

# Anti-CARM1 Antibody

Rabbit polyclonal antibody to CARM1 Catalog # AP61368

## **Product Information**

Application WB
Primary Accession Q86X55
Other Accession Q9WVG6

Reactivity Human, Mouse, Rat

HostRabbitClonalityPolyclonalCalculated MW65854

# **Additional Information**

**Gene ID** 10498

Other Names PRMT4; Histone-arginine methyltransferase CARM1; Coactivator-associated

arginine methyltransferase 1; Protein arginine N-methyltransferase 4

**Target/Specificity** Recognizes endogenous levels of CARM1 protein.

**Dilution** WB~~WB (1/500 - 1/1000)

**Format** Liquid in 0.42% Potassium phosphate, 0.87% Sodium chloride, pH 7.3, 30%

glycerol, and 0.09% (W/V) sodium azide.

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

## **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, PubMed:16497732, PubMed: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:<u>12237300</u>, PubMed:<u>16497732</u>, PubMed:<u>19405910</u>). 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). 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:<u>33412112</u>).

#### **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)

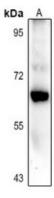
### **Tissue Location**

Overexpressed in prostate adenocarcinomas and high- grade prostatic intraepithelial neoplasia

# **Background**

KLH-conjugated synthetic peptide encompassing a sequence within the center region of human CARM1. The exact sequence is proprietary.

# **Images**



Western blot analysis of CARM1 expression in mouse kidney (A) whole cell lysates.

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