

**Product Name: LRRK2 (16M6) Rabbit Monoclonal Antibody**  
**Catalog #: AMRe13445**



## Summary

<b>Production Name</b>	LRRK2 (16M6) Rabbit Monoclonal Antibody
<b>Description</b>	Rabbit Monoclonal Antibody
<b>Host</b>	Rabbit
<b>Application</b>	WB,ELISA
<b>Reactivity</b>	Human,Mouse,Rat

## Performance

<b>Conjugation</b>	Unconjugated
<b>Modification</b>	Unmodified
<b>Isotype</b>	IgG
<b>Clonality</b>	Monoclonal
<b>Form</b>	Liquid
<b>Storage</b>	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.
<b>Buffer</b>	Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.02% New type preservative N and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid freeze / thaw cycle.
<b>Purification</b>	Affinity purification

## Immunogen

<b>Gene Name</b>	LRRK2
<b>Alternative Names</b>	Leucine-rich repeat serine/threonine-protein kinase 2; Dardarin; PARK8; ROCO2; RIPK7; LRRK2
<b>Gene ID</b>	120892.0
<b>SwissProt ID</b>	Q5S007.

## Application

<b>Dilution Ratio</b>	WB 1:500-1:2000
<b>Molecular Weight</b>	286kDa

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## Background

LRRK2 positively regulates autophagy through a calcium-dependent activation of the CaMKK/AMPK signaling pathway. The process involves activation of nicotinic acid adenine dinucleotide phosphate (NAADP) receptors, increase in lysosomal pH, and calcium release from lysosomes. Together with RAB29, plays a role in the retrograde trafficking pathway for recycling proteins, such as mannose 6 phosphate receptor (M6PR), between lysosomes and the Golgi apparatus in a retromer-dependent manner. Regulates neuronal process morphology in the intact central nervous system (CNS). Plays a role in synaptic vesicle trafficking. Phosphorylates PRDX3. Has GTPase activity. May play a role in the phosphorylation of proteins central to Parkinson disease. Serine/threonine-protein kinase which phosphorylates a broad range of proteins involved in multiple processes such as neuronal plasticity, autophagy, and vesicle trafficking (PubMed:[20949042](http://www.uniprot.org/citations/20949042), PubMed:[22012985](http://www.uniprot.org/citations/22012985), PubMed:[26824392](http://www.uniprot.org/citations/26824392), PubMed:[29125462](http://www.uniprot.org/citations/29125462), PubMed:[28720718](http://www.uniprot.org/citations/28720718), PubMed:[29127255](http://www.uniprot.org/citations/29127255), PubMed:[30398148](http://www.uniprot.org/citations/30398148), PubMed:[29212815](http://www.uniprot.org/citations/29212815), PubMed:[30635421](http://www.uniprot.org/citations/30635421), PubMed:[21850687](http://www.uniprot.org/citations/21850687), PubMed:[23395371](http://www.uniprot.org/citations/23395371), PubMed:[17114044](http://www.uniprot.org/citations/17114044), PubMed:[24687852](http://www.uniprot.org/citations/24687852), PubMed:[26014385](http://www.uniprot.org/citations/26014385), PubMed:[25201882](http://www.uniprot.org/citations/25201882)). Is a key regulator of RAB GTPases by regulating the GTP/GDP exchange and interaction partners of RABs through phosphorylation (PubMed:[26824392](http://www.uniprot.org/citations/26824392), PubMed:[28720718](http://www.uniprot.org/citations/28720718), PubMed:[29127255](http://www.uniprot.org/citations/29127255), PubMed:[30398148](http://www.uniprot.org/citations/30398148), PubMed:[29212815](http://www.uniprot.org/citations/29212815), PubMed:[29125462](http://www.uniprot.org/citations/29125462), PubMed:[30635421](http://www.uniprot.org/citations/30635421)). Phosphorylates RAB3A, RAB3B, RAB3C, RAB3D, RAB5A, RAB5B, RAB5C, RAB8A, RAB8B, RAB10, RAB12, RAB35, and RAB43 (PubMed:[26824392](http://www.uniprot.org/citations/26824392), PubMed:[26824392](http://www.uniprot.org/citations/26824392), PubMed:[26824392](http://www.uniprot.org/citations/26824392)).



