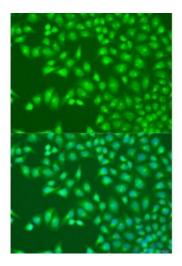


## **Anti-UBA3** Antibody





**Description** 

The modification of proteins with ubiquitin is an important cellular mechanism for targeting abnormal or short-lived proteins for degradation. Ubiquitination involves at least three classes of enzymes: ubiquitinactivating enzymes, or E1s, ubiquitin-conjugating enzymes, or E2s, and ubiquitin-protein ligases, or E3s. This gene encodes a member of the E1 ubiquitin-activating enzyme family. The encoded enzyme associates with AppBp1, an amyloid beta precursor protein binding protein, to form a heterodimer, and then the enzyme complex activates NEDD8, a ubiquitin-like protein, which regulates cell division, signaling and embryogenesis. Multiple alternatively spliced transcript variants encoding distinct isoforms have been found for this gene.

Model STJ117313

**Host** Rabbit

**Reactivity** Human

**Applications** IF

Immunogen Recombinant fusion protein containing a sequence corresponding to amino

acids 234-463 of human UBA3 (NP\_003959.3).

**Gene ID** 9039

Gene Symbol <u>UBA3</u>

**Dilution range** IF 1:50 - 1:200

Tissue Specificity Ubiquitously expressed

**Purification** Affinity purification

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

**Protein Name** NEDD8-activating enzyme E1 catalytic subunit

Molecular Weight 51.852 kDa

**Clonality** Polyclonal

**Conjugation** Unconjugated

**Isotype** IgG

**Formulation** PBS with 0.02% sodium azide, 50% glycerol, pH7.3.

**Storage Instruction** Store at -20C. Avoid freeze / thaw cycles.

Database Links HGNC:12470OMIM:603172Reactome:R-HSA-5607761

Alternative Names NEDD8-activating enzyme E1 catalytic subunit

**Function** Catalytic 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, Down-regulates steroid receptor

activity, Necessary for cell cycle progression,

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