contact information

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