

**Goat Anti-PRMT5 Antibody**  
Peptide-affinity purified goat antibody  
Catalog # AF1866a

**Specification**

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**Goat Anti-PRMT5 Antibody - Product Information**

Application	WB
Primary Accession	<a href="#">O14744</a>
Other Accession	<a href="#">NP_001034708</a> , <a href="#">10419</a> , <a href="#">27374 (mouse)</a> , <a href="#">364382 (rat)</a>
Reactivity	Human
Predicted	Mouse, Rat, Pig, Dog
Host	Goat
Clonality	Polyclonal
Concentration	100ug/200ul
Isotype	IgG
Calculated MW	72684

**Goat Anti-PRMT5 Antibody - Additional Information**

**Gene ID** 10419

**Other Names**

Protein arginine N-methyltransferase 5, 2.1.1.-, 72 kDa ICh-binding protein, Histone-arginine N-methyltransferase PRMT5, 2.1.1.125, Jak-binding protein 1, Shk1 kinase-binding protein 1 homolog, SKB1 homolog, SKB1Hs, Protein arginine N-methyltransferase 5, N-terminally processed, PRMT5, HRMT1L5, IBP72, JBP1, SKB1

**Format**

0.5 mg IgG/ml in Tris saline (20mM Tris pH7.3, 150mM NaCl), 0.02% sodium azide, with 0.5% bovine serum albumin

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

**Precautions**

Goat Anti-PRMT5 Antibody is for research use only and not for use in diagnostic or therapeutic procedures.

**Goat Anti-PRMT5 Antibody - Protein Information**

**Name** PRMT5

**Synonyms** HRMT1L5, IBP72, JBP1, SKB1

**Function**

Arginine methyltransferase that can both catalyze the formation of omega-N monomethylarginine (MMA) and symmetrical dimethylarginine (sDMA), with a preference for the formation of MMA (PubMed:<a href="http://www.uniprot.org/citations/10531356" target="\_blank">10531356</a>, PubMed:<a href="http://www.uniprot.org/citations/11152681" target="\_blank">11152681</a>, PubMed:<a href="http://www.uniprot.org/citations/11747828" target="\_blank">11747828</a>, PubMed:<a href="http://www.uniprot.org/citations/12411503" target="\_blank">12411503</a>, PubMed:<a href="http://www.uniprot.org/citations/15737618" target="\_blank">15737618</a>, PubMed:<a href="http://www.uniprot.org/citations/17709427" target="\_blank">17709427</a>, PubMed:<a href="http://www.uniprot.org/citations/20159986" target="\_blank">20159986</a>, PubMed:<a href="http://www.uniprot.org/citations/20810653" target="\_blank">20810653</a>, PubMed:<a href="http://www.uniprot.org/citations/21081503" target="\_blank">21081503</a>, PubMed:<a href="http://www.uniprot.org/citations/21258366" target="\_blank">21258366</a>, PubMed:<a href="http://www.uniprot.org/citations/21917714" target="\_blank">21917714</a>, PubMed:<a href="http://www.uniprot.org/citations/22269951" target="\_blank">22269951</a>). Specifically mediates the symmetrical dimethylation of arginine residues in the small nuclear ribonucleoproteins Sm D1 (SNRPD1) and Sm D3 (SNRPD3); such methylation being required for the assembly and biogenesis of snRNP core particles (PubMed:<a href="http://www.uniprot.org/citations/11747828" target="\_blank">11747828</a>, PubMed:<a href="http://www.uniprot.org/citations/12411503" target="\_blank">12411503</a>, PubMed:<a href="http://www.uniprot.org/citations/17709427" target="\_blank">17709427</a>). Methylates SUPT5H and may regulate its transcriptional elongation properties (PubMed:<a href="http://www.uniprot.org/citations/12718890" target="\_blank">12718890</a>). May methylate the N-terminal region of MBD2 (PubMed:<a href="http://www.uniprot.org/citations/16428440" target="\_blank">16428440</a>). Mono- and dimethylates arginine residues of myelin basic protein (MBP) in vitro. May play a role in cytokine-activated transduction pathways. Negatively regulates cyclin E1 promoter activity and cellular proliferation. Methylates histone H2A and H4 'Arg-3' during germ cell development (By similarity). Methylates histone H3 'Arg-8', which may repress transcription (By similarity). Methylates the Piwi proteins (PIWIL1, PIWIL2 and PIWIL4), methylation of Piwi proteins being required for the interaction with Tudor domain-containing proteins and subsequent localization to the meiotic nuage (By similarity). Methylates RPS10. Attenuates EGF signaling through the MAPK1/MAPK3 pathway acting at 2 levels. First, monomethylates EGFR; this enhances EGFR 'Tyr-1197' phosphorylation and PTPN6 recruitment, eventually leading to reduced SOS1 phosphorylation (PubMed:<a href="http://www.uniprot.org/citations/21258366" target="\_blank">21258366</a>, PubMed:<a href="http://www.uniprot.org/citations/21917714" target="\_blank">21917714</a>). Second, methylates RAF1 and probably BRAF, hence destabilizing these 2 signaling proteins and reducing their catalytic activity (PubMed:<a href="http://www.uniprot.org/citations/21917714" target="\_blank">21917714</a>). Required for induction of E-selectin and VCAM-1, on the endothelial cells surface at sites of inflammation. Methylates HOXA9 (PubMed:<a href="http://www.uniprot.org/citations/22269951" target="\_blank">22269951</a>). Methylates and regulates SRGAP2 which is involved in cell migration and differentiation (PubMed:<a href="http://www.uniprot.org/citations/20810653" target="\_blank">20810653</a>). Acts as a transcriptional corepressor in CRY1-mediated repression of the core circadian component PER1 by regulating the H4R3 dimethylation at the PER1 promoter (By similarity). Methylates GM130/GOLGA2, regulating Golgi ribbon formation (PubMed:<a href="http://www.uniprot.org/citations/20421892" target="\_blank">20421892</a>). Methylates H4R3 in genes involved in glioblastomagenesis in a CHTOP- and/or TET1-dependent manner (PubMed:<a href="http://www.uniprot.org/citations/25284789" target="\_blank">25284789</a>). Symmetrically methylates POLR2A, a modification that allows the recruitment to POLR2A of proteins including SMN1/SMN2 and SETX. This is required for resolving RNA-DNA hybrids created by RNA polymerase II, that form R-loop in transcription terminal regions, an important step in proper transcription termination (PubMed:<a href="http://www.uniprot.org/citations/26700805" target="\_blank">26700805</a>). Along with LYAR, binds the promoter of gamma-globin HBG1/HBG2 and represses its expression (PubMed:<a href="http://www.uniprot.org/citations/25092918" target="\_blank">25092918</a>). Symmetrically methylates NCL (PubMed:<a href="http://www.uniprot.org/citations/21081503" target="\_blank">21081503</a>). Methylates p53/TP53; methylation might possibly affect

