

GAPDH Monoclonal Antibody(2B8), AbFluor 594 Conjugated

CatalogNo: YM2047

Key Features

Host Species Reactivity Applications
• Mouse • WB,IF,IHC

Human, Rat, Mouse, Mk, Dg, Ch, Hamster, Rabbit, Pig, sheep, Insect, Yeast

Isotype Conjugate
• IgG1 • AbFluor 594

Recommended Dilution Ratios

Optimal working dilutions should be determined experimentally by the investigator Suggested starting dilutions are as follows:IHC 1:200 IF 1:200.

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) AbFluor™ 594 Conjugated specially designed for your

Immunofluorescence 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 . 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

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Please scan the QR code to access additional product information:

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