

# METTL3 Knockdown U2OS cell line, Heterozygous

Catalog No.: RM50187

## Basic Information

### Catalog No.

RM50187

### Category

Cell Lysate

### Parental Cell line

U2OS

### Genotype

Knockdown

## Gene Information

### Gene Symbol

METTL3

### Species

Human

### Gene ID

56339

### Swiss Prot

Q86U44

### Synonyms

M6A; IME4; Spo8; MT-A70; hMETTL3;  
METTL3

## Contact

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## Background

This gene encodes the 70 kDa subunit of MT-A which is part of N6-adenosine-methyltransferase. This enzyme is involved in the posttranscriptional methylation of internal adenosine residues in eukaryotic mRNAs, forming N6-methyladenosine.

## Product Information

### Description

METTL3 Knockdown cell line is engineered from U2OS cell line with Gene-Editing Technology.

Allele-1:139bp deletion in exon2

Allele-2:138bp deletion in exon2

Mammalian cells such as human, rat and mouse cells are normally diploid with two alleles. Homozygote: both alleles were knocked out, mRNA has no signal, no expression of proteins. Heterozygote: only one allele was knocked out, the mRNA transcript levels was decreased compared to wild type, and the protein expression levels was also lower than that of the wild type.

### Packaging

1 vial parental cell line and 1 vial knockout cell line

### Shipping Conditions

Dry ice

### Amount

1~5x10<sup>6</sup> cells/vial.

### Storage

Stored in liquid nitrogen for a long time less than -130°C. Minimizing freeze-thaw cycles.

### Protocol

Upon arrival, it should be maintained in DMEM medium with 10%(v/v) fetal bovine serum and 100U penicillin-streptomycin, at 37°C with 5% CO<sub>2</sub> condition.

1. Thaw the vial in 37°C water bath, and shake it to melt as soon as possible.
2. Transfer the cell suspension to a 15mL conical tube with pre-warmed 5mL complete medium and centrifuge 1000rpm for approximately 5 minutes at room temperature.
3. Remove and discard the supernatant.
4. Resuspend the cell pellet with 1mL pre-warmed complete medium and seed in 10cm dish.
5. Add 8-10mL of complete medium.
6. Incubate the culture at 37°C incubator with 5% CO<sub>2</sub>.
7. A subcultivation ratio of 1:2-1:4 is recommended.

## Sequencing data

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WT CATTGCTCCAACC\*\*\*\*\*CATTGCCCACTGAT  
Mut CATTGCTCCAACC\*\*\*Deletion\*\*\*CATTGCCCACTGAT  
Allele-1: 139bp deletion in exon2

Genome sequence analysis of PCR products from parental (WT) and METTL3 knockdown (KD) U2OS cells, using sanger sequencing.

WT ATTGCTCCAACCT\*\*\*\*\*CATTGCCCACTGAT  
Mut ATTGCTCCAACCT\*\*\*Deletion\*\*\*CATTGCCCACTGAT  
Allele-2: 138bp deletion in exon2