



# SPT16 Monoclonal Antibody

<b>Catalog No</b>	YP-mAb-02038
<b>Isotype</b>	IgG
<b>Reactivity</b>	Human;Mouse
<b>Applications</b>	WB
<b>Gene Name</b>	SUPT16H
<b>Protein Name</b>	FACT complex subunit SPT16
<b>Immunogen</b>	The antiserum was produced against synthesized peptide derived from human SUPT16H. AA range:941-990
<b>Specificity</b>	SPT16 Monoclonal Antibody detects endogenous levels of SPT16 protein.
<b>Formulation</b>	Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.
<b>Source</b>	Monoclonal, Mouse,IgG
<b>Purification</b>	The antibody was affinity-purified from mouse antiserum by affinity-chromatography using epitope-specific immunogen.
<b>Dilution</b>	WB 1:500-1:2000
<b>Concentration</b>	1 mg/ml
<b>Purity</b>	≥90%
<b>Storage Stability</b>	-20°C/1 year
<b>Synonyms</b>	SUPT16H; FACT140; FACTP140; FACT complex subunit SPT16; Chromatin-specific transcription elongation factor 140 kDa subunit; FACT 140 kDa subunit; FACTp140; Facilitates chromatin transcription complex subunit SPT16; hSPT16
<b>Observed Band</b>	119kD
<b>Cell Pathway</b>	Nucleus . Chromosome . Colocalizes with RNA polymerase II on chromatin. Recruited to actively transcribed loci.
<b>Tissue Specificity</b>	Ubiquitous.
<b>Function</b>	caution:Although related to the peptidase M24 family, this protein lacks conserved active site residues suggesting that it may lack peptidase activity.,domain:The Glu-rich acidic region in C-terminus is essential for FACT activity.,function:Component of the FACT complex, a general chromatin factor that acts to reorganize nucleosomes. The FACT complex is involved in multiple processes that require DNA as a template such as mRNA elongation, DNA replication and DNA repair. During transcription elongation the FACT complex acts as a histone chaperone that both destabilizes and restores nucleosomal structure. It facilitates the passage of RNA polymerase II and transcription by promoting the dissociation of one histone H2A-H2B dimer from the nucleosome, then subsequently promotes the reestablishment of the nucleosome following the



passage of RNA polymerase II. The FACT complex is probably also

#### Background

Transcription of protein-coding genes can be reconstituted on naked DNA with only the general transcription factors and RNA polymerase II. However, this minimal system cannot transcribe DNA packaged into chromatin, indicating that accessory factors may facilitate access to DNA. One such factor, FACT (facilitates chromatin transcription), interacts specifically with histones H2A/H2B to effect nucleosome disassembly and transcription elongation. FACT is composed of an 80 kDa subunit and a 140 kDa subunit; this gene encodes the 140 kDa subunit. [provided by RefSeq, Feb 2009],

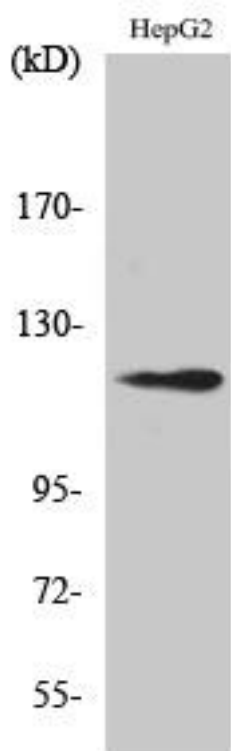
#### matters needing attention

Avoid repeated freezing and thawing!

#### Usage suggestions

This product can be used in immunological reaction related experiments. For more information, please consult technical personnel.

### Products Images



Western Blot analysis of various cells using SPT16 Monoclonal Antibody