

PRMT1 Antibody
Purified Mouse Monoclonal Antibody (Mab)
Catalog # AP52749**Specification**

PRMT1 Antibody - Product Information

Application	WB
Primary Accession	Q99873
Reactivity	Human
Host	Mouse
Clonality	Monoclonal
Isotype	IgG2b
Calculated MW	42 KDa

PRMT1 Antibody - Additional Information**Gene ID** 3276**Other Names**

ANM 1;ANM1;ANM1_HUMAN;complete cds;HCP 1;HCP1;Heterogeneous nuclear ribonucleoprotein methyltransferase 1 like 2;Heterogeneous nuclear ribonucleoproteins methyltransferase like 2;Heterogeneous nuclear ribonucleoproteins methyltransferase like2;Histone-arginine N-methyltransferase PRMT1;HMT 2;HMT1 (hnRNP methyltransferase;HMT1 (hnRNP methyltransferase S. cerevisiae) like 2;HMT1 hnRNP methyltransferase;HMT1 hnRNP methyltransferase like 2 (S. cerevisiae);HMT1 hnRNP methyltransferase like 2;HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae);HMT2;HRMT1 L2;HRMT1L 2;HRMT1L2;Human mRNA for suppressor for yeast mutant;Human mRNA for suppressor for yeast mutant complete cds;Interferon receptor 1 bound protein 4;Interferon receptor 1 bound protein4;Interferon receptor 1-bound protein 4;Interferon receptor 1bound protein 4;IR1 B4;IR1B 4;IR1B4;Mrmt 1;Mrmt1;PRMT 1;PRMT1;Protein arginine methyltransferase 1;Protein arginine N methyltransferase 1;Protein arginine N methyltransferase1;Protein arginine N-methyltransferase 1;R1B4;S. cerevisiae like 2.

Dilution

WB~~1:1000

Format

Purified mouse monoclonal antibody in PBS(pH 7.4) containing with 0.09% (W/V) sodium azide and 50% glycerol.

Storage

Store at -20 °C.Stable for 12 months from date of receipt

PRMT1 Antibody - Protein Information**Name** PRMT1 ([HGNC:5187](#))**Function**

Arginine methyltransferase that methylates (mono and asymmetric dimethylation) the guanidino

nitrogens of arginyl residues present in proteins such as ESR1, histone H2, H3 and H4, FMR1, ILF3, HNRNPA1, HNRNPD, NFATC2IP, SUPT5H, TAF15, EWS, HABP4, SERBP1, RBM15, FOXO1, CHTOP, MAP3K5/ASK1, MICU1 and NPRL2 (PubMed:10749851, PubMed:15741314, PubMed:16879614, PubMed:18951090, PubMed:22095282, PubMed:25284789, PubMed:26575292, PubMed:26876602, PubMed:27642082, PubMed:30765518, PubMed:31257072, PubMed:38006878). Constitutes the main enzyme that mediates monomethylation and asymmetric dimethylation of histone H4 'Arg-3' (H4R3me1 and H4R3me2a, respectively), a specific tag for epigenetic transcriptional activation. May be involved in the regulation of TAF15 transcriptional activity, act as an activator of estrogen receptor (ER)-mediated transactivation, play a key role in neurite outgrowth and act as a negative regulator of megakaryocytic differentiation, by modulating p38 MAPK pathway. Methylates RBM15, promoting ubiquitination and degradation of RBM15 (PubMed:26575292). Methylates MRE11 and TP53BP1, promoting the DNA damage response (PubMed:15741314, PubMed:16294045, PubMed:29651020). Methylates FOXO1 and retains it in the nucleus increasing its transcriptional activity (PubMed:18951090). Methylates CHTOP and this methylation is critical for its 5-hydroxymethylcytosine (5hmC)-binding activity (PubMed:25284789). Methylates MAP3K5/ASK1 at 'Arg-78' and 'Arg-80' which promotes association of MAP3K5 with thioredoxin and negatively regulates MAP3K5 association with TRAF2, inhibiting MAP3K5 stimulation and MAP3K5-induced activation of JNK (PubMed:22095282). Methylates H4R3 in genes involved in glioblastomagenesis in a CHTOP- and/or TET1- dependent manner (PubMed:25284789). Plays a role in regulating alternative splicing in the heart (By similarity). Methylates NPRL2 at 'Arg-78' leading to inhibition of its GTPase activator activity and then the GATOR1 complex and consequently inducing timely mTORC1 activation under methionine-sufficient conditions (PubMed:38006878).

