

Product datasheet for MC220418

Prkg1 (NM_011160) Mouse Untagged Clone

Product data:

Product Type: Expression Plasmids

Product Name: Prkg1 (NM_011160) Mouse Untagged Clone

Tag: Tag Free

Symbol: Prkg1

Synonyms: AW125416; CGKI; Gm19690; Prkg1b; Prkgr1b

Mammalian Cell Neomycin

Selection:

Vector:pCMV6-Entry (PS100001)E. coli Selection:Kanamycin (25 ug/mL)

OriGene Technologies, Inc.

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Fully Sequenced ORF:

>MC220418 representing NM_011160

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCCGCCGCGATCGCC

ATGGGCACCCTGCGGGATTTACAGTATGCGCTCCAGGAGAAGATCGAGGAGCTGAGGCAGCGGGATGCTC CAAGTATCGCTCGGTGATCCGGCCGGCCACCCAGCAGCCGCAGAAGCAGAGCGCCAGCACCTTGCAGGGT GAACCGCGCACCAAGCGCCAGGCGATCTCCGCGGAGCCCACCGCCTTCGACATCCAGGATCTCAGCCACG TGACCCTGCCCTTCTACCCCAAGAGCCCACAGTCGAAGGATCTCATAAAGGAGGCTATCCTTGACAATGA AAGGACAGCTGCATCAAGGAAGGCGATGTGGGGTCACTGGTGTACGTCATGGAAGATGGGAAGGTTG AAGTCACAAAAGAAGGCGTGAAGCTCTGCACCATGGGTCCTGGAAAAGTGTTCGGGGAGCTGGCTATACT TGTTTTCAAACAATAATGATGAGGACAGGACTCATCAAGCATACCGAGTACATGGAATTTTTAAAAAGTG TGAAAATGGAGAATATATCATCAGGCAGGGTGCAAGAGGAGACACCTTCTTCATCATCAGTAAAGGGCAG GTGAATGTTACTCGAGAAGACTCACCAAGTGAAGACCCAGTCTTCCTTAGAACTTTAGGGAAGGGAAGATT GGTTTGGAGAGAAAGCGTTGCAGGGGGGAGGATGTGAGAACAGCAAATGTTATTGCCGCAGAAGCTGTCAC CTGCCTTGTGATTGACAGAGCTCTTTCAAGCATTTGATTGGAGGACTGGACGATGTTTCTAACAAAGCA TATGAGGACGCAGAAGCAAAAGCAAAATATGAAGCCGAAGCTGCCTTCTTCGCCAACCTGAAGCTGTCTG ATTTCAACATCATTGACACCCTTGGAGTTGGAGGTTTCGGACGAGTAGAGCTGGTCCAGTTGAAAAGTGA AGAATCCAAAACATTTGCAATGAAGATCCTCAAGAAACGCCACATTGTGGACACCAGACAGCAGGAGCAC ATCCGCTCAGAGAAGCAGATCATGCAGGGGGCTCATTCTGACTTCATTGTGAGGCTGTACAGGACATTTA AAGACAGCAAATACTTGTATATGTTGATGGAAGCGTGCCTGGGTGGAGAGCTCTGGACTATTCTCAGGGA TAGGGGTTCGTTTGAAGACTCAACAACCAGGTTTTACACTGCATGTGTGGTAGAAGCATTCGCCTATCTG CATTCCAAAGGAATCATTTACAGGGACCTCAAGCCGGAGAATCTCATCCTAGATCATCGAGGCTATGCCA AGAATATGTAGCCCCAGAGATCATCCTGAACAAGGCCATGACATTTCAGCTGACTACTGGTCACTAGGA ATTCTGATGTATGAGCTTCTGACTGGCAGCCCACCTTTCTCAGGCCCAGATCCAATGAAAACCTACAATA AAAACTATGCAGGGACAACCCATCAGAGAGGTTAGGAAATTTGAAAAACGGAGTCAAAGACATTCAGAAA CACAAGTGGTTTGAGGGCTTTAATTGGGAAGGCTTAAGAAAAGGCACCTTGACACCTCCCATAATTCCAA GTGTTGCGTCACCCACAGACACAAGCAATTTTGACAGTTTCCCTGAGGACAGCGATGAGCCACCACCTGA TGACAACTCAGGCTGGGACATAGACTTCTAA

ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATTACAAGGATGACGACGATAAGGTTTAA

Restriction Sites: Sgfl-Mlul
ACCN: NM_011160
Insert Size: 2061 bp

OTI Disclaimer: Our molecular clone sequence data has been matched to the reference identifier above as a

point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative

RNA splicing form or single nucleotide polymorphism (SNP).

Components: The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube

containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).





Reconstitution Method: 1. Centrifuge at 5,000xg for 5min.

2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.

3. Close the tube and incubate for 10 minutes at room temperature.

4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid

at the bottom.

5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of

shipping when stored at -20°C.

Note: Plasmids are not sterile. For experiments where strict sterility is required, filtration with

0.22um filter is required.

RefSeq: <u>NM 011160.3, NP 035290.1</u>

RefSeq Size: 6915 bp
RefSeq ORF: 2061 bp
Locus ID: 19091
UniProt ID: P0C605
Cytogenetics: 19 C1



Gene Summary:

Serine/threonine protein kinase that acts as key mediator of the nitric oxide (NO)/cGMP signaling pathway. GMP binding activates PRKG1, which phosphorylates serines and threonines on many cellular proteins. Numerous protein targets for PRKG1 phosphorylation are implicated in modulating cellular calcium, but the contribution of each of these targets may vary substantially among cell types. Proteins that are phosphorylated by PRKG1 regulate platelet activation and adhesion, smooth muscle contraction, cardiac function, gene expression, feedback of the NO-signaling pathway, and other processes involved in several aspects of the CNS like axon guidance, hippocampal and cerebellar learning, circadian rhythm and nociception. Smooth muscle relaxation is mediated through lowering of intracellular free calcium, by desensitization of contractile proteins to calcium, and by decrease in the contractile state of smooth muscle or in platelet activation. Regulates intracellular calcium levels via several pathways: phosphorylates MRVI1/IRAG and inhibits IP3-induced Ca(2+) release from intracellular stores, phosphorylation of KCNMA1 (BKCa) channels decreases intracellular Ca(2+) levels, which leads to increased opening of this channel. PRKG1 phosphorylates the canonical transient receptor potential channel (TRPC) family which inactivates the associated inward calcium current. Another mode of action of NO/cGMP/PKGI signaling involves PKGI-mediated inactivation of the Ras homolog gene family member A (RhoA). Phosphorylation of RHOA by PRKG1 blocks the action of this protein in myriad processes: regulation of RHOA translocation; decreasing contraction; controlling vesicle trafficking, reduction of myosin light chain phosphorylation resulting in vasorelaxation. Activation of PRKG1 by NO signaling alters also gene expression in a number of tissues. In smooth muscle cells, increased cGMP and PRKG1 activity influence expression of smooth muscle-specific contractile proteins, levels of proteins in the NO/cGMP signaling pathway, down-regulation of the matrix proteins osteopontin and thrombospondin-1 to limit smooth muscle cell migration and phenotype. Regulates vasodilator-stimulated phosphoprotein (VASP) functions in platelets and smooth muscle.[UniProtKB/Swiss-Prot Function] Transcript Variant: This variant (2) encodes the longer isoform (beta). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.