

ACTG2 Rabbit pAb

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Product Information

Application WB, IHC-P, IHC-F, IF, E

Primary Accession P63267

Reactivity Rat, Pig, Mouse, Chicken

Host Rabbit
Clonality Polyclonal
Calculated MW 41877
Physical State Liquid

Immunogen KLH conjugated synthetic peptide derived from human ACTG2/Gamma 2 actin

Epitope Specificity 21-120/376

Isotype IgG

Post-translational

modifications

Purity affinity purified by Protein A

Buffer 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol.

SUBCELLULAR LOCATION Cytoplasm, cytoskeleton. Belongs to the actin family.

SUBUNITPolymerization of globular actin (G-actin) leads to a structural filament (F-actin) in the form of a two-stranded helix. Each actin can bind to 4 others.

Oxidation of Met-45 and Met-48 by MICALs (MICAL1, MICAL2 or MICAL3) to form methionine sulfoxide promotes actin filament depolymerization. MICAL1 and MICAL2 produce the (R)-S-oxide form. The (R)-S-oxide form is reverted by MSRB1 and MSRB2, which promote actin repolymerization. Monomethylation at Lys-85 (K84me1) regulates actin-myosin interaction and

actomyosin-dependent processes. Demethylation by ALKBH4 is required for maintaining actomyosin dynamics supporting normal cleavage furrow

ingression during cytokinesis and cell migration.

Important Note This product as supplied is intended for research use only, not for use in

human, therapeutic or diagnostic applications.

Background Descriptions Actins are highly conserved proteins that are involved in various types of cell

motility and in the maintenance of the cytoskeleton. Three types of actins, alpha, beta and gamma, have been identified in vertebrates. Alpha actins are found in muscle tissues and are a major constituent of the contractile apparatus. The beta and gamma actins co-exist in most cell types as components of the cytoskeleton and as mediators of internal cell motility. This gene encodes actin gamma 2; a smooth muscle actin found in enteric tissues. Alternative splicing results in multiple transcript variants encoding distinct isoforms. Based on similarity to peptide cleavage of related actins, the

mature protein of this gene is formed by removal of two N-terminal

peptides.[provided by RefSeq, Dec 2010]

Additional Information

Gene ID 72

Other Names Actin, gamma-enteric smooth muscle, 3.6.4.-, Alpha-actin-3, Gamma-2-actin,

Smooth muscle gamma-actin, Actin, gamma-enteric smooth muscle,

intermediate form, ACTG2, ACTA3, ACTL3, ACTSG

Dilution WB=1:500-2000,IHC-P=1:100-500,IHC-F=1:100-500,ICC/IF=1:100-500,IF=1:100-

500.ELISA=1:5000-10000

Storage Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. When

reconstituted in sterile pH 7.4 0.01M PBS or diluent of antibody the antibody

is stable for at least two weeks at 2-4 °C.

Protein Information

Name ACTG2

Synonyms ACTA3, ACTL3, ACTSG

Function Actins are highly conserved proteins that are involved in various types of cell

motility and are ubiquitously expressed in all eukaryotic cells.

Cellular Location Cytoplasm, cytoskeleton.

Tissue Location In the intestine, abundantly expressed in smooth muscle cells of muscularis

mucosa and muscularis propria. Also detected in intestinal vascular smooth

muscle cells

Background

Actins are highly conserved proteins that are involved in various types of cell motility and in the maintenance of the cytoskeleton. Three types of actins, alpha, beta and gamma, have been identified in vertebrates. Alpha actins are found in muscle tissues and are a major constituent of the contractile apparatus. The beta and gamma actins co-exist in most cell types as components of the cytoskeleton and as mediators of internal cell motility. This gene encodes actin gamma 2; a smooth muscle actin found in enteric tissues. Alternative splicing results in multiple transcript variants encoding distinct isoforms. Based on similarity to peptide cleavage of related actins, the mature protein of this gene is formed by removal of two N-terminal peptides.[provided by RefSeq, Dec 2010]

References

Miwa T., et al. Nucleic Acids Res. 18:4263-4263(1990). Miwa T., et al. Mol. Cell. Biol. 11:3296-3306(1991).

Ota T., et al. Nat. Genet. 36:40-45(2004).

Ebert L., et al. Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.

Hillier L.W., et al. Nature 434:724-731(2005).

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