

**CLN3 Antibody (Center)**  
**Purified Rabbit Polyclonal Antibody (Pab)**  
**Catalog # AP20845c**

**Specification**

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**CLN3 Antibody (Center) - Product Information**

Application	WB,E
Primary Accession	<a href="#">Q13286</a>
Other Accession	<a href="#">Q60HHQ</a>
Reactivity	Human, Mouse
Predicted	Monkey
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Antigen Region	250-284

**CLN3 Antibody (Center) - Additional Information**

**Gene ID** 1201

**Other Names**

Battenin, Batten disease protein, Protein CLN3, CLN3, BTS

**Target/Specificity**

This CLN3 antibody is generated from a rabbit immunized with a KLH conjugated synthetic peptide between 250-284 amino acids from the Central region of human CLN3.

**Dilution**

WB~~1:500

**Format**

Purified polyclonal antibody supplied in PBS with 0.09% (W/V) sodium azide. 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**

CLN3 Antibody (Center) is for research use only and not for use in diagnostic or therapeutic procedures.

**CLN3 Antibody (Center) - Protein Information**

**Name** CLN3 ([HGNC:2074](#))

**Synonyms** BTS

**Function** Mediates microtubule-dependent, anterograde transport connecting the Golgi network, endosomes, autophagosomes, lysosomes and plasma membrane, and participates in several cellular processes such as regulation of lysosomal pH, lysosome protein degradation, receptor-mediated endocytosis, autophagy, transport of proteins and lipids from the TGN, apoptosis and synaptic transmission (PubMed:[10924275](#), PubMed:[18817525](#), PubMed:[18317235](#), PubMed:[22261744](#), PubMed:[15471887](#), PubMed:[20850431](#)). Facilitates the proteins transport from trans-Golgi network (TGN)-to other membrane compartments such as transport of microdomain-associated proteins to the plasma membrane, IGF2R transport to the lysosome where it regulates the CTSD release leading to regulation of CTSD maturation and thereby APP intracellular processing (PubMed:[10924275](#), PubMed:[18817525](#)). Moreover regulates CTSD activity in response to osmotic stress (PubMed:[23840424](#), PubMed:[28390177](#)). Also binds galactosylceramide and transports it from the trans Golgi to the rafts, which may have immediate and downstream effects on cell survival by modulating ceramide synthesis (PubMed:[18317235](#)). At the plasma membrane, regulates actin-dependent events including filopodia formation, cell migration, and pinocytosis through ARF1-CDC42 pathway and also the cytoskeleton organization through interaction with MYH10 and fodrin leading to the regulation of the plasma membrane association of Na<sup>+</sup>, K<sup>+</sup> ATPase complex (PubMed:[20850431](#)). Regulates synaptic transmission in the amygdala, hippocampus, and cerebellum through regulation of synaptic vesicles density and their proximity to active zones leading to modulation of short-term plasticity and age-dependent anxious behavior, learning and memory (By similarity). Regulates autophagic vacuoles (AVs) maturation by modulating the trafficking between endocytic and autophagolysosomal/lysosomal compartments, which involves vesicle fusion leading to regulation of degradation process (By similarity). Participates also in cellular homeostasis of compounds such as, water, ions, amino acids, proteins and lipids in several tissue namely in brain and kidney through regulation of their transport and synthesis (PubMed:[17482562](#)).

#### Cellular Location

Lysosome membrane; Multi-pass membrane protein. Late endosome. Lysosome. Golgi apparatus. Golgi apparatus membrane. Golgi apparatus, Golgi stack. Golgi apparatus, trans-Golgi network. Cell membrane Recycling endosome. Membrane raft. Membrane, caveola. Early endosome membrane. Synapse, synaptosome {ECO:0000250|UniProtKB:Q61124}. Late endosome membrane {ECO:0000250|UniProtKB:Q61124}. Cytoplasmic vesicle, autophagosome {ECO:0000250|UniProtKB:Q61124}. Note=CLN3 is not present in late endosomes/lysosomes in fibroblasts and neurons (PubMed:15240864) Trafficks from cell membrane to Golgi via endosomes (PubMed:15240864) Osmotic stress changes the subcellular localization of CLN3 (PubMed:23840424). Trafficks to intracellular compartments via the plasma membranet through AP3M1-dependent mechanisms (PubMed:14644441) Excluded from the synaptic vesicles (By similarity) {ECO:0000250|UniProtKB:Q61124, ECO:0000269|PubMed:14644441, ECO:0000269|PubMed:15240864, ECO:0000269|PubMed:23840424}

#### Tissue Location

Expressed in the cortical brain, pancreas, spleen, and testis with weaker expression in the peripheral nerve (at protein level). Highly expressed in gray matter (at protein level)

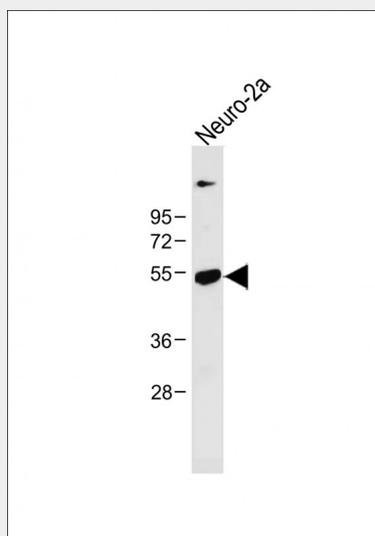
### CLN3 Antibody (Center) - Protocols

Provided below are standard protocols that you may find useful for product applications.

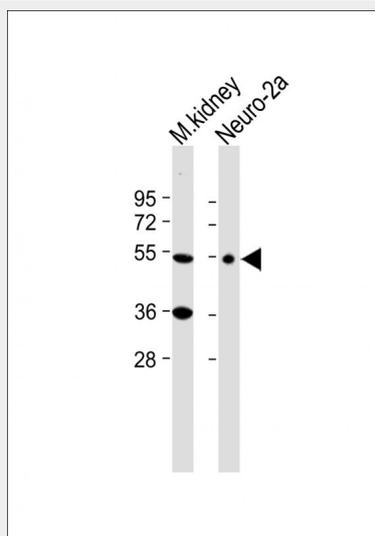
- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)

- [Cell Culture](#)

## CLN3 Antibody (Center) - Images



Anti-CLN3 Antibody (Center) at 1:1000 dilution + Neuro-2a whole cell lysate Lysates/proteins at 20  $\mu$ g per lane. Secondary Goat Anti-Rabbit IgG, (H+L), Peroxidase conjugated at 1/10000 dilution. Predicted band size : 48 kDa Blocking/Dilution buffer: 5% NFDm/TBST.



All lanes : Anti-CLN3 Antibody (Center) at 1:500 dilution Lane 1: Mouse kidney lysate Lane 2: Neuro-2a whole cell lysate Lysates/proteins at 20  $\mu$ g per lane. Secondary Goat Anti-Rabbit IgG, (H+L), Peroxidase conjugated at 1/10000 dilution. Predicted band size : 48 kDa Blocking/Dilution buffer: 5% NFDm/TBST.

## CLN3 Antibody (Center) - Background

Involved in microtubule-dependent, anterograde transport of late endosomes and lysosomes.

## CLN3 Antibody (Center) - References

- Lerner T.J., et al. Cell 82:949-957(1995).  
 Mitchison H.M., et al. Genomics 40:346-350(1997).  
 LaFauci G., et al. Submitted (JUL-1997) to the EMBL/GenBank/DDBJ databases.

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