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# **EMRE Knockout 293T Cell Line, Homozygous**

Catalog No.: RM02149

### **Basic Information**

## Catalog No.

RM02149

## Category

Cell Line

#### **Parental Cell line**

293T

#### Genotype

Knockout

# **Gene Information**

### **Gene Symbol**

**EMRE** 

#### **Species**

Human

#### **Gene ID**

91689

#### **Swiss Prot**

Q9H4I9

# Synonyms

C22orf32; DDDD; EMRE; dJ186O1.1

#### **Contact**

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# **Background**

### **Product Information**

#### Description

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

Allele-1:55bp deletion in exon1

Allele-2:56bp deletion in exon1

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**

**Amount** 

Dry ice

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

#### Storage

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

#### Protoco

Upon arrival, it should be maintained in DMEM medium with 10%(v/v) fetal bovine serum and 100U penicillin-streptomycin, at  $37^{\circ}C$  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

WT CGATGTCTCCGCCG\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*\*CAGCGGCGCCATTT
Mut CGATGTCTCCGCCG\*\*\*Deletion\*\*\*CAGCGGCGCCATTT
Allele-1: 55bp deletion in exon1

WT GCGATGTCTCCGCC\*\*\*\*\*\*\*\*\*CAGCGGCGCCATTT
Mut GCGATGTCTCCGCC\*\*\*Deletion\*\*\*CAGCGGCGCCATTT

Allele-2: 56bp deletion in exon1

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