# SANTA CRUZ BIOTECHNOLOGY, INC.

# RPRC1 (E-14): sc-163319



### BACKGROUND

Chromosome 1 is the largest human chromosome spanning about 260 million base pairs and making up 8% of the human genome. There are about 3,000 genes on chromosome 1, and considering the great number of genes there are also a large number of diseases associated with chromosome 1. Notably, the rare aging disease Hutchinson-Gilford progeria is associated with the LMNA gene which encodes lamin A. When defective, the LMNA gene product can build up in the nucleus and cause characteristic nuclear blebs. The mechanism of rapidly enhanced aging is unclear and is a topic of continuing exploration. The MUTYH gene is located on chromosome 1 and is partially responsible for familial adenomatous polyposis. Stickler syndrome, Parkinsons, Gaucher disease and Usher syndrome are also associated with chromosome 1. A breakpoint has been identified in 1q which disrupts the DISC1 gene and is linked to schizophrenia. Aberrations in chromosome 1 are found in a variety of cancers including head and neck cancer, malignant melanoma and multiple myeloma. The RPRC1 gene product has been provisionally designated RPRC1 pending further characterization.

## REFERENCES

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- 3. Weise, A., et al. 2005. New insights into the evolution of chromosome 1. Cytogenet. Genome Res. 108: 217-222.
- 4. Lans, H., et al. 2006. Cell biology: aging nucleus gets out of shape. Nature 440: 32-34.
- 5. Gregory, S.G., et al. 2006. The DNA sequence and biological annotation of human chromosome 1. Nature 441: 315-321.
- 6. Hennah, W., et al. 2006. Genes and schizophrenia: beyond schizophrenia: the role of DISC-1 in major mental illness. Schizophr. Bull. 32: 409-416.
- 7. Marzin, Y., et al. 2006. Chromosome 1 abnormalities in multiple myeloma. Anticancer Res. 26: 953-959.
- 8. McClintock, D., et al. 2006. Hutchinson-Gilford progeria mutant Lamin A primarily targets human vascular cells as detected by an anti-Lamin A G608G antibody. Proc. Natl. Acad. Sci. USA 103: 2154-2159.

#### CHROMOSOMAL LOCATION

Genetic locus: MAP7D1 (human) mapping to 1p34.3; Mtap7d1 (mouse) mapping to 4 D2.2.

#### SOURCE

RPRC1 (E-14) is an affinity purified goat polyclonal antibody raised against a peptide mapping near the C-terminus of RPRC1 of human origin.

#### PRODUCT

Each vial contains 200 µg lgG in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

Blocking peptide available for competition studies, sc-163319 P, (100 µg peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% BSA).

### **APPLICATIONS**

RPRC1 (E-14) is recommended for detection of RPRC1 of mouse, rat and human origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000), 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).

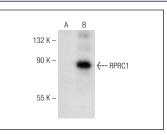
Suitable for use as control antibody for RPRC1 siRNA (h): sc-88733, RPRC1 siRNA (m): sc-153115, RPRC1 shRNA Plasmid (h): sc-88733-SH, RPRC1 shRNA Plasmid (m): sc-153115-SH, RPRC1 shRNA (h) Lentiviral Particles: sc-88733-V and RPRC1 shRNA (m) Lentiviral Particles: sc-153115-V.

Molecular Weight (predicted) of RPRC1: 93 kDa.

Molecular Weight (observed) of RPRC1: 102 kDa.

Positive Controls: RPRC1 (h): 293T Lysate: sc-111888.

#### DATA





RPRC1 (E-14): sc-163319. Western blot analysis of RPRC1 expression in non-transfected: sc-117752 (A) and human RPRC1 transfected: sc-111888 (B) 293T whole cell lysates

RPRC1 (E-14): sc-163319. Immunoperoxidase staining of formalin fixed, paraffin-embedded human heart muscle tissue showing cytoplasmic staining of mvocvtes

#### **STORAGE**

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

#### PROTOCOLS

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