

## **Anti-ULA1** antibody



**Description** Unconjugated Rabbit polyclonal to ULA1

Model STJ191045

**Host** Rabbit

**Reactivity** Human, Mouse, Rat

**Applications** ELISA, WB

Immunogen Synthesized peptide derived from human ULA1 protein.

**Immunogen Region** 390-470aa

**Gene ID** <u>8883</u>

Gene Symbol NAE1

**Dilution range** WB 1:500-2000 ELISA 1:5000-20000

**Specificity** ULA1 Polyclonal Antibody detects endogenous levels of protein.

**Tissue Specificity** Ubiquitous in fetal tissues. Expressed throughout the adult brain.

**Purification** ULA1 antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

**Note** For Research Use Only (RUO).

Protein Name NEDD8-activating enzyme E1 regulatory subunit Amyloid beta precursor

protein-binding protein 1, 59 kDa APP-BP1 Amyloid protein-binding protein

1 Proto-oncogene protein 1

Molecular Weight 58 kDa

**Clonality** Polyclonal

Conjugation Unconjugated

**Isotype IgG** 

Liquid form in PBS containing 50% glycerol, and 0.02% sodium azide. **Formulation** 

1 mg/ml Concentration

Store at -20°C, and avoid repeat freeze-thaw cycles. **Storage Instruction** 

**Database Links** HGNC:621OMIM:603385

NEDD8-activating enzyme E1 regulatory subunit Amyloid beta precursor **Alternative Names** 

protein-binding protein 1, 59 kDa APP-BP1 Amyloid protein-binding protein

1 Proto-oncogene protein 1

**Function** Regulatory subunit of the dimeric UBA3-NAE1 E1 enzyme. E1 activates

> NEDD8 by first adenylating its C-terminal glycine residue with ATP, thereafter linking this residue to the side chain of the catalytic cysteine, yielding a NEDD8-UBA3 thioester and free AMP. E1 finally transfers NEDD8 to the catalytic cysteine of UBE2M. Necessary for cell cycle progression through the S-M checkpoint. Overexpression of NAE1 causes

apoptosis through deregulation of NEDD8 conjugation.

Cell membrane. Colocalizes with APP in lipid rafts. **Cellular Localization** 

Ubiquitinated by TRIP12, leading to its degradation by the proteasome. Post-translational

**Modifications** 

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