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EEF1B2 Knockout 293T Cell Line, Homozygous

Catalog No.: RM02235

Basic Information

Catalog No.

RM02235

Category

Cell Line

Parental Cell line

293T

Genotype

Knockout

Background

This gene encodes a translation elongation factor. The protein is a guanine nucleotide exchange factor involved in the transfer of aminoacylated tRNAs to the ribosome. Alternative splicing results in three transcript variants which differ only in the 5' UTR. [provided by RefSeq, Jul 2008]

Gene Information

Gene Symbol

EEF1B2

Species

Human

Gene ID

1933

Swiss Prot

P24534

Synonyms

EEF1B; EEF1B1; EF1B

Contact

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Product Information

Description

 ${\tt EEF1B2}\ {\tt Knockout}\ {\tt cell\ line}\ {\tt is\ engineered\ from\ 293T\ cell\ line\ with\ Gene-Editing\ Technology.}$ ${\tt Allele-1:exon1\ was\ deleted}$

Allele-2:exon1 was deleted

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

 $1\sim5x10^6$ 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_2 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 CCTTTTTCCTCTCT***********GGACGGGCTGAGTC Mut CCTTTTTCCTCTCT***Deletion***GGACGGGCTGAGTC Allele-1: exon1 was deleted

WT CCTTTTTCCTCTCT****GGACGGGCTGAGTC
Mut CCTTTTTCCTCTCT***Deletion***GGACGGGCTGAGTC

Allele-2: exon1 was deleted

Genome sequence analysis of PCR products from parental (WT) and EEF1B2 knockout (KO) 293T cells, using sanger sequencing.