

# mSHMT (F-11): sc-390641

## BACKGROUND

Mammalian serine hydroxymethyltransferase (SHMT) is a tetrameric, pyridoxal phosphate (PLP)-dependent enzyme that catalyzes the reversible interconversion of serine and tetrahydrofolate to glycine and methylenetetrahydrofolate in the cytoplasm (cSHMT, SHMT1) and mitochondria (mSHMT, SHMT2). cSHMT preferentially supplies one-carbon units for thymidylate biosynthesis, depletes methylenetetrahydrofolate pools for S-adenosylmethionine (SAM) synthesis by synthesizing serine, sequesters 5-methyltetrahydrofolate, and inhibits SAM synthesis. Sheep liver cytosolic recombinant SHMT (scSHMT) Lys71, Arg80 and Asp89 residues influence intra-subunit ionic interactions essential for catalytic activity; Tyr72, Asp227 and His356 residues in the active site interact with PLP and maintain the tetrameric structure. The cDNA for the mitochondrial enzyme encodes a mature protein of 474 residues.

## CHROMOSOMAL LOCATION

Genetic locus: SHMT2 (human) mapping to 12q13.3; Shmt2 (mouse) mapping to 10 D3.

## SOURCE

mSHMT (F-11) is a mouse monoclonal antibody raised against amino acids 435-504 mapping at the C-terminus of mSHMT of human origin.

## PRODUCT

Each vial contains 200 µg IgG<sub>1</sub> kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

mSHMT (F-11) is available conjugated to agarose (sc-390641 AC), 500 µg/0.25 ml agarose in 1 ml, for IP; to HRP (sc-390641 HRP), 200 µg/ml, for WB, IHC(P) and ELISA; to either phycoerythrin (sc-390641 PE), fluorescein (sc-390641 FITC), Alexa Fluor<sup>®</sup> 488 (sc-390641 AF488), Alexa Fluor<sup>®</sup> 546 (sc-390641 AF546), Alexa Fluor<sup>®</sup> 594 (sc-390641 AF594) or Alexa Fluor<sup>®</sup> 647 (sc-390641 AF647), 200 µg/ml, for WB (RGB), IF, IHC(P) and FCM; and to either Alexa Fluor<sup>®</sup> 680 (sc-390641 AF680) or Alexa Fluor<sup>®</sup> 790 (sc-390641 AF790), 200 µg/ml, for Near-Infrared (NIR) WB, IF and FCM.

## APPLICATIONS

mSHMT (F-11) is recommended for detection of mSHMT of mouse, rat and human origin by Western Blotting (starting dilution 1:100, dilution range 1:100-1:1000), immunoprecipitation [1-2 µg per 100-500 µg of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500), immunohistochemistry (including paraffin-embedded sections) (starting dilution 1:50, dilution range 1:50-1:500) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

Suitable for use as control antibody for mSHMT siRNA (h): sc-40942, mSHMT siRNA (m): sc-40943, mSHMT shRNA Plasmid (h): sc-40942-SH, mSHMT shRNA Plasmid (m): sc-40943-SH, mSHMT shRNA (h) Lentiviral Particles: sc-40942-V and mSHMT shRNA (m) Lentiviral Particles: sc-40943-V.

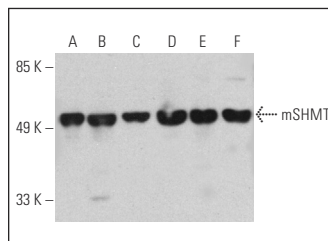
Molecular Weight of mSHMT: 52 kDa.

Positive Controls: A-431 whole cell lysate: sc-2201, c4 whole cell lysate: sc-364186 or Raji whole cell lysate: sc-364236.

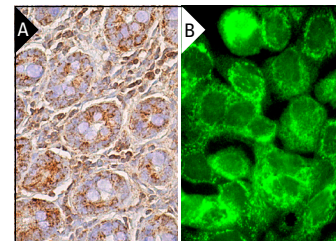
## STORAGE

Store at 4° C, **\*\*DO NOT FREEZE\*\***. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

## DATA



mSHMT (F-11) HRP: sc-390641 HRP. Direct western blot analysis of mSHMT expression in A-431 (A), Raji (B), WEHI-231 (C), EOC 20 (D), F9 (E) and c4 (F) whole cell lysates.



mSHMT (F-11): sc-390641. Immunoperoxidase staining of formalin fixed, paraffin-embedded human colon tissue showing cytoplasmic staining of glandular cells and endothelial cells (A). Immunofluorescence staining of formalin-fixed A-431 cells showing mitochondrial, cytoplasmic and nuclear localization (B).

## SELECT PRODUCT CITATIONS

- Yang, X., et al. 2018. SHMT2 desuccinylation by SIRT5 drives cancer cell proliferation. *Cancer Res.* 78: 372-386.
- Adamus, A., et al. 2018. GCSH antisense regulation determines breast cancer cells' viability. *Sci. Rep.* 8: 15399.
- Moreno-Felici, J., et al. 2019. Phosphoenolpyruvate from glycolysis and PEPCK regulate cancer cell fate by altering cytosolic Ca<sup>2+</sup>. *Cells* 9: 18.
- Son, S.I., et al. 2020. Garcinol is an HDAC11 inhibitor. *ACS Chem. Biol.* 15: 2866-2871.
- Liu, C., et al. 2021. Cytoplasmic SHMT2 drives the progression and metastasis of colorectal cancer by inhibiting β-catenin degradation. *Theranostics* 11: 2966-2986.
- Aslan, M., et al. 2021. Oncogene-mediated metabolic gene signature predicts breast cancer outcome. *NPJ Breast Cancer* 7: 141.
- Pranzini, E., et al. 2022. SHMT2-mediated mitochondrial serine metabolism drives 5-FU resistance by fueling nucleotide biosynthesis. *Cell Rep.* 40: 111233.
- Hwang, Y., et al. 2023. Co-inhibition of glutaminolysis and one-carbon metabolism promotes ROS accumulation leading to enhancement of chemotherapeutic efficacy in anaplastic thyroid cancer. *Cell Death Dis.* 14: 515.

## RESEARCH USE

For research use only, not for use in diagnostic procedures.

## PROTOCOLS

See our web site at [www.scbt.com](http://www.scbt.com) for detailed protocols and support products.

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