

**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide**  
**Synthetic peptide**  
**Catalog # BP1078a****Specification**

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**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - Product Information**

Primary Accession [O92830](#)  
Other Accession [O8N1A2](#)

**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - 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**

The synthetic peptide sequence used to generate the antibody [AP1078a](/product/products/AP1078a) was selected from the N-term region of human GCN5L2. A 10 to 100 fold molar excess to antibody is recommended. Precise conditions should be optimized for a particular assay.

**Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - Protein Information**

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

**Function**

Protein lysine acyltransferase that can act as a acetyltransferase, glutaryltransferase, succinyltransferase or malonyltransferase, depending on the context (PubMed:[29211711](http://www.uniprot.org/citations/29211711), PubMed:[35995428](http://www.uniprot.org/citations/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](http://www.uniprot.org/citations/29211711)). Succinylation of histones gives a specific tag for epigenetic transcription activation (PubMed:[29211711](http://www.uniprot.org/citations/29211711)). Association

with the 2-oxoglutarate dehydrogenase complex, which provides succinyl-CoA, is required for histone succinylation (PubMed:<a href="http://www.uniprot.org/citations/29211711" target="\_blank">29211711</a>). 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:<a href="http://www.uniprot.org/citations/17301242" target="\_blank">17301242</a>, PubMed:<a href="http://www.uniprot.org/citations/19103755" target="\_blank">19103755</a>, PubMed:<a href="http://www.uniprot.org/citations/29211711" target="\_blank">29211711</a>). Has significant histone acetyltransferase activity with core histones, but not with nucleosome core particles (PubMed:<a href="http://www.uniprot.org/citations/17301242" target="\_blank">17301242</a>, PubMed:<a href="http://www.uniprot.org/citations/19103755" target="\_blank">19103755</a>, PubMed:<a href="http://www.uniprot.org/citations/21131905" target="\_blank">21131905</a>). Has a strong preference for acetylation of H3 at 'Lys-9' (H3K9ac) (PubMed:<a href="http://www.uniprot.org/citations/21131905" target="\_blank">21131905</a>). Acetylation of histones gives a specific tag for epigenetic transcription activation (PubMed:<a href="http://www.uniprot.org/citations/17301242" target="\_blank">17301242</a>, PubMed:<a href="http://www.uniprot.org/citations/19103755" target="\_blank">19103755</a>, PubMed:<a href="http://www.uniprot.org/citations/29211711" target="\_blank">29211711</a>). 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:<a href="http://www.uniprot.org/citations/29973595" target="\_blank">29973595</a>, PubMed:<a href="http://www.uniprot.org/citations/31527837" target="\_blank">31527837</a>). 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, PPARGC1A, PLK4 and TBX5 (PubMed:<a href="http://www.uniprot.org/citations/17301242" target="\_blank">17301242</a>, PubMed:<a href="http://www.uniprot.org/citations/16753578" target="\_blank">16753578</a>, PubMed:<a href="http://www.uniprot.org/citations/27796307" target="\_blank">27796307</a>, PubMed:<a href="http://www.uniprot.org/citations/29174768" target="\_blank">29174768</a>). Involved in heart and limb development by mediating acetylation of TBX5, acetylation regulating nucleocytoplasmic shuttling of TBX5 (PubMed:<a href="http://www.uniprot.org/citations/29174768" target="\_blank">29174768</a>). Acts as a negative regulator of centrosome amplification by mediating acetylation of PLK4 (PubMed:<a href="http://www.uniprot.org/citations/27796307" target="\_blank">27796307</a>). Acts as a negative regulator of gluconeogenesis by mediating acetylation and subsequent inactivation of PPARGC1A (PubMed:<a href="http://www.uniprot.org/citations/16753578" target="\_blank">16753578</a>, PubMed:<a href="http://www.uniprot.org/citations/23142079" target="\_blank">23142079</a>). 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:<a href="http://www.uniprot.org/citations/31542297" target="\_blank">31542297</a>).

### 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.

**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - Protocols**

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

- [Blocking Peptides](#)

**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - Images****GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - Background**

GCN5L2 functions as a histone acetyltransferase (HAT) to promote transcriptional activation. Acetylation of histones gives a specific tag for epigenetic transcription activation. This protein has significant histone acetyltransferase activity with core histones, but not with nucleosome core particles.

**GCN5 (GCN5L2) Antibody (N-term) Blocking peptide - References**

Kikuchi, H., et al., Gene 347(1):83-97 (2005).Gangisetty, O., et al., Oncogene 23(14):2559-2563 (2004).Lusic, M., et al., EMBO J. 22(24):6550-6561 (2003).Fulco, M., et al., Mol. Cell 12(1):51-62 (2003).Liu, X., et al., J. Biol. Chem. 278(22):20405-20412 (2003).