

# FAM160B2 Knockdown HuH-7 Cell Line, Heterozygous

Catalog No.: RM02234

### **Basic Information**

### Catalog No.

RM02234

### Category

Cell Line

### **Parental Cell line**

HuH-7

### Genotype

Knockdown

### **Gene Information**

### **Gene Symbol**

FAM160B2

### **Species**

Human

#### **Gene ID**

64760

### **Swiss Prot**

Q86V87

### Synonyms

RAI16

### **Contact**

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|-----------|---------------------------|
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### **Background**

### **Product Information**

#### Description

FAM160B2 Knockdown HuH-7 cell line is engineered from HuH-7 cell line with Gene-Editing Technology.

Allele-1:2bp deletion in exon3

Allele-2:141bp 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**

 ${\bf 1}$  vial parental cell line and  ${\bf 1}$  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 TGAAAGCACCCCGGCCAAGAAGACAGACAT Mut TGAAAGCACCCCCG - -AAGAAGACAGACAT

Allele-1: 2bp deletion in exon3

WT GAAAGCACCCCGC\*\*\*\*\*\*\*\*\*\*\*\*\*GCTGGGCAAGGCCG
Mut GAAAGCACCCCCGC\*\*\*Deletion\*\*\*GCTGGGCAAGGCCG

Allele-2: 141bp deletion in exon3

Genome sequence analysis of PCR products from parental (WT) and FAM160B2 Knockdown (KD) Huh-7 cells, using sanger sequencing.