

# FTO Knockout 293T Cell Lysate, Homozygous

Catalog No.: RM50208

## Basic Information

**Catalog No.**

RM50208

**Category**

Cell Lysate

**Parental Cell line**

293T

**Genotype**

Knockout

## Gene Information

**Gene Symbol**

58kDa

**Species**

Human

**Gene ID**

79068

**Swiss Prot**

Q9C0B1

**Synonyms**

ALKBH9; BMIQ14; GDFD

## Contact

 | 400-999-6126

 | [cn.market@abclonal.com.cn](mailto:cn.market@abclonal.com.cn)

 | [www.abclonal.com.cn](http://www.abclonal.com.cn)

## Background

This gene is a nuclear protein of the AlkB related non-haem iron and 2-oxoglutarate-dependent oxygenase superfamily but the exact physiological function of this gene is not known. Other non-heme iron enzymes function to reverse alkylated DNA and RNA damage by oxidative demethylation. Studies in mice and humans indicate a role in nervous and cardiovascular systems and a strong association with body mass index, obesity risk, and type 2 diabetes. [provided by RefSeq, Jul 2011]

## Product Information

**Description**

FTO Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing technology.

Allele-1:50bp deletion in exon3

Allele-2:52bp 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 Lysate and 1 vial knockout cell Lysate

**Shipping Conditions**

4°C

**Amount**

50µL, 2µg/µL.

**Storage**

Lysate is stable for 12 months when stored at -20°C. Minimizing freeze-thaw cycles.

**Protocol**

To be used as WB control. Lysate is supplied in 1× SDS sample buffer (2% SDS, 60 mM Tris-HCl pH 6.8, 10% Glycerol, 0.02% Bromophenol blue, 60 mM beta-mercaptoethanol). Lysate should be boiled for 3 - 5 minutes before loading onto gel.

## Sequencing data

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WT GCTTATTCGGG\*\*\*\*\*CTCATTGGTAA  
Mut GCTTATTCGGG\*\*\*Deletion(50bp)\*\*\*CTCATTGGTAA  
Allele-1: 50bp deletion in exon3

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

WT GCTTATTCGGG\*\*\*\*\*CATTGGTAATCC  
Mut GCTTATTCGGG\*\*\*Deletion(52bp)\*\*\*CATTGGTAATCC  
Allele-2: 52bp deletion in exon3