

Product datasheet for TA382215

Spt6 (SUPT6H) Rabbit Polyclonal Antibody

Product data:

Product Type: Primary Antibodies

Applications: ICC/IF, IP, WB

Recommended Dilution: WB,1:500 - 1:2000

IF,1:50 - 1:200 IP,1:50 - 1:200

Reactivity: Human, Mouse, Rat

Modifications: Unmodified

Host: Rabbit

Isotype: IgG

Clonality: Polyclonal

Immunogen: Recombinant fusion protein containing a sequence corresponding to amino acids 1500-1726

of human SUPT6H (NP_003161.2).

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

Concentration: lot specific

Purification: Affinity purification

Conjugation: Unconjugated

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

Stability: Shelf life: one year from despatch.

Predicted Protein Size: 61kDa/84kDa/199kDa

Gene Name: SPT6 homolog, histone chaperone

Database Link: Entrez Gene 6830 Human

Q7KZ85



OriGene Technologies, Inc. 9620 Medical Center Drive, Ste 200

CN: techsupport@origene.cn

Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com



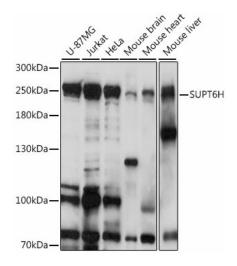
Background:

Transcription elongation factor which binds histone H3 and plays a key role in the regulation of transcription elongation and mRNA processing. Enhances the transcription elongation by RNA polymerase II (RNAPII and is also required for the efficient activation of transcriptional elongation by the HIV-1 nuclear transcriptional activator, Tat. Besides chaperoning histones in transcription, acts to transport and splice mRNA by forming a complex with IWS1 and the C-terminal domain (CTD of the RNAPII subunit RPB1 (POLR2A. The SUPT6H:IWS1:CTD complex recruits mRNA export factors (ALYREF/THOC4, EXOSC10 as well as histone modifying enzymes (such as SETD2, to ensure proper mRNA splicing, efficient mRNA export and elongation-coupled H3K36 methylation, a signature chromatin mark of active transcription. SUPT6H via its association with SETD1A, regulates both class-switch recombination and somatic hypermutation through formation of H3K4me3 epigenetic marks on activation-induced cytidine deaminase (AICDA target loci. Promotes the activation of the myogenic gene program by entailing erasure of the repressive H3K27me3 epigenetic mark through stabilization of the chromatin interaction of the H3K27 demethylase KDM6A.

Synonyms:

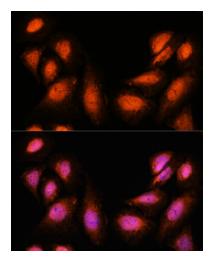
emb-5; hSPT6; KIAA0162; MGC87943; SPT6; SPT6H

Product images:

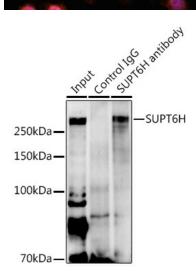


Western blot analysis of extracts of various cell lines, using SUPT6H antibody (TA382215) at 1:1000 dilution. | Secondary antibody: HRP Goat Anti-Rabbit IgG (H+L) at 1:10000 dilution. | Lysates/proteins: 25ug per lane. | Blocking buffer: 3% nonfat dry milk in TBST. | Detection: ECL Basic Kit. | Exposure time:





Immunofluorescence analysis of U-2 OS cells using SUPT6H antibody (TA382215) at dilution of 1:100. Blue: DAPI for nuclear staining.



Immunoprecipitation analysis of 600ug extracts of Mouse heart using 3ug SUPT6H antibody (TA382215). Western blot was performed from the immunoprecipitate using SUPT6H antibody (TA382215) at a dilition of 1:500.