

**AFG3L2 Polyclonal Antibody**  
Catalog # AP74280**Specification****AFG3L2 Polyclonal Antibody - Product Information**

Application	WB
Primary Accession	<a href="#">O9Y4W6</a>
Reactivity	Human
Host	Rabbit
Clonality	Polyclonal

**AFG3L2 Polyclonal Antibody - Additional Information**

Gene ID 10939

**Other Names**

AFG3-like protein 2 (EC 3.4.24.-) (Paraplegin-like protein)

**Dilution**

WB~~WB 1:500-2000, ELISA 1:10000-20000

**Format**

Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.09% (W/V) sodium azide.

**Storage Conditions**

-20°C

**AFG3L2 Polyclonal Antibody - Protein Information****Name** AFG3L2 {ECO:0000303|PubMed:10395799, ECO:0000312|HGNC:HGNC:315}**Function**

Catalytic component of the m-AAA protease, a protease that plays a key role in proteostasis of inner mitochondrial membrane proteins, and which is essential for axonal and neuron development (PubMed: [19748354](http://www.uniprot.org/citations/19748354) target="\_blank">19748354</a>, PubMed: [28396416](http://www.uniprot.org/citations/28396416) target="\_blank">28396416</a>, PubMed: [29932645](http://www.uniprot.org/citations/29932645) target="\_blank">29932645</a>, PubMed: [30683687](http://www.uniprot.org/citations/30683687) target="\_blank">30683687</a>, PubMed: [31327635](http://www.uniprot.org/citations/31327635) target="\_blank">31327635</a>, PubMed: [37917749](http://www.uniprot.org/citations/37917749) target="\_blank">37917749</a>, PubMed: [38157846](http://www.uniprot.org/citations/38157846) target="\_blank">38157846</a>). AFG3L2 possesses both ATPase and protease activities: the ATPase activity is required to unfold substrates, threading them into the internal proteolytic cavity for hydrolysis into small peptide fragments (PubMed: [19748354](http://www.uniprot.org/citations/19748354) target="\_blank">19748354</a>, PubMed: [31327635](http://www.uniprot.org/citations/31327635) target="\_blank">31327635</a>). The m-AAA protease carries out quality control in the inner membrane of the mitochondria by mediating degradation of mistranslated or misfolded polypeptides (PubMed:

<http://www.uniprot.org/citations/26504172> target="\_blank">26504172</a>, PubMed:<a href="http://www.uniprot.org/citations/30683687" target="\_blank">30683687</a>, PubMed:<a href="http://www.uniprot.org/citations/34718584" target="\_blank">34718584</a>). The m-AAA protease complex also promotes the processing and maturation of mitochondrial proteins, such as MRPL32/bL32m, PINK1 and SP7 (PubMed:<a href="http://www.uniprot.org/citations/22354088" target="\_blank">22354088</a>, PubMed:<a href="http://www.uniprot.org/citations/29932645" target="\_blank">29932645</a>, PubMed:<a href="http://www.uniprot.org/citations/30252181" target="\_blank">30252181</a>). Mediates protein maturation of the mitochondrial ribosomal subunit MRPL32/bL32m by catalyzing the cleavage of the presequence of MRPL32/bL32m prior to assembly into the mitochondrial ribosome (PubMed:<a href="http://www.uniprot.org/citations/29932645" target="\_blank">29932645</a>). Required for SPG7 maturation into its active mature form after SPG7 cleavage by mitochondrial-processing peptidase (MPP) (PubMed:<a href="http://www.uniprot.org/citations/30252181" target="\_blank">30252181</a>). Required for the maturation of PINK1 into its 52kDa mature form after its cleavage by mitochondrial- processing peptidase (MPP) (PubMed:<a href="http://www.uniprot.org/citations/22354088" target="\_blank">22354088</a>). Acts as a regulator of calcium in neurons by mediating degradation of SMDT1/EMRE before its assembly with the uniporter complex, limiting the availability of SMDT1/EMRE for MCU assembly and promoting efficient assembly of gatekeeper subunits with MCU (PubMed:<a href="http://www.uniprot.org/citations/27642048" target="\_blank">27642048</a>, PubMed:<a href="http://www.uniprot.org/citations/28396416" target="\_blank">28396416</a>). Promotes the proteolytic degradation of GHITM upon hyperpolarization of mitochondria: progressive GHITM degradation leads to respiratory complex I degradation and broad reshaping of the mitochondrial proteome by AFG3L2 (PubMed:<a href="http://www.uniprot.org/citations/35912435" target="\_blank">35912435</a>). Also acts as a regulator of mitochondrial glutathione homeostasis by mediating cleavage and degradation of SLC25A39 (PubMed:<a href="http://www.uniprot.org/citations/37917749" target="\_blank">37917749</a>, PubMed:<a href="http://www.uniprot.org/citations/38157846" target="\_blank">38157846</a>). SLC25A39 cleavage is prevented when SLC25A39 binds iron-sulfur (PubMed:<a href="http://www.uniprot.org/citations/37917749" target="\_blank">37917749</a>, PubMed:<a href="http://www.uniprot.org/citations/38157846" target="\_blank">38157846</a>). Involved in the regulation of OMA1-dependent processing of OPA1 (PubMed:<a href="http://www.uniprot.org/citations/17615298" target="\_blank">17615298</a>, PubMed:<a href="http://www.uniprot.org/citations/29545505" target="\_blank">29545505</a>, PubMed:<a href="http://www.uniprot.org/citations/30252181" target="\_blank">30252181</a>, PubMed:<a href="http://www.uniprot.org/citations/30683687" target="\_blank">30683687</a>, PubMed:<a href="http://www.uniprot.org/citations/32600459" target="\_blank">32600459</a>). May act by mediating processing of OMA1 precursor, participating in OMA1 maturation (PubMed:<a href="http://www.uniprot.org/citations/29545505" target="\_blank">29545505</a>).

