# SANTA CRUZ BIOTECHNOLOGY, INC.

# Mad2 (yS-19): sc-6332



# BACKGROUND

Cell cycle progression is subject to arrest at the mitotic spindle assembly checkpoint in response to incorrect spindle fiber assembly. MAD2 (for mitotic arrest-deficient) is a component of the mitotic spindle checkpoint. Cells with mutated MAD2 do not undergo mitotic arrest in response to incorrect spindle fiber assembly, which results in missegregation and eventual cell death. A breast carcinoma cell line with reduced MAD2 expression, T47D, was shown to complete mitosis in the presence of nocodazole, an inhibitor of mitotic spindle assembly. MAD2 is localized to unattached kinetochores during prometaphase and disassociates upon spindle fiber attachment, indicating that MAD2 regulates kinetochore binding to the spindle fibers. Human MAD2 has also been shown to associate with Insulin receptor (IR), but not IGFIR, implicating MAD2 as a mediator for IR-specific signaling. MAD2B, a MAD2 homolog, is required for the execution of the mitotic checkpoint monitoring the kinetochore-spindle attachment process and if the process is not complete, MAD2B delays the onset of anaphase.

#### REFERENCES

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- 2. Glotzer, M. 1996. Mitosis: don't get mad, get even. Curr. Biol. 6: 1592-1594.
- Chen, R.H., et al. 1996. Association of spindle assembly checkpoint component XMAD2 with unattached kinetochores. Science 274: 242-246.
- 4. Li, Y. et al. 1996. Identification of a human mitotic checkpoint gene: hsMAD2. Science 274: 246-248.
- O'Neill, T.J., et al. 1997. Interaction of MAD2 with the carboxyl terminus of the Insulin receptor but not with the IGFIR. Evidence for release from the Insulin receptor after activation. J. Biol. Chem. 272: 10035-10040.
- Liu, S.T., et al. 2003. Human CENP-I specifies localization of CENP-F, MAD1 and MAD2 to kinetochores and is essential for mitosis. Nat. Cell Biol. 5: 341-345.
- Michel, L., et al. 2004. Complete loss of the tumor suppressor MAD2 causes premature cyclin B degradation and mitotic failure in human somatic cells. Proc. Natl. Acad. Sci. USA 101: 4459-4464.

#### SOURCE

Mad2 (yS-19) is an affinity purified goat polyclonal antibody raised against a peptide mapping at the N-terminus of Mad2 of *Saccharomyces cerevisiae* origin.

#### PRODUCT

Each vial contains 200  $\mu g$  lgG in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

Blocking peptide available for competition studies, sc-6332 P, (100  $\mu$ g peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% BSA).

#### **STORAGE**

Store at 4° C, \*\*D0 NOT FREEZE\*\*. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.

#### APPLICATIONS

Mad2 (yS-19) is recommended for detection of Mad2 of *Saccharomyces cerevisiae* origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000) and immunoprecipitation [1-2 µg per 100-500 µg of total protein (1 ml of cell lysate)].

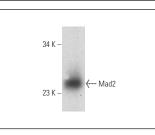
Molecular Weight of Mad2: 25 kDa.

Positive Controls: *Saccaromyces cerevisiae* whole cell lysate or EGY48 whole cell lysate: sc-364775.

#### **RECOMMENDED SECONDARY REAGENTS**

To ensure optimal results, the following support (secondary) reagents are recommended: 1) Western Blotting: use donkey anti-goat IgG-HRP: sc-2020 (dilution range: 1:2000-1:100,000) or Cruz Marker<sup>™</sup> compatible donkey anti-goat IgG-HRP: sc-2033 (dilution range: 1:2000-1:5000), Cruz Marker<sup>™</sup> Molecular Weight Standards: sc-2035, TBS Blotto A Blocking Reagent: sc-2333 and Western Blotting Luminol Reagent: sc-2048. 2) Immunoprecipitation: use Protein A/G PLUS-Agarose: sc-2003 (0.5 ml agarose/2.0 ml).

#### DATA



Mad2 (yS-19): sc-6332. Western blot analysis of Mad2 expression in EGY48 whole cell lysate.

## **RESEARCH USE**

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

## PROTOCOLS

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

# MONOS Satisfation Guaranteed

Try Mad2 (C-3): sc-365813, our highly recommended monoclonal alternative to Mad2 (yS-19).