

## Cortactin (Acetyl Lys235) Polyclonal Antibody

Catalog No: YK0057

**Reactivity:** Human; Mouse; Rat

**Applications:** WB;ELISA

Target: Cortactin

Fields: >>Tight junction;>>Bacterial invasion of epithelial cells;>>Pathogenic

Escherichia coli infection;>>Shigellosis;>>Proteoglycans in cancer

Gene Name: CTTN

**Protein Name:** Src substrate cortactin

Q14247

Q60598

Human Gene Id: 2017

**Human Swiss Prot** 

No:

Mouse Gene Id: 13043

**Mouse Swiss Prot** 

No:

Immunogen: Synthesized acetyl-peptide derived from the Internal region of human Cortactin

around the acetylation site of K235.

**Specificity:** Acetyl-Cortactin (K235) Polyclonal Antibody detects endogenous levels of

Cortactin protein only when acetylation at K235.

**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. ELISA: 1:10000. Not yet tested in other applications.

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

chromatography using epitope-specific immunogen.

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Concentration: 1 mg/ml

**Storage Stability:** -15°C to -25°C/1 year(Do not lower than -25°C)

Observed Band: 62kD

**Cell Pathway:** Tight junction; Pathogenic Escherichia coli infection;

**Background:** cortactin(CTTN) Homo sapiens This gene is overexpressed in breast cancer

and squamous cell carcinomas of the head and neck. The encoded protein is localized in the cytoplasm and in areas of the cell-substratum contacts. This gene has two roles: (1) regulating the interactions between components of adherenstype junctions and (2) organizing the cytoskeleton and cell adhesion structures of epithelia and carcinoma cells. During apoptosis, the encoded protein is degraded in a caspase-dependent manner. The aberrant regulation of this gene contributes to tumor cell invasion and metastasis. Three splice variants that encode different

isoforms have been identified for this gene. [provided by RefSeq, May 2010],

**Function:** function: May contribute to the organization of cell structure. The SH3 motif may

function as a binding region to cytoskeleton. Tyrosine phosphorylation in

transformed cells may contribute to cellular growth regulation and transformation.,online information:Cortactin entry,similarity:Contains 1 SH3 domain.,similarity:Contains 7 cortactin repeats.,subcellular location:Associated with membrane ruffles and lamellipodia.,subunit:Interacts with SHANK2 and

SHANK3 via its SH2 domain. Also interacts with FGD1 (By similarity). Interacts

with PLXDC2.,

Subcellular Cytoplasm, cytoskeleton . Cell projection, lamellipodium . Cell projection, ruffle.

Location : Cell projection, dendrite . Cell projection . Cell membrane ; Peripheral membrane

protein; Cytoplasmic side. Cell projection, podosome. Cell junction. Cell

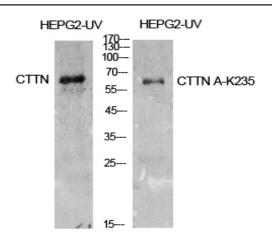
junction, focal adhesion. Membrane, clathrin-coated pit. Cell projection, dendritic spine. Cytoplasm, cell cortex. Colocalizes transiently with PTK2/FAK1 at focal adhesions (By similarity). Associated with membrane ruffles and lamellipodia. In the presence of CTTNBP2NL, colocalizes with stress fibers (By similarity). In the presence of CTTNBP2, localizes at the cell cortex (By similarity). In response to

neuronal activation by glutamate, redistributes from dendritic spines to the

dendritic shaft (By similarity). Colocalizes with DNM2 at the basis

**Expression:** Coronary artery, Epithelium, Mammary gland, Placenta, Platelet, Testis,

## **Products Images**



Western Blot analysis of HepG2-UV cells using Acetyl-Cortactin (K235) Polyclonal Antibody. Antibody was diluted at 1:500. Secondary antibody(catalog#:RS0002) was diluted at 1:20000