

# 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](mailto:cn.market@abclonal.com.cn)

🌐 | [www.abclonal.com.cn](http://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

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

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

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