

B23 (NA24): sc-53175

BACKGROUND

The transport of proteins across the nuclear envelope is a selective, multi-step process involving several cytoplasmic factors. Proteins must be recognized as import substrates, dock at the nuclear pore complex, and translocate across the nuclear envelope in an ATP-dependent fashion. Several cytosolic and nuclear proteins that are central to this process have been identified. For example, two cytosolic factors critically involved in the recognition and docking process are the karyopherin α and karyopherin β proteins. The karyopherin holoenzyme is a heterodimer of α and β subunits. The nuclear protein B23 (also referred to as nucleophosmin) is involved in ribosomal assembly and rRNA transport. B23 is an abundant protein that is highly phosphorylated by Cdc2 kinase during mitosis.

REFERENCES

1. Moroianu, J., et al. 1995. Protein export from the nucleus requires the GTPase Ran and GTP hydrolysis. *Proc. Natl. Acad. Sci. USA* 92: 4318-4322.
2. Chou, Y.H., et al. 1995. Cell cycle phase-dependent changes of localization and oligomerization states of nucleophosmin/B23. *Biochem. Biophys. Res. Commun.* 217: 313-325.

CHROMOSOMAL LOCATION

Genetic locus: NPM1 (human) mapping to 5q35.1; Npm1 (mouse) mapping to 11 A4.

SOURCE

B23 (NA24) is a mouse monoclonal antibody raised against a recombinant protein corresponding to the N-terminus of B23 of human origin.

PRODUCT

Each vial contains 200 μ g IgG₁ kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

APPLICATIONS

B23 (NA24) is recommended for detection of B23 of mouse, rat and human origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000), immunoprecipitation [1-2 μ g per 100-500 μ g of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500) and immunohistochemistry (including paraffin-embedded sections) (starting dilution 1:50, dilution range 1:50-1:500).

Suitable for use as control antibody for B23 siRNA (h): sc-29771, B23 siRNA (m): sc-29772, B23 shRNA Plasmid (h): sc-29771-SH, B23 shRNA Plasmid (m): sc-29772-SH, B23 shRNA (h) Lentiviral Particles: sc-29771-V and B23 shRNA (m) Lentiviral Particles: sc-29772-V.

Molecular Weight of B23: 40 kDa.

Positive Controls: MCF7 whole cell lysate: sc-2206, K-562 whole cell lysate: sc-2203 or CCRF-CEM cell lysate: sc-2225.

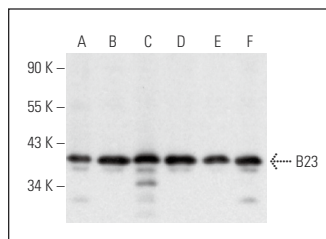
RESEARCH USE

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

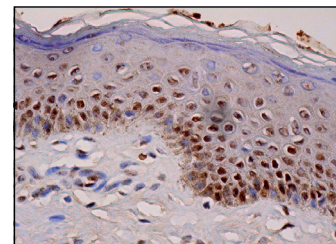
STORAGE

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

DATA



B23 (NA24): sc-53175. Western blot analysis of B23 expression in HEL 92.1.7 (A), LNCaP (B), HeLa (C), K-562 (D), CCRF-CEM (E) and MCF7 (F) whole cell lysates.



B23 (NA24): sc-53175. Immunoperoxidase staining of formalin fixed, paraffin-embedded human vulva/anal skin tissue showing nucleolar and nuclear staining of epidermal cells.

SELECT PRODUCT CITATIONS

1. Kikuta, K., et al. 2009. Nucleophosmin as a candidate prognostic biomarker of Ewing's sarcoma revealed by proteomics. *Clin. Cancer Res.* 15: 2885-2894.
2. Piovan, C., et al. 2012. Oncosuppressive role of p53-induced miR-205 in triple negative breast cancer. *Mol. Oncol.* 6: 458-472.
3. Haga, A., et al. 2013. Interactomic approach for evaluating nucleophosmin-binding proteins as biomarkers for Ewing's sarcoma. *Electrophoresis* 34: 1670-1678.
4. Raman, N., et al. 2014. mTOR signaling regulates nucleolar targeting of the SUMO-specific isopeptidase SENP3. *Mol. Cell. Biol.* 34: 4474-4484.
5. Caudron-Herger, M., et al. 2015. Alu element-containing RNAs maintain nucleolar structure and function. *EMBO J.* 34: 2758-2774.
6. Bober, J., et al. 2016. Identification of new FGF1 binding partners-implications for its intracellular function. *IUBMB Life* 68: 242-251.
7. Brodská, B., et al. 2017. Localization of AML-related nucleophosmin mutant depends on its subtype and is highly affected by its interaction with wild-type NPM. *PLoS ONE* 12: e0175175.
8. Sabbir, M.G. 2018. Loss of Ca²⁺/calmodulin dependent protein kinase 2 leads to aberrant transferrin phosphorylation and trafficking: a potential biomarker for Alzheimer's disease. *Front. Mol. Biosci.* 5: 99.
9. Caudron-Herger, M., et al. 2019. R-DeeP: proteome-wide and quantitative identification of RNA-dependent proteins by density gradient ultracentrifugation. *Mol. Cell* 75: 184-199.e10.



See **B23 (E-3): sc-271737** for B23 antibody conjugates, including AC, HRP, FITC, PE, and Alexa Fluor® 488, 546, 594, 647, 680 and 790.