

# UCHL1 / PGP9.5 Antibody

Purified Mouse Monoclonal Antibody (Mab) Catalog # AP52772

### **Product Information**

Application WB, ICC
Primary Accession P09936
Reactivity Human
Host Mouse
Clonality Monoclonal
Isotype IgG2b
Calculated MW 24824

### **Additional Information**

**Gene ID** 7345

Other Names UCHL1 / PGP9.5;UCHL1;B220;CD 45;CD45;cd45 antigen;ec3.1.3.48;GP

180;GP180;Human homolog of severe combined immunodeficiency due to PTPRC deficiency;L CA;L-CA;lca;Leukocyte common antigen;LY 5;LY5;Protein tyrosine phosphatase receptor type C;Protein tyrosine phosphatase receptor type c polypeptide;PTPRC;PTPRC\_HUMAN;Receptor-type tyrosine-protein phosphatase C;SCID due to PTPRC deficiency;t200;T200 glycoprotein;T200

leukocyte common antigen.

**Dilution** WB~~1:1000 ICC~~1:300

**Format** Purified mouse monoclonal in buffer containing 0.1M Tris-Glycine (pH 7.4,

150 mM NaCl) with 0.09% (W/V) sodium azide, 50% glycerol

**Storage** Store at -20 °C.Stable for 12 months from date of receipt

### **Protein Information**

Name UCHL1

**Function** Deubiquitinase that plays a role in the regulation of several processes such

as maintenance of synaptic function, cardiac function, inflammatory response or osteoclastogenesis (PubMed:22212137, PubMed:23359680). Abrogates the ubiquitination of multiple proteins including WWTR1/TAZ, EGFR, HIF1A and

beta-site amyloid precursor protein cleaving enzyme 1/BACE1

(PubMed:22212137, PubMed:25615526). In addition, recognizes and

hydrolyzes a peptide bond at the C-terminal glycine of ubiquitin to maintain a

stable pool of monoubiquitin that is a key requirement for the ubiquitin-proteasome and the autophagy-lysosome pathways

(PubMed:<u>12408865</u>, PubMed:<u>8639624</u>, PubMed:<u>9774100</u>). Regulates amyloid precursor protein/APP processing by promoting BACE1 degradation resulting

in decreased amyloid beta production (PubMed:22212137). Plays a role in the immune response by regulating the ability of MHC I molecules to reach cross-presentation compartments competent for generating Ag-MHC I complexes (By similarity). Mediates the 'Lys-48'-linked deubiquitination of the transcriptional coactivator WWTR1/TAZ leading to its stabilization and inhibition of osteoclastogenesis (By similarity). Deubiquitinates and stabilizes epidermal growth factor receptor EGFR to prevent its degradation and to activate its downstream mediators (By similarity). Modulates oxidative activity in skeletal muscle by regulating key mitochondrial oxidative proteins (By similarity). Enhances the activity of hypoxia-inducible factor 1-alpha/HIF1A by abrogateing its VHL E3 ligase-mediated ubiquitination and consequently inhibiting its degradation (PubMed:25615526).

#### **Cellular Location**

Cytoplasm. Endoplasmic reticulum membrane; Lipid- anchor. Note=About 30% of total UCHL1 is associated with membranes in brain. Localizes near and/or within mitochondria to potentially interact with mitochondrial proteins {ECO:0000250|UniProtKB:Q9R0P9}

#### **Tissue Location**

Found in neuronal cell bodies and processes throughout the neocortex (at protein level). Expressed in neurons and cells of the diffuse neuroendocrine system and their tumors. Weakly expressed in ovary. Down-regulated in brains from Parkinson disease and Alzheimer disease patients.

## **Background**

Ubiquitin-protein hydrolase involved both in the processing of ubiquitin precursors and of ubiquitinated proteins. This enzyme is a thiol protease that recognizes and hydrolyzes a peptide bond at the C-terminal glycine of ubiquitin. Also binds to free monoubiquitin and may prevent its degradation in lysosomes. The homodimer may have ATP-independent ubiquitin ligase activity.

#### References

Hillier L.W.,et al.Nature 434:724-731(2005).

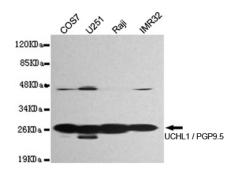
Mural R.J.,et al.Submitted (JUL-2005) to the EMBL/GenBank/DDBJ databases.

Day I.N.M.,et al.Biochem. J. 268:521-524(1990).

Choi J.,et al.J. Biol. Chem. 279:13256-13264(2004).

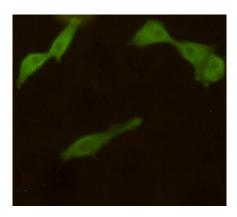
Lubec G.,et al.Submitted (DEC-2008) to UniProtKB.

# **Images**



Western blot detection of UCHL1 / PGP9.5 in U251,IMR32,Raji and COS7 cell lysates and using UCHL1 / PGP9.5 mouse mAb (1:1000 diluted).Predicted band size: 25KDa.Observed band size: 28KDa.

Immunocytochemistry stain of COS7 using UCHL1 / PGP9.5 mouse mAb (1:300).



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