# SANTA CRUZ BIOTECHNOLOGY, INC.

# Mig-6 (D-1): sc-137154



## BACKGROUND

Mitogen-inducible gene 6 protein (Mig-6), also designated Gene 33 or RALT, belongs to the Mig-6 family. The gene encoding for Mig-6 maps to chromosome 1p36.23. Mig-6 is a cytoplasmic protein acting as a feedback inhibitor of ErbB-2 mitogenic function and can suppress ErbB-2 oncogenic activity. The expression of Mig-6 is upregulated with cell growth. Mig-6 binds to the epidermal growth factor receptor (EGFR) upon EGF stimulation and is considered a negative feedback regulator of EGFR and a potential tumor suppressor. Mig-6 induces transcriptional activation of NF $\kappa$ B by binding to its inhibitor I $\kappa$ B\alpha. It enables the cell to respond persistently to chronic stress. Mig-6 mRNA levels increase in response to stress such as diabetic nephropathy, vasoactive peptides or mechanical strain. Mig-6 is expressed in liver, placenta and lung.

## **CHROMOSOMAL LOCATION**

Genetic locus: ERRFI1 (human) mapping to 1p36.23; Errfi1 (mouse) mapping to 4 E2.

# SOURCE

Mig-6 (D-1) is a mouse monoclonal antibody raised against amino acids 156-280 mapping within an internal region of Mig-6 of human origin.

## PRODUCT

Each vial contains 200  $\mu$ g lgG<sub>1</sub> kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

Mig-6 (D-1) is available conjugated to agarose (sc-137154 AC), 500  $\mu$ g/0.25 ml agarose in 1 ml, for IP; to HRP (sc-137154 HRP), 200  $\mu$ g/ml, for WB, IHC(P) and ELISA; to either phycoerythrin (sc-137154 PE), fluorescein (sc-137154 FITC), Alexa Fluor<sup>®</sup> 488 (sc-137154 AF488), Alexa Fluor<sup>®</sup> 546 (sc-137154 AF546), Alexa Fluor<sup>®</sup> 594 (sc-137154 AF594) or Alexa Fluor<sup>®</sup> 647 (sc-137154 AF647), 200  $\mu$ g/ml, for WB (RGB), IF, IHC(P) and FCM; and to either Alexa Fluor<sup>®</sup> 680 (sc-137154 AF680) or Alexa Fluor<sup>®</sup> 790 (sc-137154 AF790), 200  $\mu$ g/ml, for Near-Infrared (NIR) WB, IF and FCM.

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#### **APPLICATIONS**

Mig-6 (D-1) is recommended for detection of Mig-6 of mouse, rat and human origin by Western Blotting (starting dilution 1:100, dilution range 1:100-1:1000), immunoprecipitation [1-2  $\mu$ g per 100-500  $\mu$ g of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

Suitable for use as control antibody for Mig-6 siRNA (h): sc-45704, Mig-6 siRNA (m): sc-45705, Mig-6 shRNA Plasmid (h): sc-45704-SH, Mig-6 shRNA Plasmid (m): sc-45705-SH, Mig-6 shRNA (h) Lentiviral Particles: sc-45704-V and Mig-6 shRNA (m) Lentiviral Particles: sc-45705-V.

Molecular Weight of Mig-6: 53 kDa.

Positive Controls: NIH/3T3 whole cell lysate: sc-2210, A549 cell lysate: sc-2413 or Mig-6 (m): 293T Lysate: sc-125619.

#### STORAGE

Store at 4° C, \*\*DO NOT FREEZE\*\*. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

# DATA





Mig-6 (D-1): sc-137154. Western blot analysis of Mig-6 expression in non-transfected 293T: sc-117752 (**A**), mouse Mig-6 transfected 293T: sc-125619 (**B**) and A549 (**C**) whole cell lysates.

Mig-6 (D-1): sc-137154. Western blot analysis of Mig-6 expression in NIH/3T3 whole cell lysate ( $\pmb{A}$ ) and rat brain tissue extract ( $\pmb{B}$ ).

#### **SELECT PRODUCT CITATIONS**

- Liu, N., et al. 2012. Chk1 phosphorylates the tumour suppressor Mig-6, regulating the activation of EGF signalling. EMBO J. 31: 2365-2377.
- 2. Sun, M., et al. 2016. Type I  $\gamma$  phosphatidylinositol phosphate 5-kinase i5 controls the ubiquitination and degradation of the tumor suppressor mitogen-inducible gene 6. J. Biol. Chem. 291: 21461-21473.
- Shangguan, Y., et al. 2017. Glucocorticoid mediates prenatal caffeine exposure-induced endochondral ossification retardation and its molecular mechanism in female fetal rats. Cell Death Dis. 8: e3157.
- 4. Mancini, M., et al. 2021. Generation and characterization of a new preclinical mouse model of EGFR-driven lung cancer with MET-induced osimertinib resistance. Cancers 13: 3441.
- Bosse, K.R., et al. 2022. Serial profiling of circulating tumor DNA identifies dynamic evolution of clinically actionable genomic alterations in hgh-risk neuroblastoma. Cancer Discov. 12: 2800-2819.
- Deng, Z., et al. 2023. Temporal transcriptome features identify early skeletal commitment during human epiphysis development at single-cell resolution. iScience 26: 107200.
- 7. Chen, Y.C., et al. 2023. Glucolipotoxic stress-induced Mig-6 desensitizes EGFR signaling and promotes pancreatic  $\beta$  cell death. Metabolites 13: 627.
- Zhao, H. and Mao, H. 2024. ERRFI1 exacerbates hepatic ischemia reperfusion injury by promoting hepatocyte apoptosis and ferroptosis in a GRB2dependent manner. Mol. Med. 30: 82.

## **RESEARCH USE**

For research use only, not for use in diagnostic procedures.

#### **PROTOCOLS**

See our web site at www.scbt.com for detailed protocols and support products.