

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

☎ | 400-999-6126

✉ | cn.market@abclonal.com.cn

🌐 | www.abclonal.com.cn

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

1~5x10⁶ 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% CO₂ 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% CO₂.
7. A subcultivation ratio of 1:2-1:4 is recommended.

Sequencing data

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.