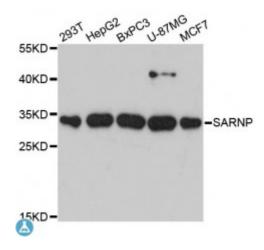


Anti-SARNP Antibody



Description This gene encodes a protein that is upregulated in response to various

cytokines. The encoded protein may play a role in cell cycle progression. A translocation between this gene and the myeloid/lymphoid leukemia gene, resulting in expression of a chimeric protein, has been associated with acute myelomonocytic leukemia. Pseudogenes exist on chromosomes 7 and 8. Alternatively spliced transcript variants have been described.

Model STJ115656

Host Rabbit

Reactivity Human

Applications WB

Immunogen Recombinant fusion protein containing a sequence corresponding to amino

acids 1-210 of human SARNP (NP_149073.1).

Gene ID 84324

Gene Symbol SARNP

Dilution range WB 1:500 - 1:2000

Tissue Specificity Low expression in spleen, liver, pancreas, testis, thymus, heart, and kidney,

Increased levels are seen in hepatocellular carcinoma and pancreatic

adenocarcinoma

Purification Affinity purification

Note For Research Use Only (RUO).

Protein Name SAP domain-containing ribonucleoprotein Cytokine-induced protein of 29

kDa Nuclear protein Hcc-1 Proliferation-associated cytokine-inducible protein

CIP29

Molecular Weight 23.671 kDa

Clonality Polyclonal

Conjugation Unconjugated

Isotype IgG

Formulation PBS with 0.02% sodium azide, 50% glycerol, pH7.3.

Storage Instruction Store at -20C. Avoid freeze / thaw cycles.

Database Links HGNC:244320MIM:610049Reactome:R-HSA-109688

Alternative Names SAP domain-containing ribonucleoprotein Cytokine-induced protein of 29

kDa Nuclear protein Hcc-1 Proliferation-associated cytokine-inducible protein

CIP29

Function Binds both single-stranded and double-stranded DNA with higher affinity for

the single-stranded form, Specifically binds to scaffold/matrix attachment region DNA, Also binds single-stranded RNA, Enhances RNA unwinding activity of DDX39A, May participate in important transcriptional or translational control of cell growth, metabolism and carcinogenesis, Component of the TREX complex which is thought to couple mRNA

transcription, processing and nuclear export, and specifically associates with spliced mRNA and not with unspliced pre-mRNA, TREX is recruited to spliced mRNAs by a transcription-independent mechanism, binds to mRNA upstream of the exon-junction complex (EJC) and is recruited in a splicing-and cap-dependent manner to a region near the 5' end of the mRNA where it functions in mRNA export to the cytoplasm via the TAP/NFX1 pathway, The TREX complex is essential for the export of Kaposi's sarcoma-associated herpesvirus (KSHV) intronless mRNAs and infectious virus production,

Cellular Localization Nucleus, Nucleus speckle

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