

## **Anti-APBB1** antibody



**Description** Unconjugated Rabbit polyclonal to APBB1

Model STJ190544

**Host** Rabbit

**Reactivity** Human, Mouse, Rat

**Applications** ELISA, WB

Immunogen Synthesized peptide derived from human APBB1 protein.

**Immunogen Region** 400-480aa

**Gene ID** <u>322</u>

Gene Symbol APBB1

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

**Specificity** APBB1 Polyclonal Antibody detects endogenous levels of protein.

Tissue Specificity Highly expressed in brain; strongly reduced in post-mortem elderly subjects

with Alzheimer disease.

**Purification** APBB1 antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

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

**Protein Name** Amyloid-beta A4 precursor protein-binding family B member 1 Protein Fe65

Molecular Weight 78 kDa

**Clonality** Polyclonal

**Conjugation** Unconjugated

**Isotype** IgG

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

**Concentration** 1 mg/ml

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

Database Links <u>HGNC:5810MIM:602709</u>

Alternative Names Amyloid-beta A4 precursor protein-binding family B member 1 Protein Fe65

**Function** Transcription coregulator that can have both coactivator and corepressor

functions. Adapter protein that forms a transcriptionally active complex with the gamma-secretase-derived amyloid precursor protein (APP) intracellular domain. Plays a central role in the response to DNA damage by translocating to the nucleus and inducing apoptosis. May act by specifically recognizing and binding histone H2AX phosphorylated on 'Tyr-142' (H2AXY142ph) at double-strand breaks (DSBs), recruiting other pro-apoptosis factors such as MAPK8/JNK1. Required for histone H4 acetylation at double-strand breaks (DSBs). Its ability to specifically bind modified histones and chromatin modifying enzymes such as KAT5/TIP60, probably explains its transcription activation activity. Function in association with TSHZ3, SET and HDAC factors as a transcriptional repressor, that inhibits the expression of CASP4. Associates with chromatin in a region surrounding the CASP4 transcriptional

start site(s).

Cellular Localization Cell membrane. Cytoplasm. Nucleus. Cell projection, growth cone Nucleus

speckle. Colocalizes with TSHZ3 in axonal growth cone . In normal conditions, it mainly localizes to the cytoplasm, while a small fraction is tethered to the cell membrane via its interaction with APP. Following exposure to DNA damaging agents, it is released from cell membrane and translocates to the nucleus. Nuclear translocation is under the regulation of APP. Colocalizes with TSHZ3 in the nucleus. Colocalizes with NEK6 at the nuclear speckles. Phosphorylation at Ser-610 by SGK1 promotes its

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localization to the nucleus.

Post-translationalPhosphorylation at Ser-610 by SGK1 promotes its localization to the nucleus .ModificationsPhosphorylated following nuclear translocation. Phosphorylation at Tyr-547by ABL1 enhances transcriptional activation activity and reduces the affinity

by ABL1 enhances transcriptional activation activity and reduces the affini

for RASD1/DEXRAS1.

St John's Laboratory Ltd

**F** +44 (0)207 681 2580

T+44 (0)208 223 3081

W http://www.stjohnslabs.com/ E info@stjohnslabs.com