

## Product datasheet for **MC229475**

### **Kcnma1 (NM\_001253369) Mouse Untagged Clone**

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Kcnma1 (NM_001253369) Mouse Untagged Clone
Tag:	Tag Free
Symbol:	Kcnma1
Synonyms:	5730414M22Rik; BKCa; MaxiK; mSlo; mSlo1; Slo; Slo1
Vector:	pCMV6-Entry (PS100001)
E. coli Selection:	Kanamycin (25 ug/mL)
Cell Selection:	Neomycin
Fully Sequenced ORF:	>MC229475 representing NM_001253369 Red=Cloning site Blue=ORF Orange=Stop codon

TTTTGTAATACGACTCACTATAGGGCGGCCGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGGATCGC**C

ATGGCAAACGGTGGCGGGCGGGCGGGCGGCAGCAGCGGGCGGGCGGGCGGGCGGGAGGCGAGCGGTC  
TTAGAATGAGCAGCAATATCCACGCGAACCATCTCAGCCTAGACGCGTCCTCCTCCTCTTCTCCTCCT  
CTCTTCTCTTCTTCTCCTCCTCCTTCTCCTCCTCGTCTCGGTCCACGAGCCCAAGATGGATGCGCTCATC  
ATACCGGTGACCATGGAGGTGCCGTGCGACAGCCGGGGCAACGCATGTGGTGGGCTTTCTTGGCCTCCT  
CCATGGTGACTTTCTTCGGGGCCTTTCATCATCTTGCTCTGGCGGACGCTCAAGTACCTGTGGACCGT  
TTGCTGCCACTGCGGGGCAAGACGAAGGAGGCCAGAGAATAAACAATGGCTCCAGCCAGGCAGATGGT  
ACTCTCAAGCCAGTGGACGAAAAAGAGGAGGTGGTGGCAGCCGAGGTCGGCTGGATGACATCTGTGAAGG  
ACTGGGCGGGGTGATGATATCCGCCCAGACACTGACTGGCAGAGTCTGGTGTGTTAGTCTTTGCTCT  
CAGCATTGGTGCCTCGTAATACTTCATAGACTCGTCAAACCAATAGAATCCTGCCAGAATTTCTAC  
AAAGATTTACATTACAGATCGACATGGCTTCAACGTGTTCTCCTCCTACTTTGGCTTCCGTTTA  
TTGCAGCCAACGATAAGCTGTGGTTCTGGCTGGAAGTGAATTCAGTAGATTCTTCCAGCCCTTC  
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CGGGGGCTCCTATAGCGCGTTAGTGAAGAAAGCACATTGTAGTCTGTGGACACACTCTGGAGAGT  
GTCTCTAACTTCTGAAGGACTTCTGCACAAGGACCGGGATGATGCAACGTGGAGATTGTCTTCTCTTC  
ACAACATCTCCCCTAACCTTGAGCTTGAGGCTCTGTTCAACGCGATTTCACTCAGGTGGAGTTTATCA  
GGGCTCTGTCCATCCACATGATCTTGCCAGAGTCAAGATAGAGTCAAGGAGTGCATGCCTGATCCTT  
GCCAATAAGTATTGCGCTGACCCGGATGCAGAAGATGCCTCCAACATCATGAGAGTGTCTCCATCAAAA



