

Anti-PRMT4 Antibody
Catalog # ABO11573**Specification**

Anti-PRMT4 Antibody - Product Information

Application	WB
Primary Accession	Q86X55
Host	Rabbit
Reactivity	Human, Mouse, Rat
Clonality	Polyclonal
Format	Lyophilized

Description

Rabbit IgG polyclonal antibody for Histone-arginine methyltransferase CARM1(CARM1) detection. Tested with WB in Human;Mouse;Rat.

Reconstitution

Add 0.2ml of distilled water will yield a concentration of 500ug/ml.

Anti-PRMT4 Antibody - Additional Information

Gene ID 10498

Other Names

Histone-arginine methyltransferase CARM1, 2.1.1.319, Coactivator-associated arginine methyltransferase 1, Protein arginine N-methyltransferase 4, CARM1, PRMT4

Calculated MW

65854 MW KDa

Application Details

Western blot, 0.1-0.5 µg/ml, Human, Rat, Mouse

Subcellular Localization

Nucleus. Cytoplasm. Mainly nuclear during the G1, S and G2 phases of the cell cycle. Cytoplasmic during mitosis, after breakup of the nuclear membrane.

Tissue Specificity

Overexpressed in prostate adenocarcinomas and high-grade prostatic intraepithelial neoplasia. .

Protein Name

Histone-arginine methyltransferase CARM1

Contents

Each vial contains 5mg BSA, 0.9mg NaCl, 0.2mg Na2HPO4, 0.05mg Thimerosal, 0.05mg NaN3.

Immunogen

A synthetic peptide corresponding to a sequence in the middle region of human PRMT4(267-284aa HLAFTDEQLYMEQFTKA), identical to the related mouse and rat sequences.

Purification

Immunogen affinity purified.

Cross Reactivity

No cross reactivity with other proteins

Storage

At -20°C for one year. After reconstitution, at 4°C for one month. It can also be aliquotted and stored frozen at -20°C for a longer time. Avoid repeated freezing and thawing.

Sequence Similarities

Belongs to the class I-like SAM-binding methyltransferase superfamily. Protein arginine N-methyltransferase family.

Anti-PRMT4 Antibody - Protein Information

Name CARM1

Synonyms PRMT4

Function

Methylates (mono- and asymmetric dimethylation) the guanidino nitrogens of arginyl residues in several proteins involved in DNA packaging, transcription regulation, pre-mRNA splicing, and mRNA stability (PubMed: [12237300](http://www.uniprot.org/citations/12237300), PubMed: [16497732](http://www.uniprot.org/citations/16497732), PubMed: [19405910](http://www.uniprot.org/citations/19405910)). Recruited to promoters upon gene activation together with histone acetyltransferases from EP300/P300 and p160 families, methylates histone H3 at 'Arg-17' (H3R17me), forming mainly asymmetric dimethylarginine (H3R17me2a), leading to activation of transcription via chromatin remodeling (PubMed: [12237300](http://www.uniprot.org/citations/12237300), PubMed: [16497732](http://www.uniprot.org/citations/16497732), PubMed: [19405910](http://www.uniprot.org/citations/19405910)). During nuclear hormone receptor activation and TCF7L2/TCF4 activation, acts synergically with EP300/P300 and either one of the p160 histone acetyltransferases NCOA1/SRC1, NCOA2/GRIP1 and NCOA3/ACTR or CTNNB1/beta-catenin to activate transcription (By similarity). During myogenic transcriptional activation, acts together with NCOA3/ACTR as a coactivator for MEF2C (By similarity). During monocyte inflammatory stimulation, acts together with EP300/P300 as a coactivator for NF-kappa-B (By similarity). Acts as a coactivator for PPARG, promotes adipocyte differentiation and the accumulation of brown fat tissue (By similarity). Plays a role in the regulation of pre-mRNA alternative splicing by methylation of splicing factors (By similarity). Also seems to be involved in p53/TP53 transcriptional activation (By similarity). Methylates EP300/P300, both at 'Arg-2142', which may loosen its interaction with NCOA2/GRIP1, and at 'Arg-580' and 'Arg-604' in the KIX domain, which impairs its interaction with CREB and inhibits CREB-dependent transcriptional activation (PubMed: [15731352](http://www.uniprot.org/citations/15731352)). Also methylates arginine residues in RNA-binding proteins PABPC1, ELAVL1 and ELAV4, which may affect their mRNA-stabilizing properties and the half-life of their target mRNAs (By similarity). Acts as a transcriptional coactivator of ACACA/acetyl-CoA carboxylase by enriching H3R17 methylation at its promoter, thereby positively regulating fatty acid synthesis (By similarity). Independently of its methyltransferase activity, involved in replication fork progression: promotes PARP1 recruitment to replication forks, leading to poly-ADP-ribosylation of chromatin at replication forks and reduced fork speed (PubMed: [33412112](http://www.uniprot.org/citations/33412112)).

