

# SETD2 Knockdown 293T Cell Line, Heterozygous

Catalog No.: RM02147

## Basic Information

### Catalog No.

RM02147

### Category

Cell Line

### Parental Cell line

293T

### Genotype

Knockdown

## Gene Information

### Gene Symbol

SETD2

### Species

Human

### Gene ID

29072

### Swiss Prot

Q9BYW2

### Synonyms

HBP231; HIF-1; HIP-1; HSPC069; HYPB;  
KMT3A; LLS; SET2; p231HBP

## Contact

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## Background

Huntington's disease (HD), a neurodegenerative disorder characterized by loss of striatal neurons, is caused by an expansion of a polyglutamine tract in the HD protein huntingtin. This gene encodes a protein belonging to a class of huntingtin interacting proteins characterized by WW motifs. This protein is a histone methyltransferase that is specific for lysine-36 of histone H3, and methylation of this residue is associated with active chromatin. This protein also contains a novel transcriptional activation domain and has been found associated with hyperphosphorylated RNA polymerase II. [provided by RefSeq, Aug 2008]

## Product Information

### Description

SETD2 Knockdown 293T Cell Line is engineered from 293T cell line with Gene-Editing Technology.

Allele-1:203bp deletion in exon3

Allele-2:207bp 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

1 vial parental cell line and 1 vial knockout cell line

### Shipping Conditions

Dry ice

### Amount

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

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WT    CCAGCTGTACCTCTT\*\*\*\*\*GATAGCAGAATCAA  
Mut    CCAGCTGTACCTCTT\*\*\*Deletion\*\*\*GATAGCAGAATCAA  
Allele-1: 203bp deletion in exon3

WT    CCAGCTGTACCTCTT\*\*\*\*\*AGCAGAATCAACAA  
Mut    CCAGCTGTACCTGG\*\*\*Deletion\*\*\*AGCAGAATCAACAA  
Allele-2: 207bp deletion in exon3

Genome sequence analysis of PCR products from parental (WT) and SETD2 Knockdown (KD) 293T cells, using sanger sequencing.