

GAPDH (2B8) Mouse mAb (FITC)

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

Liquid in PBS, pH 7.4, containing 0.02% sodium azide as preservative and 50% Glycerol.

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	2597 ;	P04406 ;

Cellular Localization Cytoplasm, cytosol . Nucleus . Cytoplasm, perinuclear region . Membrane . Cytoplasm, cytoskeleton . Translocates to the nucleus following S-nitrosylation and interaction with SIAH1, which contains a nuclear localization signal (By similarity). Postnuclear and Perinuclear regions (PubMed:12829261). .

Tissue specificity Astrocytoma,Brain,Cajal-Retzius cell,Colon adenocarcinoma,Epitheliu

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

Orders: order@immunoway.com
Support: tech@immunoway.com
Telephone: 877-594-3616 (Toll Free), 408-747-0185
Website: <http://www.immunoway.com>
Address: 2200 Ringwood Ave San Jose, CA 95131 USA



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GAPDH (2B8) Mouse mAb (FITC)

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