

## CtBP1 (phospho Ser422) Polyclonal Antibody

Catalog No: YP0815

**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

088712

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 around the phosphorylation site of Ser422. AA range:388-437

Specificity: Phospho-CtBP1 (S422) Polyclonal Antibody detects endogenous levels of

CtBP1 protein only when phosphorylated at S422.

**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: 1:10000.. 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.

Tag: orthogonal

**Sort**: 4648

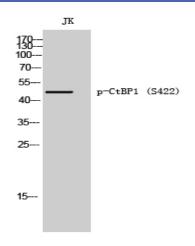
No4: 1

Host: Rabbit

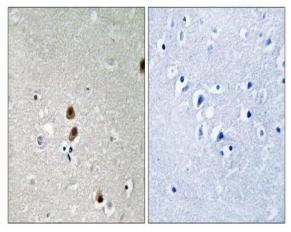
2/3

Modifications: Phospho

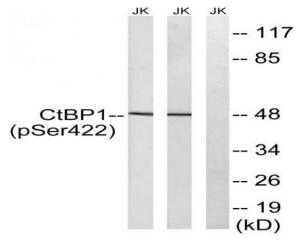
## **Products Images**



Western Blot analysis of JK cells using Phospho-CtBP1 (S422) Polyclonal Antibody



Immunohistochemistry analysis of paraffin-embedded human brain, using CtBP1 (Phospho-Ser422) Antibody. The picture on the right is blocked with the phospho peptide.



Western blot analysis of lysates from Jurkat cells treated with TNF 20ng/ml 30' and Jurkat cells treated with UV 15', using CtBP1 (Phospho-Ser422) Antibody. The lane on the right is blocked with the phospho peptide.