Cellular Location

Nucleus. Nucleus, nucleoplasm {ECO:0000250|UniProtKB:Q9JIF0}. Cytoplasm. Cytoplasm, cytosol {ECO:0000250|UniProtKB:Q9JIF0}. Lysosome membrane. Note=Mostly found in the cytoplasm Colocalizes with CHTOP within the nucleus. Low levels detected also in the chromatin fraction (By similarity). Upon methionine stimulation, localizes to the lysosome membrane in an NPRL2-dependent manner (PubMed:38006878). {ECO:0000250|UniProtKB:Q9JIF0, ECO:0000269|PubMed:38006878}

Tissue Location

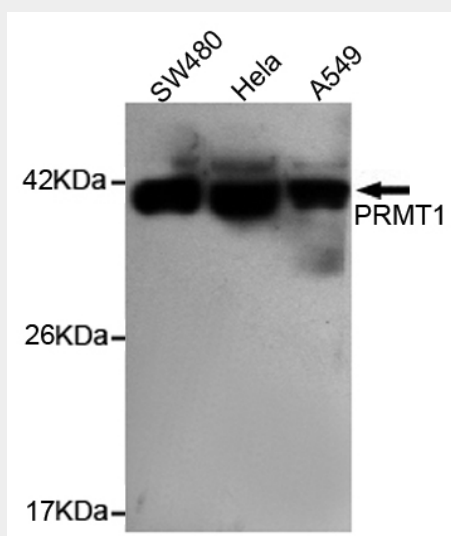
Widely expressed (PubMed:11097842). Expressed strongly in colorectal cancer cells (at protein level) (PubMed:28040436). Expressed strongly in colorectal cancer tissues compared to wild-type colon samples (at protein level) (PubMed:28040436). Expressed strongly in colorectal cancer tissues compared to wild-type colon samples (PubMed:28040436)

PRMT1 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](#)

PRMT1 Antibody - Images



Western blot detection of PRMT1 in HeLa, A549 and SW480 cell lysates using PRMT1 mouse mAb (1:1000 diluted). Predicted band size: 42KDa. Observed band size: 42KDa.

PRMT1 Antibody - Background

Arginine methyltransferase that methylates (mono and asymmetric dimethylation) the guanidino nitrogens of arginyl residues present in proteins such as ESR1, histone H2, H3 and H4, PIAS1, HNRNPA1, HNRNPD, NFATC2IP, SUPT5H, TAF15 and EWS. Constitutes the main enzyme that mediates monomethylation and asymmetric dimethylation of histone H4 'Arg-4' (H4R3me1 and H4R3me2a, respectively), a specific tag for epigenetic transcriptional activation. Together with dimethylated PIAS1, represses STAT1 transcriptional activity, in the late phase of interferon gamma (IFN-gamma) signaling. May be involved in the regulation of TAF15 transcriptional activity, act as an activator of estrogen receptor (ER)-mediated transactivation, play a key role in neurite outgrowth and act as a negative regulator of megakaryocytic differentiation, by modulating p38 MAPK pathway. Methylates FOXO1 and retains it in the nucleus increasing its transcriptional activity.

PRMT1 Antibody - References

- Scott H.S., et al. Genomics 48:330-340(1998).
Nikawa J., et al. Gene 171:107-111(1996).
Scorilas A., et al. Biochem. Biophys. Res. Commun. 278:349-359(2000).
Ota T., et al. Nat. Genet. 36:40-45(2004).
Ebert L., et al. Submitted (MAY-2004) to the EMBL/GenBank/DDBJ databases.