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[28720718](http://www.uniprot.org/citations/28720718), PubMed:[29127255](http://www.uniprot.org/citations/29127255), PubMed:[30398148](http://www.uniprot.org/citations/30398148), PubMed:[29212815](http://www.uniprot.org/citations/29212815), PubMed:[29125462](http://www.uniprot.org/citations/29125462), PubMed:[30635421](http://www.uniprot.org/citations/30635421), PubMed:[23395371](http://www.uniprot.org/citations/23395371)). Regulates the RAB3IP-catalyzed GDP/GTP exchange for RAB8A through the phosphorylation of 'Thr-72' on RAB8A (PubMed:[26824392](http://www.uniprot.org/citations/26824392)). Inhibits the interaction between RAB8A and GDI1 and/or GDI2 by phosphorylating 'Thr- 72' on RAB8A (PubMed:[26824392](http://www.uniprot.org/citations/26824392)). Regulates primary ciliogenesis through phosphorylation of RAB8A and RAB10, which promotes SHH signaling in the brain (PubMed:[29125462](http://www.uniprot.org/citations/29125462), PubMed:[30398148](http://www.uniprot.org/citations/30398148)). Together with RAB29, plays a role in the retrograde trafficking pathway for recycling proteins, such as mannose-6-phosphate receptor (M6PR), between lysosomes and the Golgi apparatus in a retromer-dependent manner (PubMed:[23395371](http://www.uniprot.org/citations/23395371)). Regulates neuronal process morphology in the intact central nervous system (CNS) (PubMed:[17114044](http://www.uniprot.org/citations/17114044)). Plays a role in synaptic vesicle trafficking (PubMed:[24687852](http://www.uniprot.org/citations/24687852)). Plays an important role in recruiting SEC16A to endoplasmic reticulum exit sites (ERES) and in regulating ER to Golgi vesicle-mediated transport and ERES organization (PubMed:[25201882](http://www.uniprot.org/citations/25201882)). Positively regulates autophagy through a calcium-dependent activation of the CaMKK/AMPK signaling pathway (PubMed:[22012985](http://www.uniprot.org/citations/22012985)). The process involves activation of nicotinic acid adenine dinucleotide phosphate (NAADP) receptors, increase in lysosomal pH, and calcium release from lysosomes (PubMed:[22012985](http://www.uniprot.org/citations/22012985)). Phosphorylates PRDX3 (PubMed:[21850687](http://www.uniprot.org/citations/21850687)). By phosphorylating APP on 'Thr-743', which promotes the production and the nuclear translocation of the APP intracellular domain (AICD), regulates dopaminergic neuron apoptosis (PubMed:[28720718](http://www.uniprot.org/citations/28720718)). Independent of its kinase activity, inhibits the proteosomal degradation of MAPT, thus promoting MAPT oligomerization and secretion (PubMed:[26014385](http://www.uniprot.org/citations/26014385)). In addition, has GTPase activity via its Roc domain which regulates LRRK2 kinase activity (PubMed:[18230735](http://www.uniprot.org/citations/18230735), PubMed:[26824392](http://www.uniprot.org/citations/26824392), PubMed:[29125462](http://www.uniprot.org/citations/29125462), PubMed:[29125462](http://www.uniprot.org/citations/29125462), PubMed:[29125462](http://www.uniprot.org/citations/29125462)).

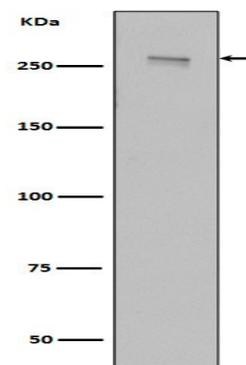
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<http://www.uniprot.org/citations/28720718> </a>, PubMed: <a href="http://www.uniprot.org/citations/29212815" target="\_blank">http://www.uniprot.org/citations/29212815 </a>).

## Research Area

## Image Data



Western blot analysis of LRRK2 in HEK293 cell lysate transfected with 3\*Flag wild type, full length LRRK2.

## Note

For research use only.