

# ENOX2 Knockout 293T Cell Lysate, Homozygous

**Catalog No.:** RM02310

## Basic Information

### Catalog No.

RM02310

### Category

Cell Lysate

### Parental Cell line

293T

### Genotype

Knockout

## Gene Information

### Gene Symbol

ENOX2

### Species

Human

### Gene ID

10495

### Swiss Prot

Q16206

### Synonyms

APK1; COVA1; tNOX

## Contact

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## Background

This gene is a tumor-specific member of the ECTO-NOX family of genes that encode cell surface NADH oxidases. The encoded protein has two enzymatic activities: catalysis of hydroquinone or NADH oxidation, and protein disulfide interchange. The protein also displays prion-like properties. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Aug 2013]

## Product Information

### Description

ENOX2 Knockout 293T Cell Line is engineered from 293T cell line with Gene-Editing technology.

Allele-1:143bp deletion in exon4

Allele-2:143bp deletion in exon4

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 TCCCACCTCCTGCA\*\*\*\*\*TCGCTTTGCTGAGG  
Mut TCCCACCTCCTGCA\*\*\*Deletion\*\*\*TCGCTTTGCTGAGG  
Allele-1: 143bp deletion in exon4  
  
WT TCCCACCTCCTGCA\*\*\*\*\*TCGCTTTGCTGAGG  
Mut TCCCACCTCCTGCA\*\*\*Deletion\*\*\*TCGCTTTGCTGAGG  
Allele-2: 143bp deletion in exon4

Genome sequence analysis of PCR products from parental (WT) and ENOX2 Knockout (KO) 293T cells, using sanger sequencing.