

SIRT6 mouse mAb

YM1275 Catalog No:

Human; Mouse; Rat; Monkey Reactivity:

Applications: WB;IF;IP

Target: SIRT6

>>Nicotinate and nicotinamide metabolism;>>Metabolic Fields:

pathways;>>Thermogenesis;>>Central carbon metabolism in cancer

Gene Name: sirt6

Human Gene Id: 51548

Human Swiss Prot

No:

Mouse Swiss Prot

No:

Purified recombinant human SIRT6 protein expressed in E.coli. Immunogen:

This antibody detects endogenous levels of SIRT6 and does not cross-react with **Specificity:**

related proteins.

Q8N6T7

P59941

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

Source: Monoclonal, Mouse

Dilution: wb 1:500 IF icc 1:100

The antibody was affinity-purified from mouse ascites by affinity-**Purification:**

chromatography using epitope-specific immunogen.

Concentration: 1 mg/ml

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

Observed Band: 42kD



Background:

This gene encodes a member of the sirtuin family of NAD-dependent enzymes that are implicated in cellular stress resistance, genomic stability, aging and energy homeostasis. The encoded protein is localized to the nucleus, exhibits ADP-ribosyl transferase and histone deacetylase activities, and plays a role in DNA repair, maintenance of telomeric chromatin, inflammation, lipid and glucose metabolism. Alternative splicing results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Mar 2016],

Function:

catalytic activity:NAD(+) + protein-L-arginine = nicotinamide + N(omega)-(ADP-D-ribosyl)-protein-L-arginine.,catalytic activity:NADP(+) + protein-L-arginine = nicotinamide + N(omega)-((2'-phospho-ADP)-D-ribosyl)-protein-L-arginine.,cofactor:Binds 1 zinc ion per subunit.,PTM:ADP-ribosylated (-auto).,similarity:Belongs to the sirtuin family.,similarity:Contains 1 deacetylase sirtuin-type domain.,subcellular location:Predominantly nuclear.,

Subcellular Location:

Nucleus . Chromosome . Chromosome, telomere . Endoplasmic reticulum . Predominantly nuclear (PubMed:18337721). Associated with pericentric heterochromatin and telomeric heterochromatin regions (PubMed:18337721, PubMed:27043296). Localizes to DNA damage sites: directly recognizes and binds double-strand breaks (DSBs) sites via a tunnel-like structure that has high affinity for DSBs (PubMed:21680843, PubMed:23911928, PubMed:27568560, PubMed:31995034, PubMed:32538779). A fraction localizes to the endoplasmic reticulum (PubMed:23552949).

Expression:	Blood, Eye, Lung, Placenta, Spleen, Teratocarcinoma,

Tag: ip

Sort: 1

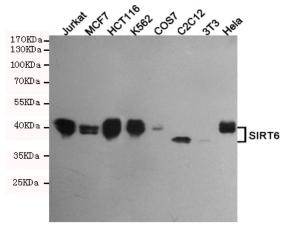
No4: 1

Host: Mouse

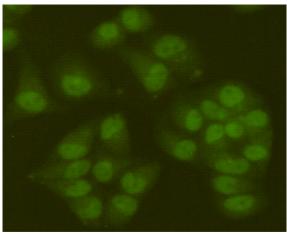
Modifications : Unmodified

Products Images

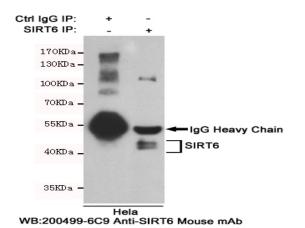
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Western blot analysis of extracts from Jurkat, MCF7, HCT116, K562, COS7, C2C12, 3T3 and Hela cell lysates using SIRT6 mouse mAb (1:500 diluted). Predicted band size: 42,36KDa. Observed band size: 42,36KDa.



Immunofluorescent analysis of Hela cells fixed fixed by 4% paraformaldehyde and using SIRT6 mouse mAb (dilution 1:100).



Immunoprecipitation analysis of Hela cell lysates using SIRT6 mouse mAb.