

# **BNIP3 Knockout 293T Cell Lysate, Homozygous**

Catalog No.: RM02480

### **Basic Information**

#### Catalog No.

RM02480

### Category

Cell Lysate

### **Parental Cell line**

293T

### Genotype

Knockout

### **Background**

This gene is encodes a mitochondrial protein that contains a BH3 domain and acts as a proapoptotic factor. The encoded protein interacts with anti-apoptotic proteins, including the E1B 19 kDa protein and Bcl2. This gene is silenced in tumors by DNA methylation. [provided by RefSeq, Dec 2014]

### **Gene Information**

### **Gene Symbol**

BNIP3

#### **Species**

Human

### Gene ID

664

### **Swiss Prot**

Q12983

### **Synonyms**

NIP3

### **Contact**

<u>a</u>		400-999-6126
$\bowtie$		cn.market@abclonal.com.cn
•	Τ	www.abclonal.com.cn

### **Product Information**

#### Description

BNIP3 Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing technology.

Allele-1:134bp deletion in exon1

Allele-2:134bp 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**

 ${\bf 1}$  vial parental cell Lysate and  ${\bf 1}$  vial knockout cell Lysate

 $\begin{array}{lll} \textbf{Shipping Conditions} & \textbf{Amount} \\ 4^{\circ} C & 50 \mu\text{L}, 2 \mu\text{g}/\mu\text{L}. \end{array}$ 

### Storage

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

#### Protoco

To be used as WB control. Lysate is supplied in  $1\times$  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 TGAGTTCCTCCGGC\*\*\*\*\*\*\*\*\*\*\*\*\*\*GCCATGTCGCAGAA
Mut TGAGTTCCTCCGGC\*\*\*Deletion\*\*\*GCCATGTCGCAGAA
Allele-1: 134bp deletion in exon1

WT TGAGTTCCTCCGGC\*\*\*\*\*\*\*\*\*\*\*\*\*GCCATGTCGCAGAA
Mut TGAGTTCCTCCGGC\*\*\*Deletion\*\*\*GCCATGTCGCAGAA

Allele-2: 134bp deletion in exon1

Genome sequence analysis of PCR products from parental (WT) and BNIP3 knockout (KO) 293T cells, using sanger sequencing.