ABclonal www.abclonal.com

FZD5 Knockout 293T Cell Line, Homozygous

Catalog No.: RM02739

Basic Information

Catalog No.

RM02739

Category

Cell Line

Parental Cell line

293T

Genotype

Knockout

Background

Members of the 'frizzled' gene family encode 7-transmembrane domain proteins that are receptors for Wnt signaling proteins. The FZD5 protein is believed to be the receptor for the Wnt5A ligand.

Gene Information

Gene Symbol

FZD5

Species

Human

Gene ID

7855

Swiss Prot

Q13467

Synonyms

HFZ5; C2orf31; FZD5

Contact

<u>a</u>	400-999-6126
×	cn.market@abclonal.com.cn
⊙	www.abclonal.com.cn

Product Information

Description

FZD5 Knockout cell line is engineered from 293T cell line with Gene-Editing Technology. Allele-1:113bp deletion in exon1

Allele-2:112bp 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

1~5x10⁶ 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% 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 ACGGTGCCCATGTG************CGCCGGACCTGCG
Mut ACGGTGCCCATGTG***Deletion***CGCCGGACCTGCG
Allele-1: 113bp deletion in exon1

WT ACGGTGCCCATGTG***********TCGCCGGACCTGC
Mut ACGGTGCCCATGTG***Deletion***TCGCCGGACCTGC
Allele-2: 112bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and FZD5 knockout (KO) 293T cells, using sanger sequencing.