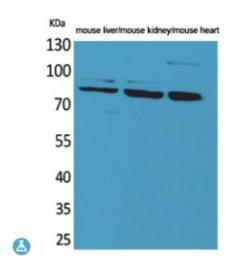
## Anti-CUL-4B antibody



**Description** Rabbit polyclonal to CUL-4B.

Model STJ96553

**Host** Rabbit

**Reactivity** Human, Mouse, Rat

**Applications** ELISA, IHC, WB

**Immunogen** Synthesized peptide derived from human CUL-4B.

Immunogen Region Internal

**Gene ID** <u>8450</u>

Gene Symbol CUL4B

**Dilution range** WB 1:500-1:2000IHC-P 1:100-300ELISA 1:20000

**Specificity** CUL-4B Polyclonal Antibody detects endogenous levels of CUL-4B 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 Cullin-4B CUL-4B

**Molecular Weight** 110 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:25550MIM:300304</u>

Alternative Names Cullin-4B CUL-4B

**Function** Core component of multiple cullin-RING-based E3 ubiquitin-protein ligase

complexes which mediate the ubiquitination and subsequent proteasomal degradation of target proteins. The functional specificity of the E3 ubiquitin-protein ligase complex depends on the variable substrate recognition subunit. CUL4B may act within the complex as a scaffold protein, contributing to catalysis through positioning of the substrate and the ubiquitin-conjugating enzyme. Plays a role as part of the E3 ubiquitin-protein ligase complex in polyubiquitination of CDT1, histone H2A, histone H3 and histone H4 in response to radiation-induced DNA damage. Targeted to UV damaged chromatin by DDB2 and may be important for DNA repair and DNA replication. Required for ubiquitination of cyclin E, and consequently, normal G1 cell cycle progression. Regulates the mammalian target-of-rapamycin (mTOR) pathway involved in control of cell growth, size and metabolism. Specific CUL4B regulation of the mTORC1-mediated pathway is dependent upon 26S proteasome function and requires interaction between CUL4B and

MLST8.

Cellular Localization Nucleus

Post-translational Modifications

Neddylated. Deneddylated via its interaction with the COP9 signalosome

(CSN) complex.

St John's Laboratory Ltd

**F** +44 (0)207 681 2580

**T** +44 (0)208 223 3081

W http://www.stjohnslabs.com/ E info@stjohnslabs.com