

FAM50A Knockout NIH/3T3 Cell Line, Homozygous

Catalog No.: RM02182

Basic Information

Catalog No.

RM02182

Category

Cell Line

Parental Cell line

NIH/3T3

Genotype

Knockout

Gene Information

Gene Symbol

FAM50A

Species

Mouse

Gene ID

108160

Synonyms

DOHXS9928E; XAP-5

Contact

 | 400-999-6126

 | cn.market@abclonal.com.cn

 | www.abclonal.com.cn

Background

Product Information

Description

FAM50A Knockout NIH/3T3 Cell Line is engineered from NIH/3T3 cell line with Gene-Editing Technology.

Allele-1:exon2 was deleted

Allele-2:exon2 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

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 GTTAAGGCA*TCGA*****TGCTCACTCTTTGTC
Mut GTTA-GGCA*TCGA***Deletion***TGCTCACTCTTTGTC
Allele-1: exon2 was deleted

Genome sequence analysis of PCR products from parental (WT) and FAM50A knockout (KO) NIH/3T3 cells, using sanger sequencing.

WT GTTAAGGCA*TCGA*****TGCTCACTCTTTGTC
Mut GTTA-GGCA*TCGA***Deletion***TGCTCACTCTTTGTC
Allele-2: exon2 was deleted