ACTACCACCCAAAGATCAGGATCATCACTCAGATGCTGCAGTATCACAACAAGGCCCATCTGCTCAACAT  
 CCCCAGCTGGAAGTGGAAAGAGGGTGATGACGCAATATGCCTTGACAGCTCAAGTTGGGTTTCATAGCC  
 CAGAGCTGTCTGGCTCAAGGCCTCTCCACAATGCTTGCCAATCTTTCTATGAGGTCATTATAAAGA  
 TTGAGGAAGACACATGGCAGAAATACTACTTGAAGGAGTCTCCAATGAAATGTACACAGAATATCTCTC  
 CAGTGCCTTCGTGGGTCTGTCTCCCTACTGTTGTGAGCTGTGTTTGTGAAGCTTAAGCTCCTGATG  
 ATAGCCATTGAGTACAAGTCTGCCAACAGAGAGAGCCGAAAGCGAATATTAATTAACCTGGGA  
 ACCACCTTAAGATCCAAGAAGTACTTTAGGATTTTTTCATCGCAAGTATGCCAAGAAGTTAAAAGGGC  
 ATTTTTTACTGCAAGGCTGTATGATGACGTACAGATCCCAAAGAATTAATAAATGTGGCTGCAGG  
 CGGCTTGAAGATGAGCAGCCCAACCTGTCCACAAAAAACAACGTAATGGGGCATGAGGAACT  
 CGCCCAACACCTCCCGAAGCTGATGAGGCATGACCCCTGTTAATTCCTGGCAATGATCAGATTGACAA  
 CATGGACTCCAATGTGAAAAAGTACGACTCCACTGGAATGTTTCACTGGTGTGCACCCAAGGAGATTGAG  
 AAAGTCATCTTGACTCGAAGTGAAGCTGCCATGACTGTCCTGAGTGGCCATGTCGTAGTCTGCATCTTG  
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 TCATGAGCTCAAACACATTGTGTTTGTGGGCTCCATTGAGTACCTCAAGAGGGAGTGGGAAACACTGCAC  
 AACTTCCCGAAAGTGTCCATATTGCCTGGTACACCATTAGTCGGGCTGATTTAAGGGCTGTCAACATCA  
 ACCTCTGTGACATGTGCGTTATCCTGTACGCAATCAGAATAATATTGATGATACTTCGCTTCAGGACAA  
 GGAATGCATCTTGGCGTCACTCAACATCAAATCTATGCAGTTTATGACAGCATCGGGTCTTGCAGGCT  
 AATTCCTCAAGGATTCACACCTCCTGGAATGGACAGATCATCACCAGCAACAGCCAGTGCACGGGATGT  
 TACGCCAGCCGTCATCACTCAACTGGGGTCAACATTCCTCATCACGGAACCTCGTAAGCCGGGCAAGTT  
 GCCTTTGGTATCAGTCAATCAGGAAAAAACAGTGGGACGCACATTCTAATGATAACGGAGTTGGTGAAT  
 GATACCAATGTTCAAGTTTTGGACCAAGACGATGACGATGACCCTGACACAGAGCTGTACCTCACACAGC  
 CCTTTGCTTGTGGGACAGCATTGCGGTCAGCGTCTGGACTCACTCATGAGCGGACATACTTCAATGA  
 CAATACCTCACCTAATACGGACCCTGGTACAGGAGGCCACACCAGAGCTCGAGGCTCTAATAGCT  
 GAGGAGAATGCACCTCGAGGAGGCTACAGCACTCCGCAGACATTGGCCAACAGGGACCGTTGCCGAGTGG  
 CCCAGTTAGCCCTGTTAGATGGTCCCTTGCAGACTTAGGGGATGGTGGTTGTTATGGTATCTGTTCTG  
 CAAAGCTCTGAAAACATATAATATGCTTTGTTTGGAAATTTACCGGCTGAGAGATGCCACCTCAGCACC  
 CCCAGCCAGTGTACAAAAGGTACGTATCACCAATCCTCCCTACGAGTTTGTAGCTCGTACCAACAGACC  
 TGATCTTCTGCCTGATGCAGTTTGACCACAACGCTGGCCAATCCCGGGCCAGTCTGTCTCATTCTCCCA  
 CTCCTCACAGTCGTCCAGTAAGAAGAGCTCCTCCGTCCTCCATCCCGTCCACAGCAAATCGGCCGAAC  
 CGGCCAAGTCCAGGGAGTCCCGGACAAACAGAACAGAAAAGAATGGTTTACAGATGA

AGCGGACCGACGCGTACGCGCCGCTCGAGCAGAACTCATCTCAGAAGAGGATCTGGCAGCAAATGATATCC  
 TGGATTACAAGGATGACGACGATAAGGTTTAA

- Restriction Sites:** SgfI-RsrII
- ACCN:** NM\_001253369
- Insert Size:** 3630 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).
- 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).

<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>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.</li></ol>
<b>RefSeq:</b>	<u>NM_001253369.1</u> , <u>NP_001240298.1</u>
<b>RefSeq Size:</b>	5037 bp
<b>RefSeq ORF:</b>	3630 bp
<b>Locus ID:</b>	16531
<b>UniProt ID:</b>	<u>Q08460</u>
<b>Cytogenetics:</b>	14 A3
<b>Gene Summary:</b>	<p>Potassium channel activated by both membrane depolarization or increase in cytosolic Ca(2+) that mediates export of K(+). It is also activated by the concentration of cytosolic Mg(2+). Its activation dampens the excitatory events that elevate the cytosolic Ca(2+) concentration and/or depolarize the cell membrane. It therefore contributes to repolarization of the membrane potential. Plays a key role in controlling excitability in a number of systems, such as regulation of the contraction of smooth muscle, the tuning of hair cells in the cochlea, regulation of transmitter release, and innate immunity. In smooth muscles, its activation by high level of Ca(2+), caused by ryanodine receptors in the sarcoplasmic reticulum, regulates the membrane potential. In cochlea cells, its number and kinetic properties partly determine the characteristic frequency of each hair cell and thereby helps to establish a tonotopic map. Kinetics of KCNMA1 channels are determined by alternative splicing, phosphorylation status and its combination with modulating beta subunits. Highly sensitive to both iberiotoxin (IbTx) and charybdotoxin (CTX).[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (13) has multiple differences in the coding region, one of which results in a frameshift, compared to variant 1. The resulting isoform (13) is shorter and has a distinct C-terminus, compared to isoform 1.</p>