## CPTC-MEN1-1 (CAB080226)

Uniprot ID: 000255

Protein name: MEN1\_HUMAN

Full name: Menin

Tissue specificity: Ubiquitous.

**Function**: Essential component of a MLL/SET1 histone methyltransferase (HMT) complex, a complex that specifically methylates 'Lys-4' of histone H3 (H3K4). Functions as a transcriptional regulator. Binds to the TERT promoter and represses telomerase expression. Plays a role in TGFB1-mediated inhibition of cell-proliferation, possibly regulating SMAD3 transcriptional activity. Represses JUND-mediated transcriptional activation on AP1 sites, as well as that mediated by NFKB subunit RELA. Positively regulates HOXC8 and HOXC6 gene expression. May be involved in normal hematopoiesis through the activation of HOXA9 expression (By similarity). May be involved in DNA repair.

Subcellular location:

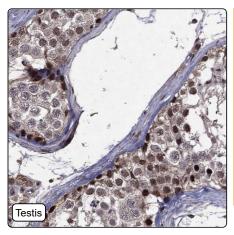
Nucleus (experimental evidence)

NOTE: Concentrated in nuclear body-like structures. Relocates to the nuclear matrix upon gamma irradiation.

Protein existence: Experimental evidence at protein level

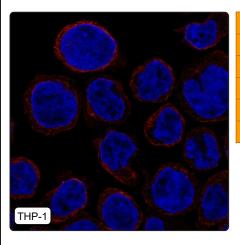
Comment:

## Immunohistochemistry



IHC protocol:	HIER pH6, Dilution 1:300	
IHC test staining:	Nuclear positivity in testis.	
Literature conformance:	Partly consistent with extensive gene/protein characterization data	
Literature significance:		
RNA similarity:	Very low consistency between antibody staining and RNA expression data	
RNA tissue specificity:	Low tissue specificity	
RNA tissue distribution:	Detected in all	
IHC Sibling similarity:	Other antibody shows dissimilar IHC staining pattern	
IHC fail comment:	ANTIBODY FAILED: Improbable histological location,Not consistent with RNA	

## Immunofluorescence



IF Overlay:	antibody (green), anti-tubulin (red) and DAPI (blue)
IF main location:	
IF additional location:	
IF approved for publication on HPA:	No
IF in THP-1:	Negative
IF in U-2 OS:	Negative

## Western blot



WB Size markers (kDa):	250, 130, 100, 70, 55, 35, 25, 15, 10
WB Lanes:	Marker (1), RT4 (2), U-251 MG (3), Plasma (4), Liver (5), Tonsil (6)
WB Target weight (kDa):	16, 16, 29, 61, 64, 67, 67, 67, 67, 68, 68, 68, 68, 72
WB Validation:	Uncertain (Single band differing more than +/-20% from predicted size in kDa and not supported by experimental and/or bioinformatic data.)