

26S Proteasome Non-ATPase Regulatory Subunit 9 (PSMD9) Antibody

Catalogue No.: abx028124



The 26S proteasome is a multicatalytic proteinase complex with a highly ordered structure composed of 2 complexes, a 20S core and a 19S regulator. The 20S core is composed of 4 rings of 28 non-identical subunits; 2 rings are composed of 7 alpha subunits and 2 rings are composed of 7 beta subunits. The 19S regulator is composed of a base, which contains 6 ATPase subunits and 2 non-ATPase subunits, and a lid, which contains up to 10 non-ATPase subunits. Proteasomes are distributed throughout eukaryotic cells at a high concentration and cleave peptides in an ATP/ubiquitin-dependent process in a non-lysosomal pathway. An essential function of a modified proteasome, the immunoproteasome, is the processing of class I MHC peptides. This gene encodes a non-ATPase subunit of the 19S regulator.

Target: 26S Proteasome Non-ATPase Regulatory Subunit 9 (PSMD9)

Clonality: Polyclonal

Reactivity: Human

Tested Applications: ELISA, WB

Host: Rabbit

Recommended dilutions: WB: 1/1000. Optimal dilutions/concentrations should be determined by the end user.

Conjugation: Unconjugated

Immunogen: KLH-conjugated synthetic peptide between 176-205 amino acids from the C-terminal region of human PSMD9.

Isotype: IgG

Form: Liquid

Purification: Purified through a protein A column, followed by peptide affinity purification.

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

Datasheet

Version: 3.0.0

Revision date: 10 Sep 2025



UniProt Primary AC: O00233 ([UniProt](#), [ExPASy](#))

Gene Symbol: PSMD9

GeneID: [5715](#)

OMIM: [603146](#)

NCBI Accession: NP_001248329.1, NM_001261400.2, NP_002804.2, NM_002813.6

KEGG: hsa:5715

Ensembl: ENSG00000110801

String: [9606.ENSP00000440485](#)

Molecular Weight: Calculated MW: 24.7 kDa

Buffer: PBS containing 0.09% sodium azide.

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