

CtBP1 Polyclonal Antibody

Catalog No: YT1148

Reactivity: Human; Mouse; Rat

Applications: WB;IHC;IF;ELISA

Target: CtBP1

Fields: >>Wnt signaling pathway;>>Notch signaling pathway;>>Pathways in

cancer;>>Chronic myeloid leukemia

Gene Name: CTBP1

Protein Name: C-terminal-binding protein 1

Q13363

O88712

Human Gene Id: 1487

Human Swiss Prot

No:

Mouse Gene Id: 13016

Mouse Swiss Prot

No:

Rat Gene Id: 29382

Rat Swiss Prot No: Q9Z2F5

Immunogen: The antiserum was produced against synthesized peptide derived from human

CtBP1. AA range:388-437

Specificity: CtBP1 Polyclonal Antibody detects endogenous levels of CtBP1 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 - 1:300. ELISA: 1:20000.. IF 1:50-200

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

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: 48kD

Cell Pathway: WNT;WNT-T CELLNotch;Pathways in cancer;Chronic myeloid leukemia;

Background: This gene encodes a protein that binds to the C-terminus of adenovirus E1A

proteins. This phosphoprotein is a transcriptional repressor and may play a role during cellular proliferation. This protein and the product of a second closely related gene, CTBP2, can dimerize. Both proteins can also interact with a polycomb group protein complex which participates in regulation of gene expression during development. Alternative splicing of transcripts from this gene

results in multiple transcript variants. [provided by RefSeq, Jul 2008],

Function: cofactor:NAD. Required for efficient interaction with E1A. Cofactor binding

induces a conformation change.,function:Involved in controlling the equilibrium between tubular and stacked structures in the Golgi complex (By similarity). Corepressor targeting diverse transcription regulators such as GLIS2. Has dehydrogenase activity.,PTM:ADP-ribosylated; when cells are exposed to brefeldin-A (BFA).,PTM:Sumoylation on Lys-428 is promoted by the E3 SUMO-protein ligase CBX4.,PTM:The level of phosphorylation appears to be regulated

during the cell cycle. Phosphorylated upon DNA damage, probably by ATM or ATR. Phosphorylation by HIPK2 on Ser-422 induces proteasomal degradation., similarity: Belongs to the D-isomer specific 2-hydroxyacid

dehydrogenase family.,subunit:Interacts with the C-terminus of adenovirus E1A protein, ELK3 and CTIP via their consensus motif P-X-[DNS]-L-[STVA]. Can form

homodime

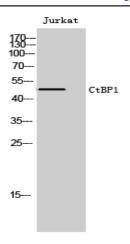
Subcellular Location:

Cytoplasm . Nucleus .

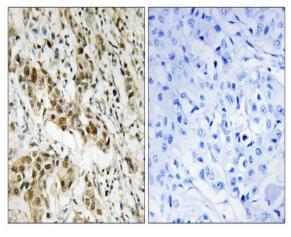
Expression:

Expressed in germinal center B-cells.

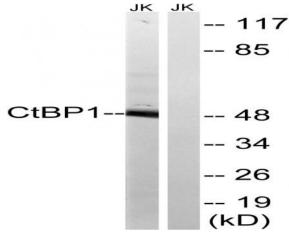
Products Images



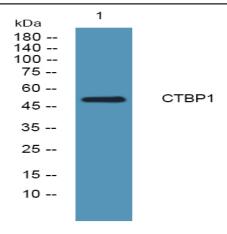
Western Blot analysis of Jurkat cells using CtBP1 Polyclonal Antibody



Immunohistochemistry analysis of paraffin-embedded human breast carcinoma tissue, using CtBP1 Antibody. The picture on the right is blocked with the synthesized peptide.



Western blot analysis of lysates from Jurkat cells, using CtBP1 Antibody. The lane on the right is blocked with the synthesized peptide.



Western blot analysis of lysates from SW480 cells, primary antibody was diluted at 1:1000, 4° over night