

## JMJD2A Mouse mAb

CatalogNo: YM0389

## **Key Features**

**Host Species** 

Reactivity

Applications

Mouse

Human

WB,IHC,IF,ELISA

• 121kD (Calculated)

### **Recommended Dilution Ratios**

WB 1:500-1:2000 IHC 1:200-1:1000 IF 1:200-1:1000 **ELISA 1:10000** 

Not yet tested in other applications.

#### Storage

Storage\* -15°C to -25°C/1 year(Do not lower than -25°C)

Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide. **Formulation** 

## **Basic Information**

Clonality Monoclonal

## Immunogen Information

Purified recombinant fragment of human JMJD2A expressed in E. Coli. **Immunogen** 

**Specificity** JMJD2A Monoclonal Antibody detects endogenous levels of JMJD2A protein.

# | Target Information

Gene name KDM4A

Protein Name

Lysine-specific demethylase 4A

Organism	Gene ID	UniProt ID
Human	<u>9682;</u>	<u>075164;</u>
Mouse		Q8BW72;

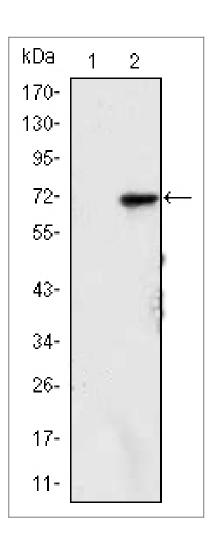
Cellular Localization Nucleus.

**Tissue specificity** Ubiquitous.

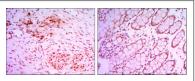
#### **Function**

cofactor:Binds 1 Fe(2+) ion per subunit., Domain:The 2 Tudor domains recognize and bind methylated histone H3 'Lvs-4' residue, Double Tudor domain has an interdigitated structure and the unusual fold is required for its ability to bind methylated histone tails. Trimethylated H3 'Lys-4' is bound in a cage of 3 aromatic residues, 2 of which are from the Tudor domain 2, while the binding specificity is determined by side-chain interactions involving residues from the Tudor domain 1. The Tudor domains are able to bind trimethylated histone H3 'Lys-4', trimethylated histone H3 'Lys-9', di- and trimethylated H4 'Lvs-20'..Function:Histone demethylase that specifically demethylates 'Lvs-9' and 'Lvs-36' residues of histone H3, thereby playing a central role in histone code. Does not demethylate histone H3 'Lys-4', H3 'Lys-27' nor H4 'Lys-20'. Demethylates trimethylated H3 'Lys-9' and H3 'Lys-36' residue, while it has no activity on mono- and dimethylated residues. Demethylation of Lys residue generates formaldehyde and succinate. Participates in transcriptional repression of ASCL2 and E2F-responsive promoters via the recruitment of histone deacetylases and NCOR1, respectively., similarity: Belongs to the JHDM3 histone demethylase family., similarity: Contains 1 ImiC domain., similarity: Contains 1 ImiN domain., similarity: Contains 2 PHD-type zinc fingers., similarity: Contains 2 Tudor domains., subunit: Interacts with histone deacetylase proteins HDAC1, HDAC2 and HDAC3. Interacts with RB and NCOR1. Interacts with HTLV-1 Tax protein., tissue specificity: Ubiquitous.,

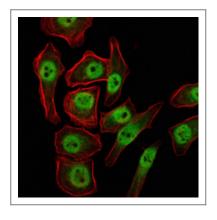
### **Validation Data**



Western Blot analysis using JMJD2A Monoclonal Antibody against HEK293 (1) and JMJD2A-hlgGFc transfected HEK293 (2) cell lysate.



Immunohistochemistry analysis of paraffin-embedded colon cancer tissues (left) and human larynx cancer tissues (right) with DAB staining using JMJD2A Monoclonal Antibody.



Immunofluorescence analysis of NTERA-2 cells using JMJD2A Monoclonal Antibody (green). Red: Actin filaments have been labeled with Alexa Fluor-555 phalloidin.

## | Contact information

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Please scan the QR code to access additional product information:

JMJD2A Mouse mAb

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