

Mi2-α Monoclonal Antibody

Catalog No: YM1062

Reactivity: Human; Mouse; Rat; Bovine; Dog; Pig

Applications: WB;IHC;IF;FCM

Target: Mi2-a

Gene Name: CHD3

Protein Name: Chromodomain-helicase-DNA-binding protein 3

Human Gene Id: 1107

Human Swiss Prot

No:

Immunogen: Purified recombinant human Mi2-α (C-terminus) protein fragments expressed in

E.coli.

Q12873

Specificity: Mi2-a Monoclonal Antibody detects endogenous levels of Mi2-a protein.

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

Source: Monoclonal, Mouse

Dilution: WB 1:1000 - 1:2000. IHC 1:500 - 1:1000. IF 1:100 - 1:500. Flow cytometry:

1:100 - 1:200. Not yet tested in other applications.

Purification : Affinity purification

Concentration: 1 mg/ml

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

Molecularweight: 227kD

Background: This gene encodes a member of the CHD family of proteins which are

characterized by the presence of chromo (chromatin organization modifier)

domains and SNF2-related helicase/ATPase domains. This protein is one of the components of a histone deacetylase complex referred to as the Mi-2/NuRD complex which participates in the remodeling of chromatin by deacetylating histones. Chromatin remodeling is essential for many processes including transcription. Autoantibodies against this protein are found in a subset of patients with dermatomyositis. Three alternatively spliced transcripts encoding different isoforms have been described. [provided by RefSeq, Jul 2008],

Function:

disease:One of the main antigens reacting with anti-MI-2 positive sera of dermatomyositis.,function:Probable transcription regulator.,sequence caution:Differs from position 1967 onward for unknown reasons.,similarity:Belongs to the SNF2/RAD54 helicase family.,similarity:Contains 1 helicase ATP-binding domain.,similarity:Contains 1 helicase C-terminal domain.,similarity:Contains 2 chromo domains.,similarity:Contains 2 PHD-type zinc fingers.,subunit:Central component of the nucleosome remodeling and histone deacetylase (NuRD) repressive complex. Interacts with TRIM28 and SERBP1. Interacts via its C-terminal region with HABP4.,tissue specificity:Widely expressed.,

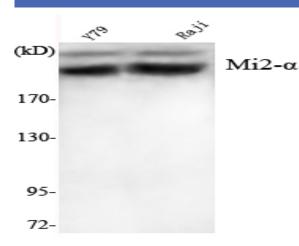
Subcellular Location:

Nucleus, PML body . Nucleus . Cytoplasm, cytoskeleton, microtubule organizing center, centrosome . Associates with centrosomes in interphase and mitosis. .

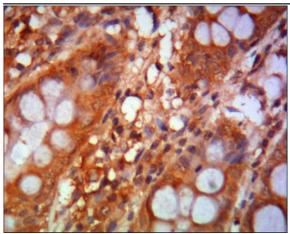
Expression:

Widely expressed.

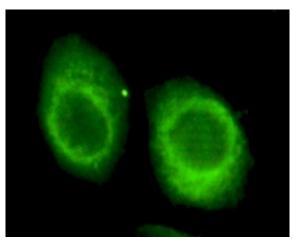
Products Images



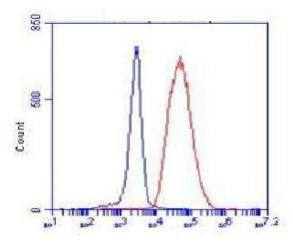
Western Blot analysis using Mi2-α Monoclonal Antibody against Y7P, Raji cell lysate.



Immunohistochemistry analysis of paraffin-embedded human colon using Mi2- α Monoclonal Antibody.



Immunofluorescence analysis of HeLa cells using Mi2- $\!\alpha$ Monoclonal Antibody.



Flow cytometric analysis of K562 cells stained with Mi2- α Monoclonal Antibody (red), followed by FITC-conjugated goat anti-mouse IgG. Blue line histogram represents the isotype control, normal mouse IgG.