

**CARD4 Antibody (C-term)**  
**Affinity Purified Rabbit Polyclonal Antibody (Pab)**  
**Catalog # AP8720b****Specification**

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**CARD4 Antibody (C-term) - Product Information**

Application	FC, IHC-P, WB,E
Primary Accession	<a href="#">O9Y239</a>
Reactivity	Human
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Calculated MW	107691
Antigen Region	923-951

**CARD4 Antibody (C-term) - Additional Information****Gene ID** 10392**Other Names**

Nucleotide-binding oligomerization domain-containing protein 1, Caspase recruitment domain-containing protein 4, NOD1, CARD4

**Target/Specificity**

This CARD4 antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 923-951 amino acids from the C-terminal region of human CARD4.

**Dilution**

FC~~1:10~50

IHC-P~~1:50~100

WB~~1:1000

E~~Use at an assay dependent concentration.

**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**

CARD4 Antibody (C-term) is for research use only and not for use in diagnostic or therapeutic procedures.

**CARD4 Antibody (C-term) - Protein Information****Name** NOD1 {ECO:0000303|PubMed:10329646, ECO:0000312|HGNC:HGNC:16390}

**Function** Pattern recognition receptor (PRR) that detects bacterial peptidoglycan fragments and other danger signals and thus participates in both innate and adaptive immune responses (PubMed:[11058605](#), PubMed:[12791997](#), PubMed:[12796777](#), PubMed:[15044951](#), PubMed:[16172124](#), PubMed:[19043560](#), PubMed:[22672233](#), PubMed:[27099311](#)). Specifically recognizes and binds gamma-D-glutamyl-meso-diaminopimelic acid (iE- DAP), a dipeptide present in peptidoglycan of Gram-negative bacteria (PubMed:[12791997](#), PubMed:[12796777](#), PubMed:[12871942](#), PubMed:[16172124](#), PubMed:[16211083](#)). Preferentially binds iE-DAP in tripeptide-containing muropeptides (MurNAc-TriDAP or TriDAP) (PubMed:[16211083](#)). Ligand binding triggers oligomerization that facilitates the binding and subsequent activation of the proximal adapter receptor-interacting RIPK2 (PubMed:[12791997](#), PubMed:[12796777](#), PubMed:[17054981](#)). Following recruitment, RIPK2 undergoes 'Met-1'- (linear) and 'Lys-63'-linked polyubiquitination by E3 ubiquitin-protein ligases XIAP, BIRC2, BIRC3 and the LUBAC complex, becoming a scaffolding protein for downstream effectors, triggering activation of the NF-kappa-B and MAP kinases signaling (PubMed:[10880512](#), PubMed:[12791997](#), PubMed:[19043560](#)). This in turn leads to the transcriptional activation of hundreds of genes involved in immune response (PubMed:[10880512](#), PubMed:[19043560](#)). Also acts as a regulator of antiviral response elicited by dsRNA and the expression of RLR pathway members by targeting IFIH1 and TRAF3 to modulate the formation of IFIH1-MAVS and TRAF3-MAVS complexes leading to increased transcription of type I IFNs (PubMed:[32169843](#)). Also acts as a regulator of autophagy via its interaction with ATG16L1, possibly by recruiting ATG16L1 at the site of bacterial entry (By similarity). Besides recognizing pathogens, also involved in the endoplasmic reticulum stress response: acts by sensing and binding to the cytosolic metabolite sphingosine-1-phosphate generated in response to endoplasmic reticulum stress, initiating an inflammation process that leads to activation of the NF-kappa-B and MAP kinases signaling (PubMed:[27007849](#), PubMed:[33942347](#)). In addition, plays a role in insulin trafficking in beta cells in a cell-autonomous manner (By similarity). Mechanistically, upon recognizing cognate ligands, NOD1 and RIPK2 localize to insulin vesicles where they recruit RAB1A to direct insulin trafficking through the cytoplasm (By similarity).

