

DcpS (A-12): sc-393226

BACKGROUND

Eukaryotic cells primarily utilize exoribonucleases and decapping enzymes to degrade their mRNA. DcpS is a scavenger pyrophosphatase that hydrolyzes the residual cap structure following 3' to 5' decay of an mRNA. Following mRNA degradation DcpS releases N-7 methyl guanosine monophosphate and 5'-diphosphate terminated cap or mRNA products. The central histidine within the DcpS HIT motif is critical for decapping activity and defines the HIT motif as a new mRNA decapping domain, making DcpS the first member of the HIT family of proteins with a defined biological function. HIT proteins are homodimeric and contain two conserved 100-amino-acid HIT fold domains with independent active sites that are each sufficient to bind and hydrolyze cognate substrates.

REFERENCES

1. Fireman, P. 1992. Diagnosis of sinusitis in children: emphasis on the history and physical examination. *J. Allergy Clin. Immunol.* 90: 433-436.
2. Wang, Z. and Kiledjian, M. 2001. Functional link between the mammalian exosome and mRNA decapping. *Cell* 107: 751-762.
3. Liu, H., et al. 2002. The scavenger mRNA decapping enzyme DcpS is a member of the HIT family of pyrophosphatases. *EMBO J.* 21: 4699-4708.
4. Wang, Z., et al. 2002. The hDcp2 protein is a mammalian mRNA decapping enzyme. *Proc. Natl. Acad. Sci. USA* 99: 12663-12668.
5. Gu, M., et al. 2004. Insights into the structure, mechanism, and regulation of scavenger mRNA decapping activity. *Mol. Cell* 14: 67-80.

CHROMOSOMAL LOCATION

Genetic locus: DCPS (human) mapping to 11q24.2; Dcps (mouse) mapping to 9 A4.

SOURCE

DcpS (A-12) is a mouse monoclonal antibody specific for an epitope mapping between amino acids 97-124 within an internal region of DcpS of human origin.

PRODUCT

Each vial contains 200 µg IgG_{2b} kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

DcpS (A-12) is available conjugated to agarose (sc-393226 AC), 500 µg/0.25 ml agarose in 1 ml, for IP; to HRP (sc-393226 HRP), 200 µg/ml, for WB, IHC(P) and ELISA; to either phycoerythrin (sc-393226 PE), fluorescein (sc-393226 FITC), Alexa Fluor® 488 (sc-393226 AF488), Alexa Fluor® 546 (sc-393226 AF546), Alexa Fluor® 594 (sc-393226 AF594) or Alexa Fluor® 647 (sc-393226 AF647), 200 µg/ml, for WB (RGB), IF, IHC(P) and FCM; and to either Alexa Fluor® 680 (sc-393226 AF680) or Alexa Fluor® 790 (sc-393226 AF790), 200 µg/ml, for Near-Infrared (NIR) WB, IF and FCM.

Blocking peptide available for competition studies, sc-393226 P, (100 µg peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% stabilizer protein).

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APPLICATIONS

DcpS (A-12) is recommended for detection of DcpS of mouse, rat and human origin by Western Blotting (starting dilution 1:100, 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).

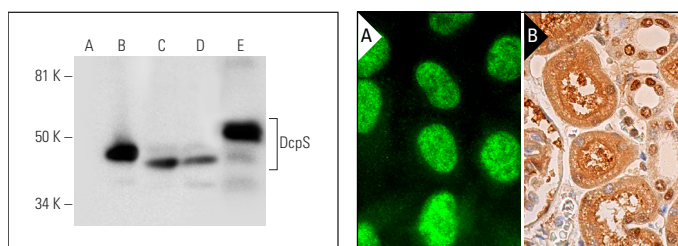
DcpS (A-12) is also recommended for detection of DcpS in additional species, including equine and porcine.

Suitable for use as control antibody for DcpS siRNA (h): sc-44389, DcpS siRNA (m): sc-44390, DcpS shRNA Plasmid (h): sc-44389-SH, DcpS shRNA Plasmid (m): sc-44390-SH, DcpS shRNA (h) Lentiviral Particles: sc-44389-V and DcpS shRNA (m) Lentiviral Particles: sc-44390-V.

Molecular Weight of DcpS: 40 kDa.

Positive Controls: DcpS (m): 293T Lysate: sc-119688, HeLa whole cell lysate: sc-2200 or human liver extract: sc-363766.

DATA



DcpS (A-12): sc-393226. Western blot analysis of DcpS expression in non-transfected 293T: sc-117752 (A), mouse DcpS transfected 293T: sc-119688 (B) and HeLa (C) whole cell lysates and mouse liver (D) and human liver (E) tissue extracts.

DcpS (A-12): sc-393226. Immunofluorescence staining of methanol-fixed HeLa cells showing nuclear localization (A). Immunoperoxidase staining of formalin fixed, paraffin-embedded human kidney tissue showing cytoplasmic staining of cells in glomeruli and nuclear and cytoplasmic staining of cells in tubules (B).

SELECT PRODUCT CITATIONS

1. Yamauchi, T., et al. 2018. Genome-wide CRISPR-Cas9 screen identifies leukemia-specific dependence on a pre-mRNA metabolic pathway regulated by DcpS. *Cancer Cell* 33: 386-400.e5.
2. Swartzel, J.C., et al. 2022. Targeted degradation of mRNA decapping enzyme DcpS by a VHL-recruiting PROTAC. *ACS Chem. Biol.* 17: 1789-1798.

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.