p53/TP53 target gene specificity (PubMed:<a href="http://www.uniprot.org/citations/19011621" target="\_blank">19011621</a>). Involved in spliceosome maturation and mRNA splicing in prophase I spermatocytes through the catalysis of the symmetrical arginine dimethylation of SNRPB (small nuclear ribonucleoprotein- associated protein) and the interaction with tudor domain-containing protein TDRD6 (By similarity).

#### Cellular Location

Cytoplasm. Nucleus. Chromosome. Golgi apparatus. Note=Localizes to promoter regions of target genes on chromosomes (PubMed:33376131). Localizes to methylated chromatin (PubMed:16428440).

#### Tissue Location

Ubiquitous..

### Goat Anti-PRMT5 Antibody - Protocols

Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

### Goat Anti-PRMT5 Antibody - Images



AF1866a staining (2 µg/ml) of Jurkat lysate (RIPA buffer, 35 µg total protein per lane). Primary incubated for 1 hour. Detected by western blot using chemiluminescence.

### Goat Anti-PRMT5 Antibody - References

The core binding factor CBF negatively regulates skeletal muscle terminal differentiation. Philipot O, et al. PLoS One, 2010 Feb 25. PMID 20195544.

The Kruppel-like zinc finger protein ZNF224 recruits the arginine methyltransferase PRMT5 on the

transcriptional repressor complex of the aldolase A gene. Cesaro E, et al. J Biol Chem, 2009 Nov 20. PMID 19741270.

PRMT5-mediated methylation of histone H4R3 recruits DNMT3A, coupling histone and DNA methylation in gene silencing. Zhao Q, et al. Nat Struct Mol Biol, 2009 Mar. PMID 19234465.

Fibroblast growth factor 2 (FGF-2) is a novel substrate for arginine methylation by PRMT5. Bruns AF, et al. Biol Chem, 2009 Jan. PMID 19086919.

Ski co-repressor complexes maintain the basal repressed state of the TGF-beta target gene, SMAD7, via HDAC3 and PRMT5. Tabata T, et al. Genes Cells, 2009 Jan. PMID 19032343.