

FOXO1 Knockdown HeLa Cell Lysate, Heterozygous

Catalog No.: RM02050

Basic Information

Catalog No.

RM02050

Category

Cell Lysate

Parental Cell line

HeLa

Genotype

Knockdown

Gene Information

Gene Symbol

FOXO1

Species

Human

Gene ID

2308

Swiss Prot

Q12778

Synonyms

FKH1; FKHR; FOXO1A

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Background

This gene belongs to the forkhead family of transcription factors which are characterized by a distinct forkhead domain. The specific function of this gene has not yet been determined; however, it may play a role in myogenic growth and differentiation. Translocation of this gene with PAX3 has been associated with alveolar rhabdomyosarcoma. [provided by RefSeq, Jul 2008]

Product Information

Description

FOXO1 Knockdown HeLa Cell Line is engineered from HeLa cell line with Gene-Editing technology.

Allele-1:1bp deletion in exon1

Allele-2:12bp deletion in exon1

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 CGGCCGCGCGGCCGCCACCGGGGGCTGTGCGGGGAC
Mut CGGCCGCGCGGCCGCCA -CGGGGGCTGTGCGGGGAC
Allele-1: 1bp deletion in exon1

WT GGCGGCGCGCCG*****GGGGCTGTGCGGG
Mut GGCGGCGCGCCG***Deletion***GGGGCTGTGCGGG
Allele-2: 12bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and FOXO1 Knockdown (KD) HeLa cells, using sanger sequencing.