Version: 3.0.0 Revision date: 12 Sep 2025



## **Ubiquitin Carboxyl-Terminal Hydrolase 14 (USP14) Antibody**

Catalogue No.:abx031550



Modification of target proteins by ubiquitin participates in a wide array of biological functions. Proteins destined for degradation or processing via the 26 S proteasome are coupled to multiple copies of ubiquitin. However, attachment of ubiquitin or ubiquitin-related molecules may also result in changes in subcellular distribution or modification of protein activity. An additional level of ubiquitin regulation, deubiquitination, is catalyzed by proteases called deubiquitinating enzymes, which fall into four distinct families. Ubiquitin C-terminal hydrolases, ubiquitin-specific processing proteases (USPs), 1 OTU-domain ubiquitin-aldehyde-binding proteins, and Jab1/Pad1/MPN-domain-containing metallo-enzymes. Among these four families, USPs represent the most widespread and represented deubiquitinating enzymes across evolution. USPs tend to release ubiquitin from a conjugated protein. They display similar catalytic domains containing conserved Cys and His boxes but divergent N-terminal and occasionally C-terminal extensions, which are thought to function in substrate recognition, subcellular localization, and protein-protein interactions.

Target: Ubiquitin Carboxyl-Terminal Hydrolase 14 (USP14)

Clonality: Polyclonal

Reactivity: Human

## **Datasheet**

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**Tested Applications:** ELISA, WB, IHC

Host: Rabbit

Recommended dilutions: WB: 1/1000, IHC-P: 1/50 - 1/100. Not tested in IHC-F. Optimal dilutions/concentrations should be

determined by the end user.

Conjugation: Unconjugated

KLH-conjugated synthetic peptide between 1-30 amino acids from the N-terminal region of human Immunogen:

USP14.

IgG Isotype:

Form: Liquid

**Purification:** Purified through a protein G column, eluted with high and low pH buffers and neutralized

immediately, followed by dialysis against PBS.

Storage: Aliquot and store at -20°C. Avoid repeated freeze/thaw cycles.

P54578 (UniProt, ExPASy) **UniProt Primary AC:** 

**NCBI Accession:** NP 001032411.1, NP 005142.1

**KEGG:** hsa:9097

String: 9606.ENSP00000261601

**Molecular Weight:** Calculated MW: 56.1 kDa

Buffer: PBS containing 0.09% sodium azide.

Predicted to react with Mouse and Cow USP14. Specificity:

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