

# KAT7 / HBO1 / MYST2 Antibody

Rabbit mAb Catalog # AP91522

# **Product Information**

**Application** WB, IHC, IF, ICC, IP, IHF

Primary Accession 095251

**Reactivity** Rat, Human, Mouse

**Clonality** Monoclonal

Other Names HBO1; HBOa; KAT7; MOZ; MYST 2; MYST protein 2; MYST2; SAS 2; SAS2 and

TIP60 protein 2; TIP60 protein 2; YBF2/SAS3; ZC2HC7;

IsotypeRabbit IgGHostRabbitCalculated MW70642

## **Additional Information**

**Dilution** WB 1:500~1:2000 IHC 1:50~1:200 ICC/IF 1:100~1:500 IP 1:50

**Purification** Affinity-chromatography

ImmunogenA synthesized peptide derived from human KAT7 / HBO1 / MYST2DescriptionComponent of the HBO1 complex which has a histone H4-specific

acetyltransferase activity, a reduced activity toward histone H3 and is responsible for the bulk of histone H4 acetylation in vivo. Through chromatin acetylation it may regulate DNA replication and act as a coactivator of

TP53-dependent transcription. Specifically represses AR-mediated

transcription.

Storage Condition and Buffer Rabbit IgG in phosphate buffered saline, pH 7.4, 150mM NaCl, 0.02% sodium

azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term.

Avoid freeze / thaw cycle.

# **Protein Information**

Name KAT7 {ECO:0000303 | PubMed:31767635, ECO:0000312 | HGNC:HGNC:17016}

**Function** Catalytic subunit of histone acetyltransferase HBO1 complexes, which

specifically mediate acetylation of histone H3 at 'Lys-14' (H3K14ac), thereby

regulating various processes, such as gene transcription, protein

ubiquitination, immune regulation, stem cell pluripotent and self-renewal

maintenance and embryonic development (PubMed: 16387653, PubMed: 21753189, PubMed: 24065767, PubMed: 26630551

PubMed:<u>21753189</u>, PubMed:<u>24065767</u>, PubMed:<u>26620551</u>, PubMed:<u>31767635</u>, PubMed:<u>31827282</u>). Some complexes also catalyze

acetylation of histone H4 at 'Lys-5', 'Lys-8' and 'Lys-12' (H4K5ac, H4K8ac and H4K12ac, respectively), regulating DNA replication initiation, regulating DNA

replication initiation (PubMed: 10438470, PubMed: 19187766,

PubMed: <u>20129055</u>, PubMed: <u>24065767</u>). Specificity of the HBO1 complexes is determined by the scaffold subunit: complexes containing BRPF scaffold

(BRPF1, BRD1/BRPF2 or BRPF3) direct KAT7/HBO1 specificity towards H3K14ac, while complexes containing JADE (JADE1, JADE2 and JADE3) scaffold direct KAT7/HBO1 specificity towards histone H4 (PubMed: 19187766, PubMed: 20129055, PubMed: 24065767, PubMed: 26620551). H3K14ac promotes transcriptional elongation by facilitating the processivity of RNA polymerase II (PubMed:31827282). Acts as a key regulator of hematopoiesis by forming a complex with BRD1/BRPF2, directing KAT7/HBO1 specificity towards H3K14ac and promoting erythroid differentiation (PubMed:21753189). H3K14ac is also required for T-cell development (By similarity). KAT7/HBO1-mediated acetylation facilitates two consecutive steps, licensing and activation, in DNA replication initiation: H3K14ac facilitates the activation of replication origins, and histone H4 acetylation (H4K5ac, H4K8ac and H4K12ac) facilitates chromatin loading of MCM complexes, promoting DNA replication licensing (PubMed: 10438470, PubMed: 11278932, PubMed: 18832067, PubMed: 19187766, PubMed: 20129055, PubMed:21856198, PubMed:24065767, PubMed:26620551). Acts as a positive regulator of centromeric CENPA assembly: recruited to centromeres and mediates histone acetylation, thereby preventing centromere inactivation mediated by SUV39H1, possibly by increasing histone turnover/exchange (PubMed:<u>27270040</u>). Involved in nucleotide excision repair: phosphorylation by ATR in response to ultraviolet irradiation promotes its localization to DNA damage sites, where it mediates histone acetylation to facilitate recruitment of XPC at the damaged DNA sites (PubMed:28719581). Acts as an inhibitor of NF-kappa-B independently of its histone acetyltransferase activity (PubMed: 16997280).

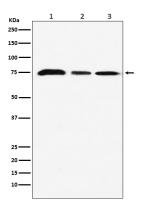
#### **Cellular Location**

Nucleus. Chromosome. Chromosome, centromere. Cytoplasm, cytosol {ECO:0000250|UniProtKB:Q5SVQ0}. Note=Associates with replication origins specifically during the G1 phase of the cell cycle (PubMed:18832067, PubMed:20129055). Localizes to transcription start sites (PubMed:21753189, PubMed:24065767). Localizes to ultraviolet- induced DNA damage sites following phosphorylation by ATR (PubMed:28719581). Localizes to centromeres in G1 phase (PubMed:27270040).

## **Tissue Location**

Ubiquitously expressed, with highest levels in testis.

# **Images**



Western blot analysis of KAT7 / HBO1 / MYST2 expression in (1)MCF7 cell lysate; (2) NIH/3T3 cell lysate; (3) C6 cell lysate.

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