## SANTA CRUZ BIOTECHNOLOGY, INC.

# SLMAP (SJ-09): sc-100957



### BACKGROUND

SLMAP (sarcolemmal membrane-associated protein) is a 828 amino acid, single-pass, type IV membrane protein. Localized to the sacrcolemma in non-replicating cells, SLMAP relocates to centrosomes in a microtubule-dependent manner during cleavage in muscle tissues. SLMAP is thought to be involved in protein-protein interactions, as well as play a role in myoblast fusion. As a homodimeric integral membrane protein, SLMAP has two leucine zippers which form a 77 amino acid coiled-coil structure and contains one forkhead-associated (FHA) domain. Coiled-coil structures may be important for the regulation of neurotransmitter release, N-type calcium channels and membrane fusion, while FHA domains are involved in nuclear signaling. SLMAP exists as eight isoforms produced by alternative splicing.

#### REFERENCES

- Bennett, M.K., et al. 1992. Syntaxin: a synaptic protein implicated in docking of synaptic vesicles at presynaptic active zones. Science 257: 255-259.
- Kutay, U., et al. 1993. A class of membrane proteins with a C-terminal anchor. Trends Cell Biol. 3: 72-75.
- Franzini-Armstrong, C. and Jorgensen, A.O. 1994. Structure and development of E-C coupling units in skeletal muscle. Annu. Rev. Physiol. 56: 509-534.

#### CHROMOSOMAL LOCATION

Genetic locus: SLMAP (human) mapping to 3p14.3; SImap (mouse) mapping to 14 A3.

#### SOURCE

SLMAP (SJ-09) is a mouse monoclonal antibody raised against recombinant SLMAP of human origin.

#### PRODUCT

Each vial contains 100  $\mu g$   $lgG_{2a}$  kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

#### **APPLICATIONS**

SLMAP (SJ-09) is recommended for detection of SLMAP of mouse, rat and human origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000), immunoprecipitation [1-2  $\mu$ g per 100-500  $\mu$ g 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 SLMAP siRNA (h): sc-78464, SLMAP siRNA (m): sc-153600, SLMAP shRNA Plasmid (h): sc-78464-SH, SLMAP shRNA Plasmid (m): sc-153600-SH, SLMAP shRNA (h) Lentiviral Particles: sc-78464-V and SLMAP shRNA (m) Lentiviral Particles: sc-153600-V.

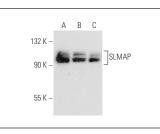
Molecular Weight of SLMAP: 95 kDa.

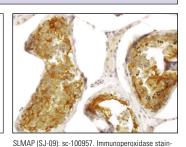
Positive Controls: HeLa whole cell lysate: sc-2200, A549 cell lysate: sc-2413 or HCT-116 whole cell lysate: sc-364175.

#### **RECOMMENDED SUPPORT REAGENTS**

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-IgG $\kappa$  BP-HRP: sc-516102 or m-IgG $\kappa$  BP-HRP (Cruz Marker): sc-516102-CM (dilution range: 1:1000-1:10000), Cruz Marker<sup>TM</sup> Molecular Weight Standards: sc-2035, UltraCruz<sup>®</sup> Blocking Reagent: sc-516214 and Western Blotting Luminol Reagent: sc-2048. 2) Immunoprecipitation: use Protein A/G PLUS-Agarose: sc-2003 (0.5 ml agarose/2.0 ml). 3) Immunofluorescence: use m-IgG $\kappa$  BP-FITC: sc-516140 or m-IgG $\kappa$  BP-PE: sc-516141 (dilution range: 1:50-1:200) with UltraCruz<sup>®</sup> Mounting Medium: sc-24941 or UltraCruz<sup>®</sup> Hard-set Mounting Medium: sc-359850. 4) Immunohistochemistry: use m-IgG $\kappa$  BP-HRP: sc-516102 with DAB, 50X: sc-24982 and Immunohistomount: sc-45086, or Organo/Limonene Mount: sc-45087.

#### DATA





SLMAP (SJ-09): sc-100957. Western blot analysis of SLMAP expression in A549 (**A**), HCT-116 (**B**) and HeLa (**C**) whole cell lysates.

SELECT PRODUCT CITATIONS

ing of formalin-fixed, paraffin-embedded human testis tissue showing membrane and cytoplasmic localization

# 1. Prévilon, M., et al. 2013. Comparative differential proteomic profiles of nonfailing and failing hearts after *in vivo* thoracic aortic constriction in

mice overexpressing FKBP12.6. Physiol. Rep. 1: e00039.

- 2. Tang, Y., et al. 2019. Architecture, substructures, and dynamic assembly of STRIPAK complexes in Hippo signaling. Cell Discov. 5: 3.
- Alì, G., et al. 2019. Whole transcriptome targeted gene quantification provides new insights on pulmonary sarcomatoid carcinomas. Sci. Rep. 9: 3536.

#### **STORAGE**

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

#### **RESEARCH USE**

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

#### PROTOCOLS

See our web site at www.scbt.com for detailed protocols and support products.