

Product datasheet for PH308648

OriGene Technologies, Inc.

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METT10D (METTL16) (NM 024086) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

METT10D MS Standard C13 and N15-labeled recombinant protein (NP_076991) **Description:**

Species: Human **HEK293 Expression Host: Expression cDNA Clone**

or AA Sequence:

RC208648

Predicted MW: 63.6 kDa

>RC208648 protein sequence **Protein Sequence:**

Red=Cloning site Green=Tags(s)

MALSKSMHARNRYKDKPPDFAYLASKYPDFKQHVQINLNGRVSLNFKDPEAVRALTCTLLREDFGLSIDI PLERLIPTVPLRLNYIHWVEDLIGHQDSDKSTLRRGIDIGTGASCIYPLLGATLNGWYFLATEVDDMCFN YAKKNVEQNNLSDLIKVVKVPQKTLLMDALKEESEIIYDFCMCNPPFFANQLEAKGVNSRNPRRPPPSSV NTGGITEIMAEGGELEFVKRIIHDSLQLKKRLRWYSCMLGKKCSLAPLKEELRIQGVPKVTYTEFCQGRT MRWALAWSFYDDVTVPSPPSKRRKLEKPRKPITFVVLASVMKELSLKASPLRSETAEGIVVVTTWIEKIL TDLKVQHKRVPCGKEEVSLFLTAIENSWIHLRRKKRERVRQLREVPRAPEDVIQALEEKKPTPKESGNSQ ELARGPQERTPCGPALREGEAAAVEGPCPSQESLSQEENPEPTEDERSEEKGGVEVLENCQGSSNGAQDQ EASEOFGSPVAERGKRLPGVAGQYLFKCLINVKKEVDDALVEMHWVEGQNRDLMNQLCTYIRNQIFRLVA

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

C-Myc/DDK Tag:

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

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

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

RefSeq: NP 076991

RefSeq Size: 5758 RefSeq ORF: 1686



METT10D (METTL16) (NM_024086) Human Mass Spec Standard - PH308648

Synonyms: METT10D

 Locus ID:
 79066

 UniProt ID:
 Q86W50

 Cytogenetics:
 17p13.3

Summary: RNA N6-methyltransferase that methylates adenosine residues at the N(6) position of a

subset of RNAs and is involved in S-adenosyl-L-methionine homeostasis by regulating expression of MAT2A transcripts (PubMed:28525753, PubMed:30197299, PubMed:30197297).

Able to N6-methylate a subset of mRNAs and U6 small nuclear RNAs (U6 snRNAs)

(PubMed:28525753). In contrast to the METTL3-METTL14 heterodimer, only able to methylate a limited number of RNAs: requires both a 5'UACAGAGAA-3' nonamer sequence and a specific RNA structure (PubMed:28525753, PubMed:30197299, PubMed:30197297). Plays a key role in S-adenosyl-L-methionine homeostasis by mediating N6-methylation of MAT2A mRNAs, altering splicing and/or stability of MAT2A transcripts: in presence of S-adenosyl-L-

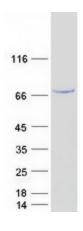
methionine, binds the 3' UTR region of MAT2A mRNA and specifically N6-methylates the first hairpin of MAT2A mRNA, impairing MAT2A expression (PubMed:28525753). In S-adenosyl-L-methionine-limiting conditions, binds the 3' UTR region of MAT2A mRNA but stalls due to the lack of a methyl donor, preventing N6-methylation and promoting expression of MAT2A (PubMed:28525753). In addition to mRNAs, also able to mediate N6-methylation of U6 small nuclear RNA (U6 snRNA): specifically N6-methylates adenine in position 43 of U6 snRNAs

(PubMed:28525753, PubMed:29051200). Also able to bind various lncRNAs

(PubMed:29051200). Specifically binds the 3'-end of the MALAT1 long non-coding RNA

(PubMed:27872311).[UniProtKB/Swiss-Prot Function]

Product images:



Coomassie blue staining of purified METTL16 protein (Cat# [TP308648]). The protein was produced from HEK293T cells transfected with METTL16 cDNA clone (Cat# [RC208648]) using MegaTran 2.0 (Cat# [TT210002]).