

# Insulin degrading enzyme Rabbit mAb

Catalog # AP76553

## Product Information

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<b>Application</b>	WB, FC
<b>Primary Accession</b>	<a href="#">P14735</a>
<b>Reactivity</b>	Rat, Human, Mouse
<b>Host</b>	Rabbit
<b>Clonality</b>	Monoclonal Antibody
<b>Isotype</b>	IgG
<b>Conjugate</b>	Unconjugated
<b>Purification</b>	Affinity Purified
<b>Calculated MW</b>	117968

## Additional Information

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<b>Gene ID</b>	3416
<b>Other Names</b>	IDE
<b>Dilution</b>	WB~~1:1000 FC~~1:10~50
<b>Format</b>	Liquid in 50mM Tris-Glycine(pH 7.4), 0.15M NaCl, 40%Glycerol, 0.01% sodium azide and 0.05% BSA.
<b>Storage</b>	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.

## Protein Information

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<b>Name</b>	IDE {ECO:0000303   PubMed:20364150, ECO:0000312   HGNC:HGNC:5381}
<b>Function</b>	Plays a role in the cellular breakdown of insulin, APP peptides, IAPP peptides, natriuretic peptides, glucagon, bradykinin, kallidin, and other peptides, and thereby plays a role in intercellular peptide signaling (PubMed: <a href="#">10684867</a> , PubMed: <a href="#">17051221</a> , PubMed: <a href="#">17613531</a> , PubMed: <a href="#">18986166</a> , PubMed: <a href="#">19321446</a> , PubMed: <a href="#">21098034</a> , PubMed: <a href="#">2293021</a> , PubMed: <a href="#">23922390</a> , PubMed: <a href="#">24847884</a> , PubMed: <a href="#">26394692</a> , PubMed: <a href="#">26968463</a> , PubMed: <a href="#">29596046</a> ). Substrate binding induces important conformation changes, making it possible to bind and degrade larger substrates, such as insulin (PubMed: <a href="#">23922390</a> , PubMed: <a href="#">26394692</a> , PubMed: <a href="#">29596046</a> ). Contributes to the regulation of peptide hormone signaling cascades and regulation of blood glucose homeostasis via its role in the degradation of insulin, glucagon and IAPP (By similarity). Plays a role in the degradation and clearance of APP-derived amyloidogenic peptides that are secreted by neurons and microglia (Probable) (PubMed: <a href="#">26394692</a> , PubMed: <a href="#">9830016</a> ). Degrades the natriuretic peptides ANP, BNP and CNP,

inactivating their ability to raise intracellular cGMP (PubMed:[21098034](#)). Also degrades an aberrant frameshifted 40-residue form of NPPA (fsNPPA) which is associated with familial atrial fibrillation in heterozygous patients (PubMed:[21098034](#)). Involved in antigen processing. Produces both the N terminus and the C terminus of MAGEA3-derived antigenic peptide (EVDPIGHLY) that is presented to cytotoxic T lymphocytes by MHC class I.

**Cellular Location**

Cytoplasm, cytosol. Cell membrane {ECO:0000250|UniProtKB:P35559}. Secreted Note=Present at the cell surface of neuron cells. The membrane-associated isoform is approximately 5 kDa larger than the known cytosolic isoform

**Tissue Location**

Detected in brain and in cerebrospinal fluid (at protein level).

## Background

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This gene encodes a zinc metallopeptidase that degrades intracellular insulin, and thereby terminates insulin's activity, as well as participating in intercellular peptide signalling by degrading diverse peptides such as glucagon, amylin, bradykinin, and kallidin. The preferential affinity of this enzyme for insulin results in insulin-mediated inhibition of the degradation of other peptides such as beta-amyloid.

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