

## **Product datasheet for PH305428**

## OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

## KHDRBS2 (NM\_152688) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** KHDRBS2 MS Standard C13 and N15-labeled recombinant protein (NP\_689901)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC205428

or AA Sequence: Predicted MW:

38.9 kDa

Protein Sequence: >RC205428 protein sequence

Red=Cloning site Green=Tags(s)

MEEEKYLPELMAEKDSLDPSFVHASRLLAEEIEKFQGSDGKKEDEEKKYLDVISNKNIKLSERVLIPVKQ YPKFNFVGKLLGPRGNSLKRLQEETGAKMSILGKGSMRDKAKEEELRKSGEAKYAHLSDELHVLIEVFAP PGEAYSRMSHALEEIKKFLVPDYNDEIRQEQLRELSYLNGSEDSGRGRGIRGRGIRIAPTAPSRGRGGAI PPPPPPGRGVLTPRGSTVTRGALPVPPVARGVPTPRARGAPTVPGYRAPPPPAHEAYEEYGYDDGYGGEY DDQTYETYDNSYATQTQSVPEYYDYGHGVSEDAYDSYAPEEWATTSSSLKAPPQRSARGGYREHPYGRY

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

Tag: C-Myc/DDK

**Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining

**Concentration:** >0.05 μg/μL as determined by microplate BCA method

Labeling Method: Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine

**Buffer:** 25 mM Tris-HCl, 100 mM glycine, pH 7.3

**Storage:** Store at -80°C. Avoid repeated freeze-thaw cycles.

**Stability:** Stable for 3 months from receipt of products under proper storage and handling conditions.

**RefSeq:** NP 689901

RefSeq Size: 2336 RefSeq ORF: 1047

Synonyms: SLM-1; SLM1

**Locus ID:** 202559



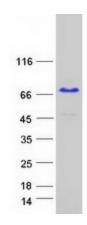
UniProt ID: Q5VWX1

**Cytogenetics:** 6q11.1

**Summary:** RNA-binding protein that plays a role in the regulation of alternative splicing and influences

mRNA splice site selection and exon inclusion. Binds both poly(A) and poly(U) homopolymers. Phosphorylation by PTK6 inhibits its RNA-binding ability (By similarity). Induces an increased concentration-dependent incorporation of exon in CD44 pre-mRNA by direct binding to purine-rich exonic enhancer. Can regulate alternative splicing of NRXN1 in the laminin G-like domain 6 containing the evolutionary conserved neurexin alternative spliced segment 4 (AS4) involved in neurexin selective targeting to postsynaptic partners. Regulates cell-type specific alternative splicing of NRXN1 at AS4 and acts synergystically with SAM68 in exon skipping. In contrast acts antagonistically with SAM68 in NRXN3 exon skipping at AS4. Its phosphorylation by FYN inhibits its ability to regulate splice site selection. May function as an adapter protein for Src kinases during mitosis.[UniProtKB/Swiss-Prot Function]

Product images:



Coomassie blue staining of purified KHDRBS2 protein (Cat# [TP305428]). The protein was produced from HEK293T cells transfected with KHDRBS2 cDNA clone (Cat# [RC205428]) using MegaTran 2.0 (Cat# [TT210002]).