

# BAP1 (C-4): sc-28383

## BACKGROUND

Mutations within the BRCA1 gene, localized to chromosome 17q, are believed to account for approximately 45% of families with increased incidence of both early-onset breast cancer and ovarian cancer. The BRCA1 gene is expressed in numerous tissues, including breast and ovary, and encodes a predicted protein of 1,863 amino acids. This protein contains a RING domain near the N-terminus and appears to encode a tumor suppressor. BARD1 (BRCA1-associated RING domain protein 1) and BAP1 (BRCA1-associated protein 1) have both been shown to bind to the N-terminus of BRCA1 and are potential mediators of tumor suppression. BARD1 contains an N-terminal RING domain and three tandem ankyrin repeats. The C-terminus of BARD1 contains a region with sequence homology to BRCA1, termed the BRCT domain. BAP1 is a ubiquitin hydrolase and has been shown to enhance BRCA1-mediated cell growth suppression.

## REFERENCES

- Hall, J.M., et al. 1990. Linkage of early-onset familial breast cancer to chromosome 17q21. *Science* 250: 1684-1689.
- Narod, S.A., et al. 1991. Familial breast-ovarian cancer locus on chromosome 17q12-q23. *Lancet* 338: 82-83.

## CHROMOSOMAL LOCATION

Genetic locus: BAP1 (human) mapping to 3p21.1; Bap1 (mouse) mapping to 14 B.

## SOURCE

BAP1 (C-4) is a mouse monoclonal antibody raised against amino acids 430-729 of BAP1 of human origin.

## PRODUCT

Each vial contains 200 µg IgG<sub>1</sub> in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

## APPLICATIONS

BAP1 (C-4) is recommended for detection of BAP1 of mouse, rat and human origin by Western Blotting (starting dilution 1:100, dilution range 1:100-1:500), immunoprecipitation [1-2 µg per 100-500 µg of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500), immunohistochemistry (including paraffin-embedded sections) (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 BAP1 siRNA (h): sc-29787, BAP1 siRNA (m): sc-29788, BAP1 shRNA Plasmid (h): sc-29787-SH, BAP1 shRNA Plasmid (m): sc-29788-SH, BAP1 shRNA (h) Lentiviral Particles: sc-29787-V and BAP1 shRNA (m) Lentiviral Particles: sc-29788-V.

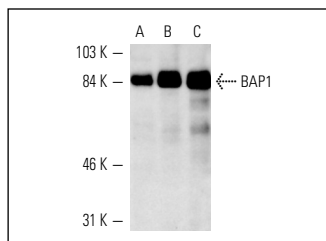
Molecular Weight of BAP1: 91 kDa.

Positive Controls: IB4 whole cell lysate: sc-364780, KNRK whole cell lysate: sc-2214 or SK-BR-3 cell lysate: sc-2218.

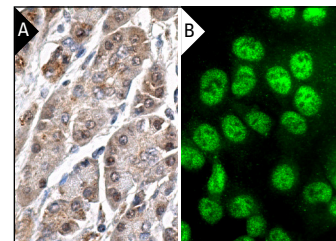
## STORAGE

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

## DATA



BAP1 (C-4): sc-28383. Western blot analysis of BAP1 expression in SK-BR-3 (A), IB4 (B) and KNRK (C) whole cell lysates.



BAP1 (C-4): sc-28383. Immunoperoxidase staining of formalin fixed, paraffin-embedded human pancreas tissue showing nuclear and cytoplasmic staining of glandular cells (A). Immunofluorescence staining of formalin-fixed HepG2 cells showing nuclear localization (B).

## SELECT PRODUCT CITATIONS

- Ventii, K.H., et al. 2008. BRCA1-associated protein-1 is a tumor suppressor that requires deubiquitinating activity and nuclear localization. *Cancer Res.* 68: 6953-6962.
- Nishikawa, H., et al. 2009. BRCA1-associated protein 1 interferes with BRCA1/BARD1 RING heterodimer activity. *Cancer Res.* 69: 111-119.
- Bott, M., et al. 2011. The nuclear deubiquitinase BAP1 is commonly inactivated by somatic mutations and 3p21.1 losses in malignant pleural mesothelioma. *Nat. Genet.* 43: 668-672.
- Wiesner, T., et al. 2012. A distinct subset of atypical Spitz tumors is characterized by BRAF mutation and loss of BAP1 expression. *Am. J. Surg. Pathol.* 36: 818-830.
- Carbone, M., et al. 2012. BAP1 cancer syndrome: malignant mesothelioma, uveal and cutaneous melanoma, and MBAITs. *J. Transl. Med.* 10: 179.
- Yoshikawa, Y., et al. 2012. Frequent inactivation of the BAP1 gene in epithelioid-type malignant mesothelioma. *Cancer Sci.* 103: 868-874.
- Gammon, B., et al. 2013. Clumped perinuclear BAP1 expression is a frequent finding in sporadic epithelioid Spitz tumors. *J. Cutan. Pathol.* 40: 538-542.
- Kerl, K., et al. 2013. Two-dimensional visualization of multicolor FISH-generated data as a helpful tool for the analysis and understanding of cytogenetic and chromosomal alterations in melanocytic lesions. *Am. J. Dermatopathol.* 35: 151-158.
- Popova T., et al. 2013. Germline BAP1 mutations predispose to renal cell carcinomas. *Am. J. Hum. Genet.* 92: 974-980.

## RESEARCH USE

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