

# PRMT5 Antibody (N-term)

Affinity Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP13773a

# **Product Information**

| Application       | IHC-P, WB, E   |
|-------------------|--|
| Primary Accession | <u>014744</u>  |
| Other Accession   | <u>Q4R5M3</u> , <u>A7YW45</u> , <u>NP_006100.2</u> , <u>NP_001034708.1</u> |
| Reactivity        | Human  |
| Predicted         | Bovine, Monkey   |
| Host              | Rabbit   |
| Clonality         | Polyclonal   |
| Isotype           | Rabbit IgG   |
| Clone Names       | RB33884  |
| Calculated MW     | 72684  |
| Antigen Region    | 76-104   |

### **Additional Information**

| Gene ID            | 10419  |
|--------------------|--|
| Other Names        | Protein arginine N-methyltransferase 5, 211-, 72 kDa ICIn-binding protein,<br>Histone-arginine N-methyltransferase PRMT5, Jak-binding protein 1, Shk1<br>kinase-binding protein 1 homolog, SKB1 homolog, SKB1Hs, Protein arginine<br>N-methyltransferase 5, N-terminally processed, PRMT5, HRMT1L5, IBP72,<br>JBP1, SKB1 |
| Target/Specificity | This PRMT5 antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 76-104 amino acids from the N-terminal region of human PRMT5.  |
| Dilution           | IHC-P~~1:100~500 WB~~1:1000 E~~Use at an assay dependent concentration.  |
| Format             | Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide.<br>This antibody is purified through a protein A column, followed by peptide<br>affinity purification.   |
| Storage            | Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.  |
| Precautions        | PRMT5 Antibody (N-term) is for research use only and not for use in diagnostic or therapeutic procedures.  |

### **Protein Information**

Name

Function

#### HRMT1L5, IBP72, JBP1, SKB1

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:10531356, PubMed:11152681, PubMed:11747828, PubMed:12411503, PubMed:15737618, PubMed:17709427, PubMed:20159986, PubMed:20810653, PubMed:21081503, PubMed:21258366, PubMed:21917714, PubMed:22269951). 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: 11747828, PubMed:<u>12411503</u>, PubMed:<u>17709427</u>). Methylates SUPT5H and may regulate its transcriptional elongation properties (PubMed:<u>12718890</u>). May methylate the N-terminal region of MBD2 (PubMed:<u>16428440</u>). 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:21258366, PubMed:21917714). Second, methylates RAF1 and probably BRAF, hence destabilizing these 2 signaling proteins and reducing their catalytic activity (PubMed:21917714). Required for induction of E-selectin and VCAM-1, on the endothelial cells surface at sites of inflammation. Methylates HOXA9 (PubMed:22269951). Methylates and regulates SRGAP2 which is involved in cell migration and differentiation (PubMed: 20810653). 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:20421892). Methylates H4R3 in genes involved in glioblastomagenesis in a CHTOP- and/or TET1-dependent manner (PubMed:25284789). 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:<u>26700805</u>). Along with LYAR, binds the promoter of gamma-globin HBG1/HBG2 and represses its expression (PubMed: 25092918). Symmetrically methylates NCL (PubMed:21081503). Methylates p53/TP53; methylation might possibly affect p53/TP53 target gene specificity (PubMed: 19011621). 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). Cytoplasm. Nucleus. Chromosome. Golgi apparatus. Note=Localizes to promoter regions of target genes on chromosomes (PubMed:33376131).

**Cellular Location** 

**Tissue Location** 

Ubiquitous..

Localizes to methylated chromatin (PubMed:16428440).

# 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.

# 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)

### Images



All lanes : Anti-PRMT5 Antibody (N-term) at 1:1000 dilution Lane 1: Jurkat whole cell lysate Lane 2: K562 whole cell lysate Lysates/proteins at 20 µg per lane. Secondary Goat Anti-Rabbit IgG, (H+L), Peroxidase conjugated at 1/10000 dilution. Predicted band size : 73 kDa Blocking/Dilution buffer: 5% NFDM/TBST.



PRMT5 Antibody (N-term) (Cat.

#AP13773a)immunohistochemistry analysis in formalin fixed and paraffin embedded human testis tissue followed by peroxidase conjugation of the secondary antibody and DAB staining.This data demonstrates the use of PRMT5 Antibody (N-term) for immunohistochemistry. Clinical relevance has not been evaluated.

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