#### **Cellular Location**

Cell membrane; Lipid-anchor. Apical cell membrane. Basolateral cell membrane. Cytoplasm. Note=Detected in the cytoplasm and at the cell membrane (PubMed:[31649195](#)). Following bacterial infection, localizes to bacterial entry sites in the cell membrane (PubMed:[31649195](#)). Recruited to the basolateral and apical membranes in polarized epithelial cells (PubMed:[19043560](#))

#### **Tissue Location**

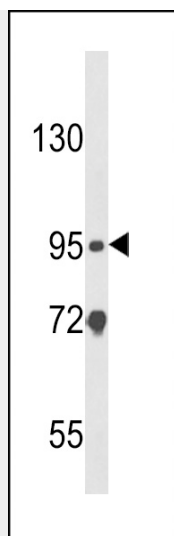
Highly expressed in adult heart, skeletal muscle, pancreas, spleen and ovary (PubMed:[10224040](#)). Also detected in placenta, lung, liver, kidney, thymus, testis, small intestine and colon (PubMed:[10224040](#)).

### **CARD4 Antibody (C-term) - Protocols**

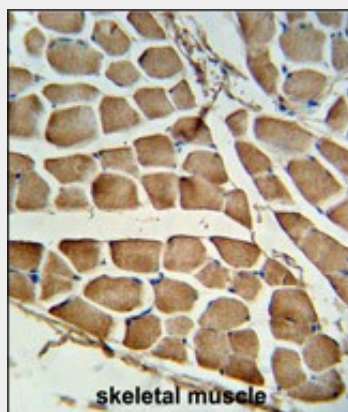
Provided below are standard protocols that you may find useful for product applications.

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

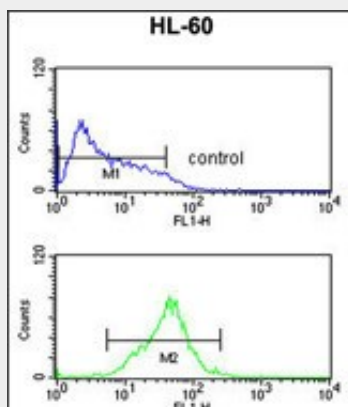
### **CARD4 Antibody (C-term) - Images**



Western blot analysis of CARD4 Antibody (C-term) (Cat. #AP8720b) in HL-60 cell line lysates (35ug/lane). CARD4 (arrow) was detected using the purified Pab.



Formalin-fixed and paraffin-embedded human skeletal muscle tissue reacted with CARD4 Antibody (C-term), which was peroxidase-conjugated to the secondary antibody, followed by DAB staining. This data demonstrates the use of this antibody for immunohistochemistry; clinical relevance has not been evaluated.



CARD4 Antibody (C-term) (Cat. #AP8720b) flow cytometric analysis of HL-60 cells (bottom histogram) compared to a negative control cell (top histogram). FITC-conjugated goat-anti-rabbit secondary antibodies were used for the analysis.

#### **CARD4 Antibody (C-term) - Background**

CARD4 is a member of the NOD (nucleotide-binding oligomerization domain) family. This member is a cytosolic protein. It contains an N-terminal caspase recruitment domain (CARD), a centrally located nucleotide-binding domain (NBD), and 10 tandem leucine-rich repeats (LRRs) in its C terminus. The CARD is involved in apoptotic signaling, LRRs participate in protein-protein interactions, and mutations in the NBD may affect the process of oligomerization and subsequent function of the LRR domain. This protein is an intracellular pattern-recognition receptor (PRR) that initiates inflammation in response to a subset of bacteria through the detection of bacterial diaminopimelic acid.

#### **CARD4 Antibody (C-term) - References**

Inohara,N., et.al., J. Biol. Chem. 274 (21), 14560-14567 (1999)  
Inohara,N.,et.al., J. Biol. Chem. 275 (36), 27823-27831 (2000)