

## **Product datasheet for MC226907**

## Kcnj11 (NM\_001204411) Mouse Untagged Clone

**Product data:** 

**Product Type:** Expression Plasmids

**Product Name:** Kcnj11 (NM\_001204411) Mouse Untagged Clone

Tag: Tag Free
Symbol: Kcnj11

Synonyms: Kir6.2; mBIR

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

Cell Selection: Neomycin

Fully Sequenced ORF: >MC226907 representing NM\_001204411

Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC

GCCGCGATCGCC

ATGGTCTGGTGGCTCATCGCCTTCGCCCACGGTGACCTGGCCCCCGGAGAGGGCACCAATGTGCCCTGCG
TCACAAGCATCCACTCCTTTTCATCTGCCTTCCTTTTCTCCATCGAGGTCCAGGTGACCATTGGTTTCGG
CGGGCGCATGGTGACAGAGGAATGTCCCCTGGCCATCCTCATTCTCATTGTGCAGAATATCGTCGGGCTG
ATGATCAACGCCATCATGCTGGGCTGCATCTTCATGAAAACGGCCCAGGCCCATCGGCGGGCAGAAACCC
TCATCTTCAGCAAGCATGCTGTGATCACCCTGCGCCATGGCCGCCTGTGCTTCATGCTGCGCGTAGGGGA
CCTCCGAAAGAGCATGATCATTAGCGCCACCATCCACATGCAGGTGGTGCCCAAGACCACCAGCCCCGAG
GCGAAGTTGTGCCTCTCCACCAGGTAGACATCCCCATGGAGAATGGCGTGGTGAAACGCCCCCGAG
GCCGAACTTCTCCACCACGTCATCGACTCCAACAGCCCGCTCTACGACCTGGCTCCTAGTGA
CCTGCACCACCACCACCAGGACCTGGAGATCATTGTCATCTTGGAAGGCGTGGTAGAAACCACGGGCATCACC
ACCCAGGCCCGCACCTCCTACCTAGCTGACGAGATTCTATGGGGGGCAGCGCTTTTGTCCCCATTGTGGCCG
AGGAGGACGGCCGCTATTCTGTGGACTACTCCAAATTTGGTAACACCATTAAAGTGCCCACACCACTCTG
CACAGCCCGCCAGCTTGATGAGGACCGCAGTCTGCTGGATGCCCTGGCCTCGCCTCGTCGCGGGGGCCC
CTGCGCAAGCCGCAGTTGTGGCCGAAGGCCAAGCCCAAGTTTAGCATCTCTCCAGATTCCTTGTCCT

 ACGCGTACGCGGCCGCTCGAGCAGAAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCCTGGATT

ACAAGGATGACGACGATAAGGTTTAA

**Restriction Sites:** Sgfl-Mlul

**ACCN:** NM\_001204411

**Insert Size:** 912 bp



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## Kcnj11 (NM\_001204411) Mouse Untagged Clone - MC226907

**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).

**OTI Annotation:** Clone contains native stop codon, and expresses the complete ORF without any c-terminal

tag.

**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.

**RefSeq:** NM 001204411.1, NP 001191340.1

RefSeq Size:2667 bpRefSeq ORF:912 bpLocus ID:16514

**Cytogenetics:** 7 29.66 cM

**Gene Summary:** This receptor is controlled by G proteins. Inward rectifier potassium channels are

characterized by a greater tendency to allow potassium to flow into the cell rather than out of it. Their voltage dependence is regulated by the concentration of extracellular potassium; as external potassium is raised, the voltage range of the channel opening shifts to more positive voltages. The inward rectification is mainly due to the blockage of outward current by internal magnesium. Can be blocked by extracellular barium. Can form cardiac and smooth muscletype KATP channels with ABCC9. KCNJ11 forms the channel pore while ABCC9 is required for

activation and regulation (By similarity), [UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (2) differs in the 5' UTR, lacks a portion of the 5' coding region and initiates translation at a downstream in-frame start codon, compared to variant 1. The encoded isoform (2) is shorter than isoform 1. Sequence Note: The RefSeq transcript and protein were derived from genomic sequence to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were

based on alignments.