

# TET3 Knockout 293T Cell Line, Homozygous

Catalog No.: RM01887

## Basic Information

### Catalog No.

RM01887

### Category

Cell Line

### Parental Cell line

293T

### Genotype

Knockout

## Gene Information

### Gene Symbol

TET3

### Species

Human

### Gene ID

200424

### Swiss Prot

O43151

### Synonyms

hCG\_40738

## Contact

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## Background

Members of the ten-eleven translocation (TET) gene family, including TET3, play a role in the DNA methylation process (Langemeijer et al., 2009 [PubMed 19923888]).[supplied by OMIM, Nov 2010]

## Product Information

### Description

TET3 Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing Technology.

Allele-1:194bp deletion in exon3

Allele-2:196bp deletion in exon3

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   GGAAGATGCCACG\*\*\*\*\*CGGCATGGTATGAA  
Mut   GGAAGATGCCACG\*\*\*Deletion\*\*\*CGGCATGGTATGAA  
Allele-1: 194bp deletion in exon3

WT   CCTGGAAGATGCC\*\*\*\*\*\*GCGGCATGGTATGA  
Mut   CCTGGAAGATGCC\*\*\*Deletion\*\*\*GCGGCATGGTATGA  
Allele-2: 196bp deletion in exon3

Genome sequence analysis of PCR products from parental (WT) and TET3 knockout (KO) 293T cells, using sanger sequencing.