

**PRMT5 Antibody (N-term) Blocking peptide**  
**Synthetic peptide**  
**Catalog # BP13773a****Specification**

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**PRMT5 Antibody (N-term) Blocking peptide - Product Information**Primary Accession [O14744](#)**PRMT5 Antibody (N-term) Blocking peptide - Additional Information****Gene ID** 10419**Other Names**

Protein arginine N-methyltransferase 5, 211-, 72 kDa ICln-binding protein, Histone-arginine N-methyltransferase PRMT5, 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

**Target/Specificity**

The synthetic peptide sequence used to generate the antibody AP13773a was selected from the N-term region of PRMT5. A 10 to 100 fold molar excess to antibody is recommended. Precise conditions should be optimized for a particular assay.

**Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

**PRMT5 Antibody (N-term) Blocking peptide - 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/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>, PubMed:<a href="http://www.uniprot.org/citations/21081503" target="\_blank">21081503</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/12411503" target="\_blank">12411503</a>, PubMed:<a href="http://www.uniprot.org/citations/11747828" target="\_blank">11747828</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/21917714" target="\_blank">21917714</a>, PubMed:<a href="http://www.uniprot.org/citations/21258366" target="\_blank">21258366</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..

**PRMT5 Antibody (N-term) Blocking peptide - Protocols**

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

- [Blocking Peptides](#)

**PRMT5 Antibody (N-term) Blocking peptide - Images****PRMT5 Antibody (N-term) Blocking peptide - Background**

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. 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. Methylates SUPT5H. Mono- and dimethylates arginine residues of myelin basic protein (MBP) in vitro. Plays a role in the assembly of snRNP core particles. May play a role in cytokine-activated transduction pathways. Negatively regulates cyclin E1 promoter activity and cellular proliferation. May regulate the SUPT5H transcriptional elongation properties. May be part of a pathway that is connected to a chloride current, possibly through cytoskeletal rearrangement. Methylates histone H2A and H4 'Arg-3' during germ cell development. Methylates histone H3 'Arg-8', which may repress transcription. 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. Methylates RPS10.

**PRMT5 Antibody (N-term) Blocking peptide - References**

Aggarwal, P., et al. Cancer Cell 18(4):329-340(2010) Rank, G., et al. Blood 116(9):1585-1592(2010) Cesaro, E., et al. J. Biol. Chem. 284(47):32321-32330(2009) Zhao, Q., et al. Nat. Struct. Mol. Biol. 16(3):304-311(2009) Bruns, A.F., et al. Biol. Chem. 390(1):59-65(2009)