

#### OriGene Technologies, Inc.

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# Product datasheet for CF808889

## SAMSN1 Mouse Monoclonal Antibody [Clone ID: OTI1F6]

### **Product data:**

Product Type:	Primary Antibodies
Clone Name:	OTI1F6
Applications:	WB
Recommended Dilution:	WB 1:500~2000
Reactivity:	Human
Host:	Mouse
lsotype:	lgG1
Clonality:	Monoclonal
Immunogen:	Human recombinant protein fragment corresponding to amino acids 1-250 of human SAMSN1(NP_071419) produced in E.coli.
Formulation:	Lyophilized powder (original buffer 1X PBS, pH 7.3, 8% trehalose)
Reconstitution Method:	For reconstitution, we recommend adding 100uL distilled water to a final antibody concentration of about 1 mg/mL. To use this carrier-free antibody for conjugation experiment, we strongly recommend performing another round of desalting process. (OriGene recommends Zeba Spin Desalting Columns, 7KMWCO from Thermo Scientific)
Purification:	Purified from mouse ascites fluids or tissue culture supernatant by affinity chromatography (protein A/G)
Conjugation:	Unconjugated
Storage:	Store at -20°C as received.
Stability:	Stable for 12 months from date of receipt.
Predicted Protein Size:	41.5 kDa
Gene Name:	SAM domain, SH3 domain and nuclear localization signals 1
Database Link:	<u>NP_071419</u> <u>Entrez Gene 64092 Human</u> <u>Q9NSI8</u>



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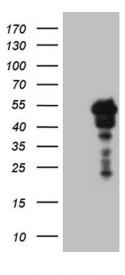
#### **GRIGENE** SAMSN1 Mouse Monoclonal Antibody [Clone ID: OTI1F6] – CF808889

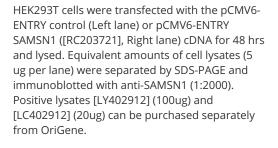
Background:SAMSN1 is a member of a novel gene family of putative adaptors and scaffold proteins<br/>containing SH3 and SAM (sterile alpha motif) domains (Claudio et al., 2001 [PubMed<br/>11536050]). [supplied by OMIM, Mar 2008]. Transcript Variant: This variant (2) contains<br/>alternate exons at the 5' end and initiates translation from an alternate start codon<br/>compared to variant 1. The resulting longer isoform (2) has a distinct N-terminus compared to<br/>isoform 1. Sequence Note: This RefSeq record was created from transcript and genomic<br/>sequence data to make the sequence consistent with the reference genome assembly. The<br/>genomic coordinates used for the transcript record were based on transcript alignments.<br/>##Evidence-Data-START## Transcript exon combination :: AF218085.2 [ECO:0000332]<br/>RNAseq introns :: mixed/partial sample support ERS025081, ERS025082 [ECO:0000350]<br/>##Evidence-Data-END## COMPLETENESS: complete on the 3' end.

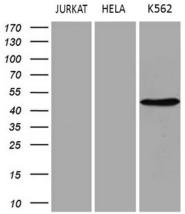
Synonyms:

HACS1; NASH1; SASH2; SH3D6B; SLy2

### **Product images:**







Western blot analysis of extracts (35ug) from 3 different cell lines by using anti-SAMSN1 monoclonal antibody (1:500).

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