SMAD4 Knockdown 293T Cell Lysate, Heterozygous

Catalog No.: RM02042



Basic Information

Catalog No. RM02042

Category Cell Lysate

Parental Cell line 293T

Genotype Knockdown

Gene Information

Gene Symbol SMAD4

Species Human

Gene ID 4089

Swiss Prot Q13485

Synonyms DPC4; JIP; MADH4; MYHRS

Contact

6	400-999-6126
\bowtie	cn.market@abclonal.com.cn
€	www.abclonal.com.cn

Background

This gene encodes a member of the Smad family of signal transduction proteins. Smad proteins are phosphorylated and activated by transmembrane serine-threonine receptor kinases in response to TGF-beta signaling. The product of this gene forms homomeric complexes and heteromeric complexes with other activated Smad proteins, which then accumulate in the nucleus and regulate the transcription of target genes. This protein binds to DNA and recognizes an 8-bp palindromic sequence (GTCTAGAC) called the Smad-binding element (SBE). The Smad proteins are subject to complex regulation by post-translational modifications. Mutations or deletions in this gene have been shown to result in pancreatic cancer, juvenile polyposis syndrome, and hereditary hemorrhagic telangiectasia syndrome. [provided by RefSeq, Oct 2009]

Product Information

Description

SMAD4 Knockdown 293T Cell Line is engineered from 293T cell line with Gene-Editing technology. Allele-1:139bp deletion in exon1

Allele-2:141bp 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℃	

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 $1 \times$ 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 TGTGCCATAGACAA***********AGAGAACATTGGAT Mut TGTGCCATAGACAA***Deletion***AGAGAACATTGGAT Allele-1: 139bp deletion in exon1

WT TGTGCCATAGACAA**********AGAACATTGGATGG Mut TGTGCCATAGACAA***Deletion***AGAACATTGGATGG Allele-2: 141bp deletion in exon1 Genome sequence analysis of PCR products from parental (WT) and SMAD4 Knockdown (KD) 293T cells, using sanger sequencing.