

# Sgms1 Antibody - middle region

Rabbit Polyclonal Antibody
Catalog # Al13800

#### **Specification**

# Sgms1 Antibody - middle region - Product Information

Application WB

Primary Accession <u>Q8VCQ6</u>

Other Accession <u>NM\_144792</u>, <u>NP\_659041</u>

Reactivity Human, Mouse, Rat, Rabbit, Pig, Horse,

Bovine, Guinea Pig, Dog

Predicted Human, Mouse, Rat, Rabbit, Pig, Horse,

**Bovine, Guinea Pig, Dog** 

Host Rabbit
Clonality Polyclonal
Calculated MW 46kDa KDa

# Sgms1 Antibody - middle region - Additional Information

**Gene ID 208449** 

Alias Symbol 9530058011Rik, Al841905, C80702, MGC30540, Mob, Sms1, Sor1, Tmem23

#### **Other Names**

Phosphatidylcholine:ceramide cholinephosphotransferase 1, 2.7.8.27, Protein Mob, Sphingomyelin synthase 1, Transmembrane protein 23, Sgms1, Tmem23

#### **Format**

Liquid. Purified antibody supplied in 1x PBS buffer with 0.09% (w/v) sodium azide and 2% sucrose.

#### **Reconstitution & Storage**

Add 50 ul of distilled water. Final anti-Sgms1 antibody concentration is 1 mg/ml in PBS buffer with 2% sucrose. For longer periods of storage, store at 20°C. Avoid repeat freeze-thaw cycles.

#### **Precautions**

Sgms1 Antibody - middle region is for research use only and not for use in diagnostic or therapeutic procedures.

## Sgms1 Antibody - middle region - Protein Information

#### Name Sgms1

Synonyms Tmem23

## **Function**

Major sphingomyelin synthase at the Golgi apparatus. Catalyzes the reversible transfer of phosphocholine moiety in sphingomyelin biosynthesis: in the forward reaction transfers phosphocholine head group of phosphatidylcholine (PC) on to ceramide (CER) to form ceramide



phosphocholine (sphingomyelin, SM) and diacylglycerol (DAG) as by-product, and in the reverse reaction transfers phosphocholine from SM to DAG to form PC and CER. The direction of the reaction depends on the levels of CER and DAG in Golgi membranes. Converts the newly synthesized CER, that is transported from the endoplasmic reticulum to the trans-Golgi by the Cer transport protein (CERT), to SM. Can form a heteromeric complex with glucosylceramide synthase (GCS) increasing SMS activity and reducing glucosylceramide synthesis, a critical mechanism that controls the metabolic fate of CER in the Golgi (By similarity). Does not use free phosphorylcholine or CDP-choline as donor. Can also transfer phosphoethanolamine head group of phosphatidylethanolamine (PE) on to CER to form ceramide phosphoethanolamine (CPE) (PubMed:<a href="http://www.uniprot.org/citations/25605874" target="\_blank">25605874</a>). Regulates receptor-mediated signal transduction via mitogenic DAG and proapoptotic CER, as well as via SM, a structural component of membrane rafts that serve as platforms for signal transduction and protein sorting (PubMed:<a href="http://www.uniprot.org/citations/16879426" target=" blank">16879426</a>, PubMed:<a href="http://www.uniprot.org/citations/22580896" target="blank">22580896</a>). Plays a role in secretory transport via regulation of DAG pool at the Golgi apparatus and its downstream effects on PRKD1 (By similarity).

#### **Cellular Location**

Golgi apparatus membrane {ECO:0000250|UniProtKB:Q86VZ5}; Multi-pass membrane protein

#### **Tissue Location**

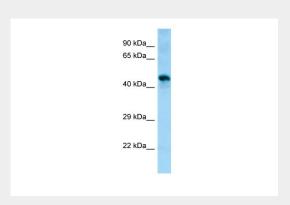
Isoform 1 is widely expressed, isoform 2 shows a more narrow distribution and isoform 3 is detected only in testis and heart.

# Sgms1 Antibody - middle region - Protocols

Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- Immunoprecipitation
- Flow Cytomety
- Cell Culture

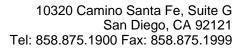
## Sgms1 Antibody - middle region - Images



WB Suggested Anti-Sgms1 Antibody Titration: 1.0 µg/ml

Positive Control: Mouse Muscle

Sgms1 Antibody - middle region - References





Yang Z.,et al.Gene 363:123-132(2005).
Carninci P.,et al.Science 309:1559-1563(2005).
Huitema K.,et al.EMBO J. 23:33-44(2004).
Yang Z.,et al.FEMS Yeast Res. 6:751-762(2006).
Li Z.,et al.Arterioscler. Thromb. Vasc. Biol. 32:1577-1584(2012).