

PSMB11 (A-5): sc-514317

BACKGROUND

In eukaryotic cells, selective breakdown of cellular proteins is ensured by their ubiquitination and subsequent degradation by the 26S Proteasome. The 26S Proteasome is a protease complex that selectively breaks down proteins that have been modified by polyubiquitin chains. It is made up of two multisubunit complexes: the 20S Proteasome chamber, which serves as the proteolytic core of the complex, and two 19S regulatory particles which recognize and unfold ubiquitinated proteins. The 20S Proteasome chamber contains α subunits (which are structural) and β subunits (which are predominantly catalytic). The outer two rings in the proteasome consist of seven α subunits each, and the inner two rings each consist of seven β subunits. PSMB11 (proteasome subunit β type-11), also known as BETA5T (Proteasome subunit β -5t) is a 300 amino acid protein that plays an important role in CD8-positive T-cells and reduces chymotrypsin-like activity in proteasomes.

REFERENCES

- Orlowski, M., Cardozo, C., Eleuteri, A.M., Kohanski, R., Kam, C.M. and Powers, J.C. 1997. Reactions of [¹⁴C]-3,4-dichloroisocoumarin with subunits of pituitary and spleen multicatalytic proteinase complexes (proteasomes). *Biochemistry* 36: 13946-13953.
- Nandi, D., Woodward, E., Ginsburg, D.B. and Monaco, J.J. 1997. Intermediates in the formation of mouse 20S proteasomes: implications for the assembly of precursor β subunits. *EMBO J.* 16: 5363-5375.
- McCusker, D., Jones, T., Sheer, D. and Trowsdale, J. 1997. Genetic relationships of the genes encoding the human proteasome β subunits and the proteasome PA28 complex. *Genomics* 45: 362-367.
- Razeghi, P., Baskin, K.K., Sharma, S., Young, M.E., Stepkowski, S., Essop, M.F. and Taegtmeier, H. 2006. Atrophy, hypertrophy, and hypoxemia induce transcriptional regulators of the ubiquitin proteasome system in the rat heart. *Biochem. Biophys. Res. Commun.* 342: 361-364.
- Kwak, M.K., Huang, B., Chang, H., Kim, J.A. and Kensler, T.W. 2007. Tissue specific increase of the catalytic subunits of the 26S proteasome by indirect antioxidant dithiolethione in mice: enhanced activity for degradation of abnormal protein. *Life Sci.* 80: 2411-2420.

CHROMOSOMAL LOCATION

Genetic locus: PSMB11 (human) mapping to 14q11.2; Psmb11 (mouse) mapping to 14 C3.

SOURCE

PSMB11 (A-5) is a mouse monoclonal antibody specific for an epitope mapping between amino acids 89-107 within an internal region of PSMB11 of human origin.

PRODUCT

Each vial contains 200 μ g IgM 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-514317 P, (100 μ g peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% stabilizer protein).

APPLICATIONS

PSMB11 (A-5) is recommended for detection of PSMB11 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 PSMB11 siRNA (h): sc-92455, PSMB11 siRNA (m): sc-152555, PSMB11 shRNA Plasmid (h): sc-92455-SH, PSMB11 shRNA Plasmid (m): sc-152555-SH, PSMB11 shRNA (h) Lentiviral Particles: sc-92455-V and PSMB11 shRNA (m) Lentiviral Particles: sc-152555-V.

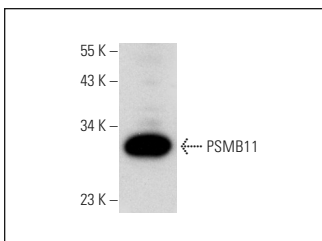
Molecular Weight of PSMB11: 33 kDa.

Positive Controls: MTE1D whole cell lysate: sc-364918.

RECOMMENDED SUPPORT REAGENTS

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-IgG κ BP-HRP: sc-516102 or m-IgG κ BP-HRP (Cruz Marker): sc-516102-CM (dilution range: 1:1000-1:10000), Cruz Marker™ Molecular Weight Standards: sc-2035, UltraCruz® Blocking Reagent: sc-516214 and Western Blotting Luminol Reagent: sc-2048. 2) Immunoprecipitation: use Protein L-Agarose: sc-2336 (0.5 ml agarose/2.0 ml). 3) Immunofluorescence: use m-IgG κ BP-FITC: sc-516140 or m-IgG 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



PSMB11 (A-5): sc-514317. Western blot analysis of PSMB11 expression in MTE1D whole cell lysate.

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.