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Phospho-DAXX (Ser668) Polyclonal Antibody

Catalog No. E-AB-21159

Note: Centrifuge before opening to ensure complete recovery of vial contents.

Description

Reactivity Human

Synthesized peptide derived from human Daxx around the phosphorylation site of **Immunogen**

Ser668

Host Rabbit **Isotype** IgG

Purification Affinity purification Conjugation **Unconjugated**

Buffer PBS with 0.02% sodium azide, 0.5% protective protein and 50% glycerol, pH7.4

Applications Recommended Dilution

WB 1:500-1:2000 **IHC** 1:100-1:300 **ELISA** 1:10000

Data



Western Blot analysis of 293T cells with Phospho-Daxx (Ser668) Polyclonal Antibody

> Observed Mw:81kDa Calculated Mw:81kDa

Preparation & Storage

Storage Store at -20°C. Avoid freeze / thaw cycles.

Background

Acts as an adapter protein in a MDM2-DAXX-USP7 complex by regulating the RING-finger E3 ligase MDM2 ubiquitination activity. Under non-stress condition, in association with the deubiquitinating USP7, prevents MDM2 selfubiquitination and enhances the intrinsic E3 ligase activity of MDM2 towards TP53, thereby promoting TP53 ubiquitination and subsequent proteasomal degradation. Upon DNA damage, its association with MDM2 and USP7 is disrupted, resulting in increased MDM2 autoubiquitination and consequently, MDM2 degradation, which leads to TP53 stabilization. Proposed to mediate activation of the JNK pathway and apoptosis via MAP3K5 in response to signaling from TNFRSF6 and TGFBR2. Interaction with HSPB1/HSP27 may prevent interaction with TNFRSF6 and MAP3K5

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and block DAXX-mediated apoptosis. In contrast, in lymphoid cells JNC activation and TNFRSF6-mediated apoptosis may not involve DAXX. Seems to regulate transcription in PML/POD/ND10 nuclear bodies together with PML and may influence TNFRSF6-dependent apoptosis thereby. Down-regulates basal and activated transcription. Seems to act as a transcriptional corepressor and inhibits PAX3 and ETS1 through direct protein-protein interaction. Modulates PAX5 activity. Its transcription repressor activity is modulated by recruiting it to subnuclear compartments like the nucleolus or PML/POD/ND10 nuclear bodies through interactions with MCSR1 and PML, respectively.

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