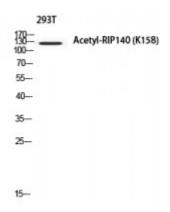


## Anti-RIP140 (Acetyl Lys158) antibody





**Description** Rabbit polyclonal to RIP140 (Acetyl Lys158).

Model STJ97698

**Host** Rabbit

**Reactivity** Human, Mouse, Rat

**Applications** ELISA, WB

Immunogen Synthesized peptide derived from human RIP140 around the acetylation site

of K158.

**Gene ID** <u>8204</u>

Gene Symbol NRIP1

**Dilution range** WB 1:500-1:2000ELISA 1:10000

**Specificity** Acetyl-RIP140 (K158) Polyclonal Antibody detects endogenous levels of

RIP140 around the acetylation site of K158 protein.

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

chromatography using epitope-specific immunogen.

**Note** For Research Use Only (RUO).

**Protein Name** Nuclear receptor-interacting protein 1 Nuclear factor RIP140 Receptor-

interacting protein 140

**Clonality** Polyclonal

**Conjugation** Unconjugated

**Isotype** IgG

**Formulation** Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

**Concentration** 1 mg/ml

**Storage Instruction** Store at -20°C, and avoid repeat freeze-thaw cycles.

Database Links <u>HGNC:8001OMIM:602490</u>

Alternative Names Nuclear receptor-interacting protein 1 Nuclear factor RIP140 Receptor-

interacting protein 140

**Function** Modulates transcriptional activation by steroid receptors such as NR3C1,

NR3C2 and ESR1. Also modulates transcriptional repression by nuclear hormone receptors. Positive regulator of the circadian clock gene expression: stimulates transcription of ARNTL/BMAL1, CLOCK and CRY1 by acting as

a coactivator for RORA and RORC.

Sequence and Domain Family Contains 9 Leu-Xaa-Xaa-Leu-Leu (LXXLL) motifs, which have different

affinities for nuclear receptors. The C-terminal LTKTNPILYYMLQK motif is required for ligand-dependent interaction with RAAR and RXRB homodimers and heterodimers, for the corepressor activity, and for the formation of an HDAC3 complex with RARA/RXRB. Contains at least four autonomous repression domains (RD1-4). RD1 functions via a histone deacetylase (HDAC)-independent mechanism, whereas RD2, RD3 and RD4 can function by HDAC-dependent or independent mechanisms, depending on cell type.

RD2 is dependent on CTBP binding.

Cellular Localization Nucleus. Localized to discrete foci and redistributes to larger nuclear domains

upon binding to ligand-bound NR3C1.

**Post-translational** Acetylation regulates its nuclear translocation and corepressive activity .

Acetylation abolishes interaction with CTBP1. Phosphorylation enhances

interaction with YWHAH.

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**Modifications** 

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