Cellular Location

Nucleus. Cytoplasm. Chromosome. Note=Mainly nuclear during the G1, S and G2 phases of the cell cycle (PubMed:19843527). Cytoplasmic during mitosis, after breakup of the nuclear membrane (PubMed:19843527) Localizes to replication forks (PubMed:33412112)

Tissue Location

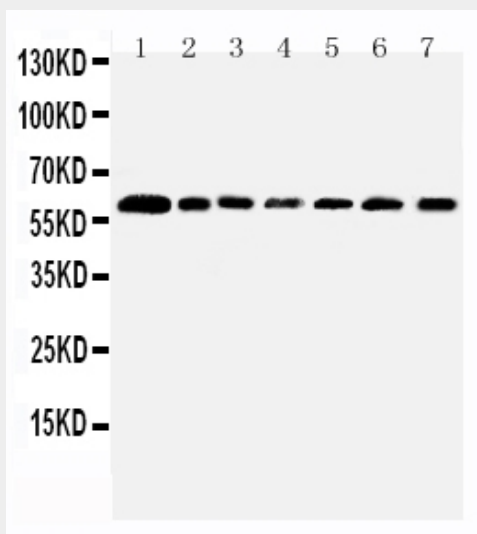
Overexpressed in prostate adenocarcinomas and high- grade prostatic intraepithelial neoplasia

Anti-PRMT4 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](#)

Anti-PRMT4 Antibody - Images



Anti-PRMT4 antibody, ABO11573, All Western blottingAll lanes: Anti-CARM1(ABO11573) at 0.5ug/mlLane 1: Rat Spleen Tissue Lysate at 40ugLane 2: Human Placenta Tissue Lysate at 40ugLane 3: Rat Kidney Tissue Lysate at 40ugLane 4: NIH Whole Cell Lysate at 40ugLane 5: HELA Whole Cell Lysate at 40ugLane 6: HL60 Whole Cell Lysate at 40ugLane 7: JURKAT Whole Cell Lysate at 40ugPredicted bind size: 60KDObserved bind size: 60KD

Anti-PRMT4 Antibody - Background

CARM1(coactivator-associated arginine methyltransferase 1), also known as PRMT4, is an enzyme encoded by the CARM1 gene found in human beings, as well as many other mammals. This gene is mapped to 19p13.2. CARM1 is a regulator of cyclin E1 and DHFR mRNA expression. Its main function includes catalyzing the transfer of a methyl group from S-adenosyl-L-methionine to the side chain nitrogens of arginine residues within proteins to form methylated arginine derivatives and S-adenosyl-L-homocysteine. CARM1 is a secondary coactivator through its association with

p160 family(SRC-1, GRIP1, AIB) of coactivators. It is responsible for moving cells toward the inner cell mass in developing blastocysts. This gene also plays an important role in androgen receptors and may play a role in prostate cancer progression.