SANTA CRUZ BIOTECHNOLOGY, INC.

GGA1 (D-8): sc-390420



BACKGROUND

The GGA family of proteins (Golgi-localized, y-adaptin ear-containing, ARFbinding proteins) are ubiquitous coat proteins that facilitate the trafficking of soluble proteins from the trans-Golgi network (TGN) to endosomes/lysosomes by means of interactions with TGN-sorting receptors, ARF (ADP-ribosylation factor), and clathrin. Members of the GGA family, GGA1, GGA2 (also known as VEAR) and GGA3, are multidomain proteins that bind mannose 6-phosphate receptors (MPRs). GGAs have modular structures with an N-terminal VHS (VPS-27, Hrs, and STAM) domain followed by a GAT (GGA and TOM1) domain, a connecting hinge segment, and a C-terminal GAE (y-adaptin ear) domain. The amino-terminal VHS domains of GGAs form complexes with the cytoplasmic domains of sorting receptors by recognizing acidic-cluster di-leucine (ACLL) sequences. GGA1 and GGA2 do not associate with each other, but they do colocalize on perinuclear membranes. The cytosolic domain of memapsin 2, but not that of memapsin 1, binds the VHS domains of GGA1 and GGA2. The human GGA1 gene maps to chromosome 22q13.1 and encodes a protein that shares 45% sequence identity with GGA2 and GGA3.

REFERENCES

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- Shiba, T., Takatsu, H., Nogi, T., Matsugaki, N., Kawasaki, M., Igarashi, N., Suzuki, M., Kato, R., Earnest, T., Nakayama, K. and Wakatsuki, S. 2002. Structural basis for recognition of acidic-cluster dileucine sequence by GGA1. Nature 415: 937-941.
- Doray, B., Ghosh, P., Griffith, J., Geuze, H.J. and Kornfeld, S. 2002. Cooperation of GGAs and AP-1 in packaging MPRs at the *trans*-Golgi network. Science 297: 1700-1703.
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CHROMOSOMAL LOCATION

Genetic locus: GGA1 (human) mapping to 22q13.1; Gga1 (mouse) mapping to 15 E1.

SOURCE

GGA1 (D-8) is a mouse monoclonal antibody specific for an epitope mapping between amino acids 567-603 near the C-terminus of GGA1 of human origin.

PRODUCT

Each vial contains 200 μg IgG_3 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-390420 P, (100 μ g peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% stabilizer protein).

APPLICATIONS

GGA1 (D-8) is recommended for detection of GGA1 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).

GGA1 (D-8) is also recommended for detection of GGA1 in additional species, including equine, canine, bovine and avian.

Suitable for use as control antibody for GGA1 siRNA (h): sc-41167, GGA1 siRNA (m): sc-41168, GGA1 shRNA Plasmid (h): sc-41167-SH, GGA1 shRNA Plasmid (m): sc-41168-SH, GGA1 shRNA (h) Lentiviral Particles: sc-41167-V and GGA1 shRNA (m) Lentiviral Particles: sc-41168-V.

Molecular Weight of GGA1: 85 kDa.

Positive Controls: GGA1 (h2): 293T Lysate: sc-128702.

RECOMMENDED SUPPORT REAGENTS

To ensure optimal results, the following support reagents are recommended: 1) Western Blotting: use m-IgGκ BP-HRP: sc-516102 or m-IgGκ 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 A/G PLUS-Agarose: sc-2003 (0.5 ml agarose/2.0 ml). 3) Immunofluorescence: use m-IgGκ BP-FITC: sc-516140 or m-IgGκ 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



GGA1 (D-8): sc-390420. Western blot analysis of GGA1 expression in non-transfected: sc-117752 (A) and human GGA1 transfected: sc-128702 (B) whole cell lysates.

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.