Product datasheet MON10216



# Mouse anti-BAP1, clone C-4 (Monoclonal)

Clone no. C-4 MONOSAN

Product name Mouse anti-BAP1, clone C-4 (Monoclonal)

**Host** Mouse

Applications IHC-P (1:50-1:500)

Species reactivity Human

Conjugate -

Immunogen Amino acids 430-729 of BAP1 of human origin

lsotype lgG1

**Clonality** Monoclonal

Clone number C-4

Size 1 ml

**Concentration** n/a

**Format** Concentrate

Storage buffer PBS with < 0.1% sodium azide and 0.1% gelatin

Storage until expiry date 2-8°C

## FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC PROCEDURES



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#### Additional info

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 Nterminus 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. Pre-treatment: Heat induced epitope retrieval in 10 mM citrate buffer, pH6.0, or in 50 mM Tris buffer pH9.5, for 20 minutes is required for IHC staining on formalin-fixed, paraffin embedded tissue sections. Control tissue Pancreas, breast carcinoma, ovarian carcinoma. Staining Nuclear and cytoplasmic.

### References

- 1. Hall, I.M., et al. 1990, Science 250: 1684-1689.
- 2 Yoshikawa, Y., et al. 2012, Cancer Sci. 103: 868-874.
- 3. Gammon, B., et al. 2013, J. Cutan. Pathol. 40: 538-542.
- 4. Kerl, K., et al. 2013, Am. J. Dermatopathol. 35: 151-158.
- 5. Popova T., et al. 2013, Am. J. Hum. Genet. 92: 974-980.

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