### **Product Name: MRCKG Rabbit Polyclonal Antibody**

Catalog #: APRab14080



### **Summary**

Production Name MRCKG Rabbit Polyclonal Antibody

**Description** Rabbit Polyclonal Antibody

Host Rabbit
Application WB

**Reactivity** Human, Rat, Mouse

### **Performance**

Conjugation	Unconjugated
Modification	Unmodified
Isotype	IgG
Clonality	Polyclonal
Form	Liquid
Storage	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.
Buffer	Liquid in PBS containing 50% glycerol, and 0.02% New type preservative N.
Purification	Affinity purification

### **Immunogen**

Gene Name CDC42BPG DMPK2

**Alternative Names** 

**Gene ID** 55561.0

**SwissProt ID** Q6DT37.Synthesized peptide derived from human protein . at AA range: 1370-1450

### **Application**

**Dilution Ratio** WB 1:500-2000 ELISA 1:5000-20000

Molecular Weight 170kD

### **Background**

catalytic activity:ATP + a protein = ADP + a phosphoprotein.,cofactor:Magnesium.,enzyme regulation:Maintained in an inactive, closed conformation by an interaction between the kinase domain and the negative autoregulatory C-terminal

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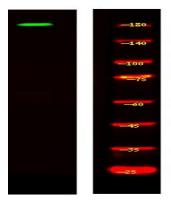
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coiled-coil region. Agonist binding to the phorbol ester binding site disrupts this, releasing the kinase domain to allow Nterminus-mediated dimerization and kinase activation by transautophosphorylation, function: May act as a downstream effector of CDC42 in cytoskeletal reorganization. Contributes to the actomyosin contractility required for cell invasion, through the regulation of MYPT1 and thus MLC2 phosphorylation., similarity: Belongs to the protein kinase superfamily. AGC Ser/Thr protein kinase family. DMPK subfamily, similarity: Contains 1 AGC-kinase C-terminal domain., similarity: Contains 1 CNH domain., similarity: Contains 1 CRIB domain., similarity: Contains 1 PH domain., similarity: Contains 1 phorbol-ester/DAGtype zinc finger, similarity: Contains 1 protein kinase domain, subcellular location: Concentrates at the leading edge of cells, subunit: Homodimer and homotetramer via the coiled coil regions. Interacts tightly with GTP-bound but not GDPbound CDC42, tissue specificity: Expressed in heart and skeletal muscle, catalytic activity: ATP + a protein = ADP + a phosphoprotein, cofactor: Magnesium, enzyme regulation: Maintained in an inactive, closed conformation by an interaction between the kinase domain and the negative autoregulatory C-terminal coiled-coil region. Agonist binding to the phorbol ester binding site disrupts this, releasing the kinase domain to allow N-terminus-mediated dimerization and kinase activation by transautophosphorylation.,function:May act as a downstream effector of CDC42 in cytoskeletal reorganization. Contributes to the actomyosin contractility required for cell invasion, through the regulation of MYPT1 and thus MLC2 phosphorylation, similarity: Belongs to the protein kinase superfamily. AGC Ser/Thr protein kinase family. DMPK subfamily, similarity: Contains 1 AGC-kinase C-terminal domain, similarity: Contains 1 CNH domain, similarity: Contains 1 CRIB domain, similarity: Contains 1 PH domain, similarity: Contains 1 phorbol-ester/DAG-type zinc finger, similarity: Contains 1 protein kinase domain, subcellular location: Concentrates at the leading edge of cells, subunit: Homodimer and homotetramer via the coiled coil regions. Interacts tightly with GTP-bound but not GDP-bound CDC42, tissue specificity: Expressed in heart and skeletal muscle.,

### **Research Area**

### **Image Data**



Western Blot analysis of HEK293 lysis, using primary antibody at 1:1000 dilution. Secondary antibody was diluted at 1:10000

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### Note

For research use only.