Proteasome subunit beta type-1 Polyclonal Antibody

Catalog No: #42253



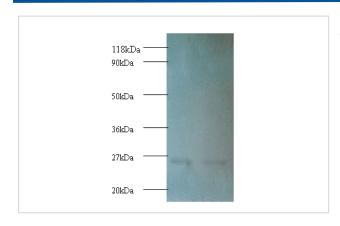
Orders: order@signalwayantibody.com Support: tech@signalwayantibody.com

| Description | Support: tech@signalwayantibody.com |
|-----------------------|--|
| Product Name | Proteasome subunit beta type-1 Polyclonal Antibody |
| Host Species | Rabbit |
| Clonality | Polyclonal |
| Purification | Caprylic Acid Ammonium Sulfate Precipitation purified |
| Applications | WB |
| Species Reactivity | Hu |
| Specificity | The antibody detects endogenous level of total Proteasome subunit beta type-1 polyclonal antibody. |
| Immunogen Type | protein |
| Immunogen Description | Recombinant human Proteasome subunit beta type-1 protein |
| Target Name | Proteasome subunit beta type-1 |
| Other Names | PSC5, PSMB1, Macropain subunit C5 Multicatalytic endopeptidase complex subunit C5 Proteasome |
| | component C5 Proteasome gamma chain |
| Accession No. | Swiss-Prot#: P20618 |
| Uniprot | P20618 |
| GeneID | 5689; |
| Calculated MW | 26.4kd |
| Formulation | Preservative: 0.03% Proclin 300 Constituents: 50% Glycerol, 0.01M PBS, PH 7.4 |
| Storage | Store at -20°C |
| | |

Application Details

Western blotting: □1:500 - 1:1000

Images



All lanes : Proteasome subunit beta 1 antibody at 2ug/mlLane 1 : EC109 whole cell lysateLane 2 : 293T whole cell lysate

SecondaryGoat polyclonal to Rabbit IgG at 1/15000 dilution

Predicted band size : 26.4 kDa Observed band size:26.4 kDa

Background

The proteasome is a multicatalytic proteinase complex which is characterized by its ability to cleave peptides with Arg, Phe, Tyr, Leu, and Glu adjacent to the leaving group at neutral or slightly basic pH. The proteasome has an ATP-dependent proteolytic activity. Cleavage of peptide bonds with very

broad specificity. The 26S proteasome consists of a 20S proteasome core and two 19S regulatory subunits. The 20S proteasome core is composed of 28 subunits that are arranged in four stacked rings, resulting in a barrel-shaped structure. The two end rings are each formed by seven alpha subunits, and the two central rings are each formed by seven beta subunits. The catalytic chamber with the active sites is on the inside of the barrel. Interacts with SERPINB2. Interacts with HIV-1 TAT protein.

References

[1]"Molecular cloning and sequence analysis of cDNAs for five major subunits of human proteasomes (multi-catalytic proteinase complexes)."Tamura T., Lee D.H., Osaka F., Fujiwara T., Shin S., Chung C.H., Tanaka K., Ichihara A.Biochim. Biophys. Acta 1089:95

Note: This product is for in vitro research use only