Vps36 (yN-14): sc-26150



The Power to Question

BACKGROUND

Vacuolar protein sorting mutant 36 (Vps36), a class E Vps protein identified in yeast, is an important component of the trafficking pathway of proteins from the Golgi to the endosome. Vps36 functions as a member of the 155 kDa ESCRT-II complex that transiently associates with the endosomal membrane to sort ubiquitinated endosomal membrane proteins as part of a coordinated cascade of events to select multivesicular body cargoes for delivery to the lumen of the lysosome. Vps36 involvement in post-Golgi endosomal-vacuolar protein sorting is indicated by the fact that Vps mutants secrete unprocessed insulin-containing fusion protein (ICFP), induce a stress-response element and repress pheromone-dependent transcription. Vps36 suppresses the phenotype of targeting defective plasma membrane ATPase (Pma1) by allowing mutant Pma1 to regain movement into the plasma membrane.

REFERENCES

- Luo, W., Chang, A. 2000. An endosome-to-plasma membrane pathway involved in trafficking of a mutant plasma membrane ATPase in yeast. Mol. Biol. Cell 11: 579-92.
- Zhang, B., Chang, A., Kjeldsen, T.B., Arvan, P. 2001. Intracellular retention of newly synthesized insulin in yeast is caused by endoproteolytic processing in the Golgi complex. J. Cell Biol. 153: 1187-98
- 3. Forsberg, H., Hammar, M., Andreasson, C., Moliner, A., Ljungdahl, P.O. 2001. Suppressors of ssy1 and ptr3 null mutations define novel amino acid sensor-independent genes in *S. Cerevisiae*. Genetics 158: 973-88.
- Babst, M., Katzmann, D.J., Snyder, W.B., Wendland, B., Emr, S.D. 2002. Endosome-associated complex, ESCRT-II, recruits transport machinery for protein sorting at the multivesicular body. Dev. Cell 3: 83-9.
- Burchett, S.A., Flanary, P., Aston, C., Jiang, L., Young, K.H., Uetz, P., Fields, S., Dohlman, H.G. 2002. Regulation of stress response signaling by the N-terminal dishevelled/EGL-10/pleckstrin domain of Sst2, a regulator of G protein signaling in *S. Cerevisiae*. J. Bio.I Chem. 277: 22156-67.

SOURCE

Vps36 (yN-14) is an affinity purified goat polyclonal antibody raised against a peptide mapping at the N-terminus of Vps36 of *Saccharomyces cerevisiae* 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-26150 P, (100 μ g peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% BSA).

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.

APPLICATIONS

Vps36 (yN-14) is recommended for detection of Vps36 of *Saccharomyces cerevisiae* origin by Western Blotting (starting dilution 1:200, dilution range 1:100-1:1000) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

RECOMMENDED SECONDARY REAGENTS

To ensure optimal results, the following support (secondary) reagents are recommended: 1) Western Blotting: use donkey anti-goat IgG-HRP: sc-2020 (dilution range: 1:2000-1:100,000) or Cruz Marker™ compatible donkey anti-goat IgG-HRP: sc-2033 (dilution range: 1:2000-1:5000), Cruz Marker™ Molecular Weight Standards: sc-2035, TBS Blotto A Blocking Reagent: sc-2333 and Western Blotting Luminol Reagent: sc-2048.

PROTOCOLS

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

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