

FTO Knockout 293T Cell Lysate, Homozygous

Catalog No.: RM02270

Basic Information

Catalog No.

RM02270

Category

Cell Lysate

Parental Cell line

293T

Genotype

Knockout

Gene Information

Gene Symbol

FTO

Species

Human

Gene ID

79068

Swiss Prot

Q9C0B1

Synonyms

ALKBH9; BMIQ14; GDFD

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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-1:52bp 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

WT CTGCTTATTCGGG*****CATTGGTAATCCAG
Mut CTGCTTATTCGGG***Deletion***CATTGGTAATCCAG
Allele-1: 52bp deletion in exon3
WT CTGCTTATTCGGG*****CATTGGTAATCCAG
Mut CTGCTTATTCGGG***Deletion***CATTGGTAATCCAG
Allele-2: 52bp deletion in exon3

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