SMUG1 (D-2): sc-377370



The Power to Overtio

BACKGROUND

The base excision repair (BER) pathway removes incorrect bases (uracil) or damaged bases (3-methyladenine) from chromatin. Each BER enzyme system addresses a specific type of base damage. Uracil-DNA glycosylases, UNG2 and SMUG1 (single-strand selective monofunctional uracil DNA glycosylase) remove uracil from both double- and single-stranded DNA in nucleosomes (chromatin core particle). The uracil-excising enzyme family shares structural and functional conservation with minimal sequence conservation. The human SMUG1 gene maps to chromosome 12q13.13.

REFERENCES

- Haushalter, K.A., et al. 1999. Identification of a new uracil-DNA glycosylase family by expression cloning using synthetic inhibitors. Curr. Biol. 9: 174-185.
- Boorstein, R.J., et al. 2001. Definitive identification of mammalian 5-hydroxymethyluracil DNA N-glycosylase activity as SMUG1. J. Biol. Chem. 276: 41991-41997.

CHROMOSOMAL LOCATION

Genetic locus: SMUG1 (human) mapping to 12q13.13; Smug1 (mouse) mapping to 15 F3.

SOURCE

SMUG1 (D-2) is a mouse monoclonal antibody specific for an epitope mapping between amino acids 39-75 near the N-terminus of SMUG1 of human origin.

PRODUCT

Each vial contains 200 $\mu g \ lgG_1$ kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

Blocking peptide available for competition studies, sc-377370 P, (100 μg peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% stabilizer protein).

APPLICATIONS

SMUG1 (D-2) is recommended for detection of SMUG1 of mouse, rat and human origin by Western Blotting (starting dilution 1:100, 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 solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

Suitable for use as control antibody for SMUG1 siRNA (h): sc-106768, SMUG1 siRNA (m): sc-153643, SMUG1 shRNA Plasmid (h): sc-106768-SH, SMUG1 shRNA Plasmid (m): sc-153643-SH, SMUG1 shRNA (h) Lentiviral Particles: sc-106768-V and SMUG1 shRNA (m) Lentiviral Particles: sc-153643-V.

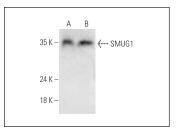
Molecular Weight of SMUG1: 34 kDa.

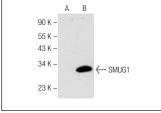
Positive Controls: SMUG1 (m): 293T Lysate: sc-123667, K-562 whole cell lysate: sc-2203 or SJRH30 cell lysate: sc-2287.

RECOMMENDED SUPPORT REAGENTS

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-lgG κ BP-HRP: sc-516102 or m-lgG κ BP-HRP (Cruz Marker): sc-516102-CM (dilution range: 1:1000-1:10000), Cruz MarkerTM Molecular Weight Standards: sc-2035, UltraCruz® Blocking Reagent: sc-516214 and Western Blotting Luminol Reagent: sc-2048. 2) Immunoprecipitation: use Protein A/G PLUS-Agarose: sc-2003 (0.5 ml agarose/2.0 ml). 3) Immunofluorescence: use m-lgG κ BP-FITC: sc-516140 or m-lgG κ BP-PE: sc-516141 (dilution range: 1:50-1:200) with UltraCruz® Mounting Medium: sc-24941 or UltraCruz® Hard-set Mounting Medium: sc-359850.

DATA





SMUG1 (D-2): sc-377370. Western blot analysis of SMUG1 expression in K-562 (**A**) and SJRH30 (**B**) whole cell lysates.

SMUG1 (D-2): sc-377370. Western blot analysis of SMUG1 expression in non-transfected: sc-117752 (A) and mouse SMUG1 transfected: sc-123667 (B) 293T whole cell lysates.

SELECT PRODUCT CITATIONS

 Tan, H.W., et al. 2019. Lasting DNA damage and aberrant DNA repair gene expression profile are associated with post-chronic cadmium exposure in human bronchial epithelial cells. Cells 8: 842.

STORAGE

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

RESEARCH USE

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

PROTOCOLS

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

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