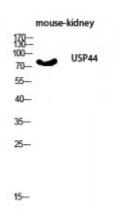


## Anti-USP44 antibody





**Description** Rabbit polyclonal to USP44.

Model STJ96206

**Host** Rabbit

**Reactivity** Human

**Applications** ELISA, WB

Immunogen Synthesized peptide derived from human USP44

Immunogen Region 180-260 aa, Internal

**Gene ID** 84101

Gene Symbol <u>USP44</u>

**Dilution range** WB 1:500-1:2000ELISA 1:40000

**Specificity** USP44 Polyclonal Antibody detects endogenous levels of USP44 protein.

**Tissue Specificity** Expressed in testis. Expressed at high levels in T-cell acute lymphoblastic

leukemia.

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

chromatography using epitope-specific immunogen.

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

**Protein Name**Ubiquitin carboxyl-terminal hydrolase 44 Deubiquitinating enzyme 44

Ubiquitin thioesterase 44 Ubiquitin-specific-processing protease 44

Molecular Weight 81 kDa

**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:20064OMIM:610993</u>

Alternative Names Ubiquitin carboxyl-terminal hydrolase 44 Deubiquitinating enzyme 44

Ubiquitin thioesterase 44 Ubiquitin-specific-processing protease 44

**Function** Deubiquitinase that plays a key regulatory role in the spindle assembly

checkpoint or mitotic checkpoint by preventing premature anaphase onset. Acts by specifically mediating deubiquitination of CDC20, a negative regulator of the anaphase promoting complex/cyclosome (APC/C).

Deubiquitination of CDC20 leads to stabilize the MAD2L1-CDC20-APC/C ternary complex (also named mitotic checkpoint complex), thereby preventing premature activation of the APC/C. Promotes association of MAD2L1 with CDC20 and reinforces the spindle assembly checkpoint. Acts as a negative

regulator of histone H2B (H2BK120ub1) ubiquitination.

Cellular Localization Nucleus. Peaks in interphase, with relatively low levels maintained throughout

mitosis.

**Post-translational** Dephosphorylated by CTDP1.; Ubiquitinated; undergoes both 'Lys-48'- and

**Modifications** 'Lys-63'-linked polyubiquitination and is degraded by the proteasome.

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