



Product Data Sheet

METTL1 Protein, Human (His)

Cat. No.: HY-P75926

Synonyms: tRNA (guanine-N(7)-)-methyltransferase; Methyltransferase-like protein 1; METTL1; C12orf1

Species: E. coli Source:

Q9UBP6 (D32-Q265) Accession:

Gene ID: 4234

Molecular Weight: Approximately 30 kDa

PROPERTIES

AA Sequence	
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DHTLRYPVKP EEMDWSELYP EFFAPLTQNQ SHDDPKDKKE KRAQAQVEFA DIGCGYGGLL VELSPLFPDT LILGLEIRVK VSDYVQDRIR ALRAAPAGGF QNIACLRSNA MKHLPNFFYK GQLTKMFFLF PDPHFKRTKH KWRIISPTLL AEYAYVLRVG GLVYTITDVL LEDLSEDPVV ELHDWMCTHF EEHPLFERVP FPAIFRRIQD GHLGTSTEEG KKVLRNGGKN PVLQ

Biological Activity

Measured in a cell proliferation assay using HepG2 cells. The ED₅₀ for this effect is 0.9259 ng/mL, corresponding to a specific activity is 1.080×10⁶ units/mg.

Appearance

Solution

Formulation

Supplied as a 0.2 µm filtered solution of 50 mM Tris, 0.5 M NaCl, 20% Glycerol, pH 8.0.

Endotoxin Level

<1 EU/µg, determined by LAL method.

Reconsititution

N/A.

Storage & Stability

Stored at -80°C for 1 year. It is stable at -20°C for 3 months after opening. It is recommended to freeze aliquots at -80°C for extended storage. Avoid repeated freeze-thaw cycles.

Shipping

Shipping with dry ice

DESCRIPTION

Background

METTL1, the catalytic component of the METTL1-WDR4 methyltransferase complex, is instrumental in mediating the formation of N(7)-methylguanine in various RNA species, including tRNAs, mRNAs, and microRNAs (miRNAs). Specifically, METTL1 catalyzes the addition of N(7)-methylguanine at position 46 (m7G46) within a significant subset of tRNAs containing the 5'-RAGGU-3' motif in the variable loop. This modification, such as m7G46, stabilizes tRNA tertiary structure and shields

tRNAs from decay. METTL1 also serves as a methyltransferase for internal N(7)-methylguanine in mRNAs, particularly in response to stress, leading to the relocalization of methylated mRNAs to stress granules and consequent translational suppression. Furthermore, METTL1 methylates specific miRNAs, including let-7, facilitating let-7 miRNA processing by disrupting inhibitory secondary structures within primary miRNA transcripts. Beyond its role in RNA modification, METTL1 emerges as a regulator of embryonic stem cell self-renewal and differentiation.

Caution: Product has not been fully validated for medical applications. For research use only.

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