

TudorSN Polyclonal Antibody

Catalog No: YT5084

Reactivity: Human; Mouse; Rat

Applications: WB;IHC;IF;ELISA

Target: TudorSN

Fields: >>Viral carcinogenesis

Gene Name: SND1

Protein Name: Staphylococcal nuclease domain-containing protein 1

Human Gene ld: 27044

Human Swiss Prot

Q7KZF4

No:

Mouse Gene ld: 56463

Mouse Swiss Prot

Q78PY7

No:

Rat Gene ld: 64635

Rat Swiss Prot No: Q66X93

Immunogen: Synthesized peptide derived from the Internal region of human TudorSN.

Specificity: TudorSN Polyclonal Antibody detects endogenous levels of TudorSN protein.

Formulation : Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Source: Polyclonal, Rabbit, IgG

Dilution: WB 1:500 - 1:2000. IHC: 1:100-300 ELISA: 1:40000.. IF 1:50-200

Purification: The antibody was affinity-purified from rabbit antiserum by affinity-

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chromatography using epitope-specific immunogen.

Concentration: 1 mg/ml

Storage Stability: -15°C to -25°C/1 year(Do not lower than -25°C)

Observed Band: 101kD

Background: This gene encodes a transcriptional co-activator that interacts with the acidic

domain of Epstein-Barr virus nuclear antigen 2 (EBNA 2), a transcriptional activator that is required for B-lymphocyte transformation. Other transcription factors that interact with this protein are signal transducers and activators of transcription, STATs. This protein is also thought to be essential for normal cell growth. A similar protein in mammals and other organisms is a component of the

RNA-induced silencing complex (RISC). [provided by RefSeq, Jul 2016],

Function: function: Functions as a bridging factor between STAT6 and the basal

transcription factor. Plays a role in PIM1 regulation of MYB activity. Functions as

a transcriptional coactivator for the Epstein-Barr virus nuclear antigen 2

(EBNA2).,PTM:Phosphorylated by PIM1 in vitro.,sequence caution:The frameshift leads to wrong initation.,similarity:Contains 1 Tudor domain.,similarity:Contains 4 TNase-like domains.,subcellular location:In IL-4 stimulated cells colocalizes with STAT6 in the nucleus. Identified by mass spectrometry in melanosome fractions from stage I to stage IV.,subunit:Binds to acidic transactivation domain of EBNA2. Interacts with EAV NSP1. Interacts with GTF2E1 and GTF2E2. Forms a ternary

complex with STAT6 and POLR2A. Interacts with STAT5.,tissue

specificity: Ubiquitously expressed.,

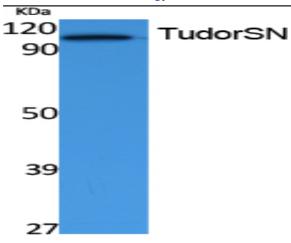
Subcellular Location : Cytoplasm . Nucleus . Melanosome . In IL-4 stimulated cells colocalizes with STAT6 in the nucleus (PubMed:12234934). Identified by mass spectrometry in

melanosome fractions from stage I to stage IV (PubMed:17081065). .

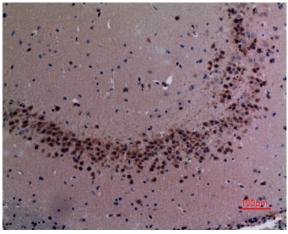
Expression: Ubiquitously expressed.

Products Images

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Western Blot analysis of extracts from Jurkat cells, using TudorSN Polyclonal Antibody. Secondary antibody(catalog#:RS0002) was diluted at 1:20000



Immunohistochemical analysis of paraffin-embedded mousebrain, antibody was diluted at 1:100