

RENT1 Antibody (Center)

Purified Rabbit Polyclonal Antibody (Pab) Catalog # AP1905c

Product Information

Application WB, E **Primary Accession** Q92900 **Other Accession** Q9EPU0 Reactivity Human **Predicted** Mouse Host Rabbit Clonality Polyclonal Isotype Rabbit IgG **Clone Names** RB8354 **Calculated MW** 124345 **Antigen Region** 583-612

Additional Information

Gene ID 5976

Other Names Regulator of nonsense transcripts 1, 364-, ATP-dependent helicase RENT1,

Nonsense mRNA reducing factor 1, NORF1, Up-frameshift suppressor 1

homolog, hUpf1, UPF1, KIAA0221, RENT1

Target/Specificity This RENT1 antibody is generated from rabbits immunized with a KLH

conjugated synthetic peptide between 583-612 amino acids from the Central

region of human RENT1.

Dilution WB~~1:1000 E~~Use at an assay dependent concentration.

Format Purified polyclonal antibody supplied in PBS with 0.05% (V/V) Proclin 300. This

antibody is purified through a protein A column, followed by peptide affinity

purification.

Storage Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store

at -20°C in small aliquots to prevent freeze-thaw cycles.

Precautions RENT1 Antibody (Center) is for research use only and not for use in diagnostic

or therapeutic procedures.

Protein Information

Name UPF1 (HGNC:9962)

Function RNA-dependent helicase required for nonsense-mediated decay (NMD) of

aberrant mRNAs containing premature stop codons and modulates the expression level of normal mRNAs (PubMed: 11163187, PubMed: 16086026, PubMed: 18172165, PubMed: 21145460, PubMed: 21419344, PubMed:24726324). Is recruited to mRNAs upon translation termination and undergoes a cycle of phosphorylation and dephosphorylation; its phosphorylation appears to be a key step in NMD (PubMed:11544179, PubMed: <u>25220460</u>). Recruited by release factors to stalled ribosomes together with the SMG1C protein kinase complex to form the transient SURF (SMG1-UPF1-eRF1-eRF3) complex (PubMed: 19417104). In EJC-dependent NMD, the SURF complex associates with the exon junction complex (EJC) (located 50-55 or more nucleotides downstream from the termination codon) through UPF2 and allows the formation of an UPF1-UPF2-UPF3 surveillance complex which is believed to activate NMD (PubMed:21419344). Phosphorylated UPF1 is recognized by EST1B/SMG5, SMG6 and SMG7 which are thought to provide a link to the mRNA degradation machinery involving exonucleolytic and endonucleolytic pathways, and to serve as adapters to protein phosphatase 2A (PP2A), thereby triggering UPF1 dephosphorylation and allowing the recycling of NMD factors (PubMed: 12554878). UPF1 can also activate NMD without UPF2 or UPF3, and in the absence of the NMD-enhancing downstream EJC indicative for alternative NMD pathways (PubMed: 18447585). Plays a role in replication-dependent histone mRNA degradation at the end of phase S; the function is independent of UPF2 (PubMed:16086026, PubMed:18172165). For the recognition of premature termination codons (PTC) and initiation of NMD a competitive interaction between UPF1 and PABPC1 with the ribosome-bound release factors is proposed (PubMed: 18447585, PubMed: 25220460). The ATPase activity of UPF1 is required for disassembly of mRNPs undergoing NMD (PubMed:21145460). Together with UPF2 and dependent on TDRD6, mediates the degradation of mRNA harboring long 3'UTR by inducing the NMD machinery (By similarity). Also capable of unwinding double-stranded DNA and translocating on single-stranded DNA (PubMed:30218034).

Cellular Location

Cytoplasm. Cytoplasm, P-body. Nucleus. Cytoplasm, perinuclear region {ECO:0000250 | UniProtKB:Q9EPU0}. Note=Hyperphosphorylated form is targeted to the P-body, while unphosphorylated protein is distributed throughout the cytoplasm. Localized in the chromatoid bodies of round spermatids (By similarity). {ECO:0000250 | UniProtKB:Q9EPU0}

Tissue Location

Ubiquitous.

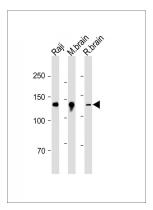
Background

RENT1 is part of a post-splicing multiprotein complex involved in both mRNA nuclear export and mRNA surveillance. mRNA surveillance detects exported mRNAs with truncated open reading frames and initiates nonsense-mediated mRNA decay (NMD). When translation ends upstream from the last exon-exon junction, this triggers NMD to degrade mRNAs containing premature stop codons. This protein is located only in the cytoplasm. When translation ends, it interacts with the protein that is a functional homolog of yeast Upf2p to trigger mRNA decapping.

References

Ohnishi, T., et al., Mol. Cell 12(5):1187-1200 (2003). Lykke-Andersen, J., Mol. Cell. Biol. 22(23):8114-8121 (2002). Carastro, L.M., et al., Nucleic Acids Res. 30(10):2232-2243 (2002). Mendell, J.T., et al., Science 298(5592):419-422 (2002). Serin, G., et al., Mol. Cell. Biol. 21(1):209-223 (2001).

Images



All lanes: Anti-RENT1 Antibody (Center) at 1:1000 dilution Lane 1: Raji whole cell lysate Lane 2: Mouse brain lysate Lane 3: Rat brain lysate Lysates/proteins at 20 µg per lane. Secondary: Goat Anti-Rabbit IgG, (H+L), Peroxidase conjugated (ASP1615) at 1/15000 dilution. Observed band size: 130 KDa Blocking/Dilution buffer: 5% NFDM/TBST.

Citations

• The intronic GABRG2 mutation, IVS6+2T-G, associated with childhood absence epilepsy altered subunit mRNA intron splicing, activated nonsense-mediated decay, and produced a stable truncated y2 subunit.

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