

## **Product datasheet for PH307607**

## OriGene Technologies, Inc.

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## RBM4 (NM\_002896) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** RBM4 MS Standard C13 and N15-labeled recombinant protein (NP\_002887)

Species:HumanExpression Host:HEK293

**Expression cDNA Clone** 

ne RC207607

or AA Sequence:

Predicted MW: 40.3 kDa

**Protein Sequence:** >RC207607 protein sequence

Red=Cloning site Green=Tags(s)

MVKLFIGNLPREATEQEIRSLFEQYGKVLECDIIKNYGFVHIEDKTAAEDAIRNLHHYKLHGVNINVEAS KNKSKTSTKLHVGNISPTCTNKELRAKFEEYGPVIECDIVKDYAFVHMERAEDAVEAIRGLDNTEFQGKR MHVQLSTSRLRTAPGMGDQSGCYRCGKEGHWSKECPIDRSGRVADLTEQYNEQYGAVRTPYTMSYGDSLY YNNAYGALDAYYKRCRAARSYEAVAAAAASVYNYAEQTLSQLPQVQNTAMASHLTSTSLDPYDRHLLPTS GAAATAAAAAAAAAAAAAVTAASTSYYGRDRSPLRRATAPVPTVGEGYGYGHESELSQASAAARNSLYDMARY

**EREQYADRARYSAF** 

**SGPTRTRPL**EQKLISEEDLAANDILDYKDDDDK**V** 

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

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 002887

RefSeq Size: 1714 RefSeq ORF: 1092

Synonyms: LARK; RBM4A; ZCCHC21; ZCRB3A

**Locus ID:** 5936





Cytogenetics:

UniProt ID: Q9BWF3

11q13.2

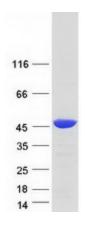
**Summary:** RNA-binding factor involved in multiple aspects of cellular processes like alternative splicing

of pre-mRNA and translation regulation. Modulates alternative 5'-splice site and exon

selection. Acts as a muscle cell differentiation-promoting factor. Activates exon skipping of the PTB pre-mRNA during muscle cell differentiation. Antagonizes the activity of the splicing factor PTBP1 to modulate muscle cell-specific exon selection of alpha tropomyosin. Binds to intronic pyrimidine-rich sequence of the TPM1 and MAPT pre-mRNAs. Required for the translational activation of PER1 mRNA in response to circadian clock. Binds directly to the 3' UTR of the PER1 mRNA. Exerts a suppressive activity on Cap-dependent translation via binding to CU-rich responsive elements within the 3' UTR of mRNAs, a process increased under stress conditions or during myocytes differentiation. Recruits EIF4A1 to stimulate IRES-dependent translation initiation in respons to cellular stress. Associates to internal ribosome entry segment (IRES) in target mRNA species under stress conditions. Plays a role for miRNA-guided RNA cleavage and translation suppression by promoting association of AGO2-containing miRNPs with their cognate target mRNAs. Associates with miRNAs during muscle cell differentiation. Binds preferentially to 5'-CGCGCG[GCA]-3' motif in vitro.[UniProtKB/Swiss-Prot Function]

**Protein Families:** Druggable Genome

## **Product images:**



Coomassie blue staining of purified RBM4 protein (Cat# [TP307607]). The protein was produced from HEK293T cells transfected with RBM4 cDNA clone (Cat# [RC207607]) using MegaTran 2.0 (Cat# [TT210002]).