

Pinin Polyclonal Antibody

Catalog # AP71914

Product Information

Application WB Primary Accession Q9H307

Reactivity Human, Mouse

HostRabbitClonalityPolyclonalCalculated MW81628

Additional Information

Gene ID 5411

Other Names PNN; DRS; MEMA; Pinin; 140 kDa nuclear and cell adhesion-related

phosphoprotein; Desmosome-associated protein; Domain-rich serine protein; DRS protein; DRSP; Melanoma metastasis clone A protein; Nuclear protein

SDK3; SR-like protein

Dilution WB~~Western Blot: 1/500 - 1/2000. ELISA: 1/20000. Not yet tested in other

applications.

Format Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.09% (W/V) sodium

azide.

Storage Conditions -20°C

Protein Information

Name PNN

Synonyms DRS, MEMA

Function Transcriptional activator binding to the E-box 1 core sequence of the

E-cadherin promoter gene; the core-binding sequence is 5'CAGGTG-3'. Capable of reversing CTBP1-mediated transcription repression. Auxiliary component of the splicing-dependent multiprotein exon junction complex (EJC) deposited at splice junction on mRNAs. The EJC is a dynamic structure consisting of core proteins and several peripheral nuclear and cytoplasmic associated factors that join the complex only transiently either during EJC assembly or during subsequent mRNA metabolism. Participates in the regulation of alternative pre-mRNA splicing. Associates to spliced mRNA within 60 nt upstream of the 5'-splice sites. Component of the PSAP complex which binds RNA in a sequence-independent manner and is proposed to be recruited to the EJC prior to or during the splicing process and to regulate specific excision of introns in specific transcription subsets. Involved in the

establishment and maintenance of epithelia cell-cell adhesion. Potential tumor suppressor for renal cell carcinoma.

Cellular Location Nucleus speckle. Cell junction, desmosome. Note=Cell-cell contact area,

predominantly desmosome of intercellular adherens junction. Not a

nucleocytoplasmic shuttling protein

Tissue Location Expressed in placenta, lung, liver, kidney, pancreas, spleen, thymus, prostate,

testis, ovary, small intestine, colon, heart, epidermis, esophagus, brain and smooth and skeletal muscle. Expressed strongly in melanoma metastasis

lesions and advanced primary tumors.

Background

Transcriptional activator binding to the E-box 1 core sequence of the E-cadherin promoter gene; the core-binding sequence is 5'CAGGTG-3'. Capable of reversing CTBP1-mediated transcription repression. Auxiliary component of the splicing- dependent multiprotein exon junction complex (EJC) deposited at splice junction on mRNAs. The EJC is a dynamic structure consisting of core proteins and several peripheral nuclear and cytoplasmic associated factors that join the complex only transiently either during EJC assembly or during subsequent mRNA metabolism. Participates in the regulation of alternative pre-mRNA splicing. Associates to spliced mRNA within 60 nt upstream of the 5'-splice sites. Component of the PSAP complex which binds RNA in a sequence-independent manner and is proposed to be recruited to the EJC prior to or during the splicing process and to regulate specific excision of introns in specific transcription subsets. Involved in the establishment and maintenance of epithelia cell- cell adhesion. Potential tumor suppressor for renal cell carcinoma.

Images



Western Blot analysis of various cells using Pinin Polyclonal Antibody

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