### Cellular Location

Mitochondrion inner membrane; Multi-pass membrane protein

### Tissue Location

Ubiquitous. Highly expressed in the cerebellar Purkinje cells.

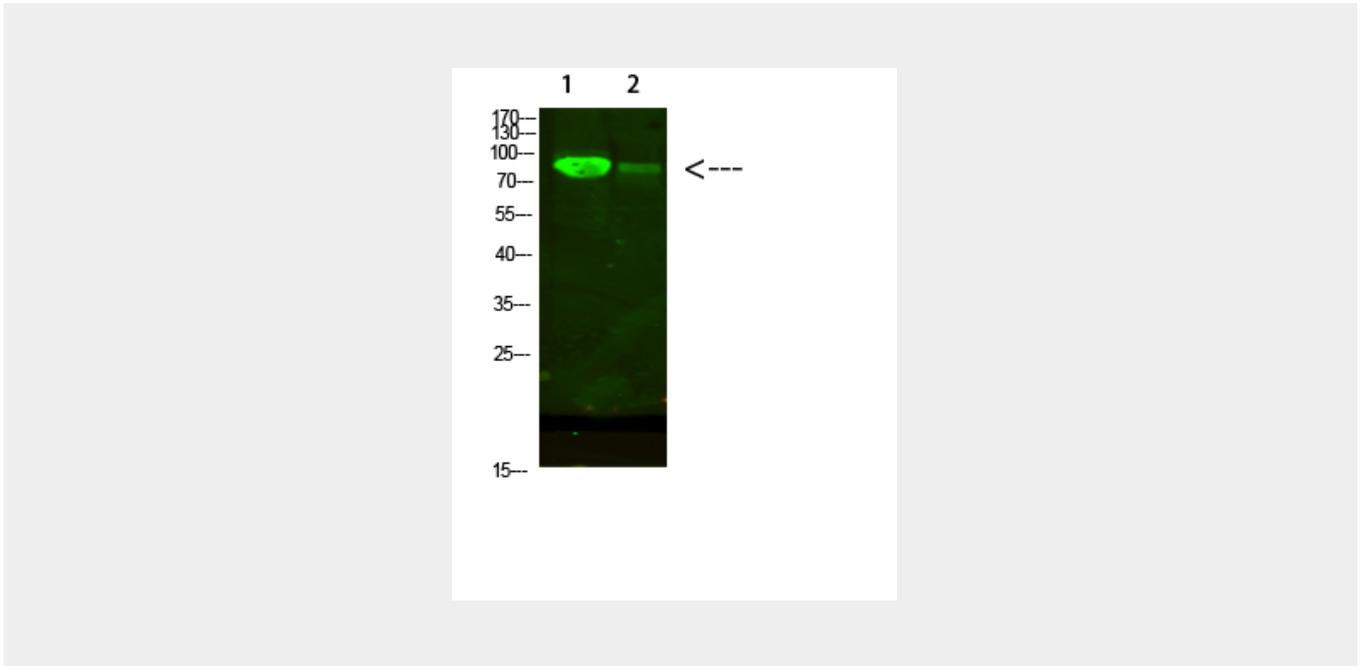
## AFG3L2 Polyclonal Antibody - Protocols

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](#)

#### **AFG3L2 Polyclonal Antibody - Images**



#### **AFG3L2 Polyclonal Antibody - Background**

ATP-dependent protease which is essential for axonal and neuron development. In neurons, mediates degradation of SMDT1/EMRE before its assembly with the uniporter complex, limiting the availability of SMDT1/EMRE for MCU assembly and promoting efficient assembly of gatekeeper subunits with MCU (PubMed:27642048). Required for the maturation of paraplegin (SPG7) after its cleavage by mitochondrial-processing peptidase (MPP), converting it into a proteolytically active mature form (By similarity).