

BAZ1B Knockout 293T Cell Lysate, Homozygous

Catalog No.: RM02488

Basic Information

Catalog No.

RM02488

Category

Cell Lysate

Parental Cell line

293T

Genotype

Knockout

Gene Information

Gene Symbol

BAZ1B

Species

Human

Gene ID

9031

Swiss Prot

Q9UIG0

Synonyms

WBSCR10; WBSCR9; WSTF

Contact

 | 400-999-6126

 | cn.market@abclonal.com.cn

 | www.abclonal.com.cn

Background

This gene encodes a member of the bromodomain protein family. The bromodomain is a structural motif characteristic of proteins involved in chromatin-dependent regulation of transcription. This gene is deleted in Williams-Beuren syndrome, a developmental disorder caused by deletion of multiple genes at 7q11.23. [provided by RefSeq, Jul 2008]

Product Information

Description

BAZ1B Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing technology.

Allele-1:73bp deletion in exon1

Allele-2:73bp 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 AGGAGTTTCCTGCC*****GGTTGGAGATCATG
Mut AGGAGTTTCCTGCC**Deletion**GGTTGGAGATCATG
Allele-1: 73bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and BAZ1B knockout (KO) 293T cells, using sanger sequencing.

WT AGGAGTTTCCTGCC*****GGTTGGAGATCATG
Mut AGGAGTTTCCTGCC**Deletion**GGTTGGAGATCATG
Allele-2: 73bp deletion in exon1