

# SAM68 Rabbit mAb

**Catalog # AP77070** 

### **Specification**

#### SAM68 Rabbit mAb - Product Information

Application
Primary Accession
Reactivity
Host
Clonality
Calculated MW

WB
O07666
Human, Mouse, Rat
Rabbit
Monoclonal Antibody
48227

#### SAM68 Rabbit mAb - Additional Information

Gene ID 10657

Other Names KHDRBS1

**Dilution** WB~~1/500-1/1000

Format Liquid

### SAM68 Rabbit mAb - Protein Information

Name KHDRBS1 (HGNC:18116)

### **Function**

Recruited and tyrosine phosphorylated by several receptor systems, for example the T-cell, leptin and insulin receptors. Once phosphorylated, functions as an adapter protein in signal transduction cascades by binding to SH2 and SH3 domain-containing proteins. Role in G2-M progression in the cell cycle. Represses CBP-dependent transcriptional activation apparently by competing with other nuclear factors for binding to CBP. Also acts as a putative regulator of mRNA stability and/or translation rates and mediates mRNA nuclear export. Positively regulates the association of constitutive transport element (CTE)-containing mRNA with large polyribosomes and translation initiation. According to some authors, is not involved in the nucleocytoplasmic export of unspliced (CTE)-containing RNA species according to (PubMed:<a

href="http://www.uniprot.org/citations/22253824" target="\_blank">22253824</a>). RNA-binding protein that plays a role in the regulation of alternative splicing and influences mRNA splice site selection and exon inclusion. Binds to RNA containing 5'-[AU]UAA- 3' as a bipartite motif spaced by more than 15 nucleotides. Binds poly(A). Can regulate CD44 alternative splicing in a Ras pathway-dependent manner (By similarity). In cooperation with HNRNPA1 modulates alternative splicing of BCL2L1 by promoting splicing toward isoform Bcl-X(S), and of SMN1 (PubMed:<a

href="http://www.uniprot.org/citations/17371836" target="\_blank">17371836</a>, PubMed:<a href="http://www.uniprot.org/citations/20186123" target="\_blank">20186123</a>). Can regulate alternative splicing of NRXN1 and NRXN3 in the laminin G-like domain 6 containing the





Tel: 858.875.1900 Fax: 858.875.1999

evolutionary conserved neurexin alternative spliced segment 4 (AS4) involved in neurexin selective targeting to postsynaptic partners. In a neuronal activity-dependent manner cooperates synergistically with KHDRBS2/SLIM-1 in regulation of NRXN1 exon skipping at AS4. The cooperation with KHDRBS2/SLIM-1 is antagonistic for regulation of NXRN3 alternative splicing at AS4 (By similarity).

### **Cellular Location**

Nucleus. Cytoplasm Membrane. Note=Predominantly located in the nucleus but also located partially in the cytoplasm

### **Tissue Location**

Ubiquitously expressed in all tissue examined. Isoform 1 is expressed at lower levels in brain, skeletal muscle, and liver whereas isoform 3 is intensified in skeletal muscle and in liver

### SAM68 Rabbit mAb - Protocols

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

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- <u>Immunoprecipitation</u>
- Flow Cytomety
- Cell Culture

## SAM68 Rabbit mAb - Images

