

#### **KAT2A Antibody (C-term)**

Affinity Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP13089B

### **Specification**

### KAT2A Antibody (C-term) - Product Information

Application WB,E
Primary Accession Q92830

Other Accession Q9IHD2, NP 066564.2, A0A0R4IXF6

Reactivity Human

Predicted Zebrafish, Mouse

Host Rabbit
Clonality Polyclonal
Isotype Rabbit IgG
Calculated MW 93926
Antigen Region 611-638

# KAT2A Antibody (C-term) - Additional Information

#### **Gene ID 2648**

#### **Other Names**

Histone acetyltransferase KAT2A, General control of amino acid synthesis protein 5-like 2, Histone acetyltransferase GCN5, HsGCN5, Lysine acetyltransferase 2A, STAF97, KAT2A, GCN5, GCN5L2, HGCN5

### Target/Specificity

This KAT2A antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 611-638 amino acids from the C-terminal region of human KAT2A.

### **Dilution**

WB~~1:1000

E~~Use at an assay dependent concentration.

#### **Format**

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. This antibody is purified through a protein A column, followed by peptide 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**

KAT2A Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

#### KAT2A Antibody (C-term) - Protein Information



# Name KAT2A {ECO:0000303|PubMed:27796307, ECO:0000312|HGNC:HGNC:4201}

Function Protein lysine acyltransferase that can act as a acetyltransferase, glutaryltransferase, succinvitransferase or malonvitransferase, depending on the context (PubMed:29211711, PubMed: 35995428). Acts as a histone lysine succinyltransferase: catalyzes succinylation of histone H3 on 'Lys-79' (H3K79succ), with a maximum frequency around the transcription start sites of genes (PubMed: 29211711). Succinylation of histones gives a specific tag for epigenetic transcription activation (PubMed: 29211711). Association with the 2-oxoglutarate dehydrogenase complex, which provides succinyl-CoA, is required for histone succinylation (PubMed: 29211711). In different complexes, functions either as an acetyltransferase (HAT) or as a succinyltransferase: in the SAGA and ATAC complexes, acts as a histone acetyltransferase (PubMed: 17301242, PubMed: 19103755, PubMed: 29211711). Has significant histone acetyltransferase activity with core histones, but not with nucleosome core particles (PubMed:17301242, PubMed:19103755, PubMed: 21131905). Has a a strong preference for acetylation of H3 at 'Lys-9' (H3K9ac) (PubMed:21131905). Acetylation of histones gives a specific tag for epigenetic transcription activation (PubMed: 17301242, PubMed: 19103755, PubMed: 29211711). Recruited by the XPC complex at promoters, where it specifically mediates acetylation of histone variant H2A.Z.1/H2A.Z, thereby promoting expression of target genes (PubMed: 29973595, PubMed: 31527837). Involved in long-term memory consolidation and synaptic plasticity: acts by promoting expression of a hippocampal gene expression network linked to neuroactive receptor signaling (By similarity). Acts as a positive regulator of T-cell activation: upon TCR stimulation, recruited to the IL2 promoter following interaction with NFATC2 and catalyzes acetylation of histone H3 at 'Lys-9' (H3K9ac), leading to promote IL2 expression (By similarity). Required for growth and differentiation of craniofacial cartilage and bone by regulating acetylation of histone H3 at 'Lys-9' (H3K9ac) (By similarity). Regulates embryonic stem cell (ESC) pluripotency and differentiation (By similarity). Also acetylates non- histone proteins, such as CEBPB, MRE11, PPARGC1A, PLK4 and TBX5 (PubMed: 16753578, PubMed: 17301242, PubMed: 27796307, PubMed: 29174768, PubMed: 38128537). Involved in heart and limb development by mediating acetylation of TBX5, acetylation regulating nucleocytoplasmic shuttling of TBX5 (PubMed: 29174768). Acts as a negative regulator of centrosome amplification by mediating acetylation of PLK4 (PubMed: <u>27796307</u>). Acts as a negative regulator of gluconeogenesis by mediating acetylation and subsequent inactivation of PPARGC1A (PubMed: 16753578, PubMed: 23142079). Also acts as a histone glutaryltransferase: catalyzes glutarylation of histone H4 on 'Lys-91' (H4K91glu), a mark that destabilizes nucleosomes by promoting dissociation of the H2A-H2B dimers from nucleosomes (PubMed: 31542297).

#### **Cellular Location**

Nucleus. Chromosome Cytoplasm, cytoskeleton, microtubule organizing center, centrosome. Note=Mainly localizes to the nucleus (PubMed:27796307). Localizes to sites of DNA damage (PubMed:25593309) Also localizes to centrosomes in late G1 and around the G1/S transition, coinciding with the onset of centriole formation (PubMed:27796307).

#### **Tissue Location**

Expressed in all tissues tested.

# KAT2A Antibody (C-term) - Protocols

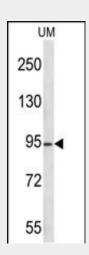
Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- <u>Immunofluorescence</u>
- Immunoprecipitation



- Flow Cytomety
- Cell Culture

# KAT2A Antibody (C-term) - Images



KAT2A Antibody (C-term) (Cat. #AP13089b) western blot analysis in uterus tumor cell line lysates (35ug/lane). This demonstrates the KAT2A antibody detected the KAT2A protein (arrow).

# KAT2A Antibody (C-term) - Background

KAT2A, or GCN5, is a histone acetyltransferase (HAT) that functions primarily as a transcriptional activator. It also functions as a repressor of NF-kappa-B (see MIM 164011) by promoting ubiquitination of the NF-kappa-B subunit RELA (MIM 164014) in a HAT-independent manner (Mao et al., 2009 [PubMed 19339690]).

### KAT2A Antibody (C-term) - References

Terreni, M., et al. Retrovirology 7, 18 (2010):
Atanassov, B.S., et al. Mol. Cell 35(3):352-364(2009)
Kelly, T.J., et al. J. Biol. Chem. 284(30):19945-19952(2009)
Mao, X., et al. Genes Dev. 23(7):849-861(2009)
Paolinelli, R., et al. Nat. Struct. Mol. Biol. 16(4):412-420(2009)