

Product datasheet for **MC203472**

Gria1 (NM_008165) Mouse Untagged Clone

Product data:

| | |
|---------------------------|---|
| Product Type: | Expression Plasmids |
| Product Name: | Gria1 (NM_008165) Mouse Untagged Clone |
| Tag: | Tag Free |
| Symbol: | Gria1 |
| Synonyms: | 2900051M01Rik; Glr-1; Glr1; GluA1; Glur-1; GluR-A; gluR-K1; Glur1; GluRA; HIPA1 |
| Mammalian Cell Selection: | Neomycin |
| Vector: | PCMV6-Kan/Neo (PCMV6KN) |
| E. coli Selection: | Kanamycin (25 ug/mL) |
| Fully Sequenced ORF: | >BC056397 |

```

AAAGAATCAAAGGGAGGGGAGGGAAGACCAAACTATGGTTGGACCAGGGCTTCTTTTTCGCCAATGTAA
AAAGGAATATGCCGTACATCTTTGCCTTTTCTGCACCGTTTTCTAGGTGCGGTTGTGGGTGCCAATTT
CCCCAACCAATATCCAGATAGGGGATTATTTCCAAACCAACAATCACAGGAACATGCGGCTTTTAGGTTT
GCTTTGTCACTACACTCACGGAGCCCCCAAGCTGCTTCCCAGATCGATATTGTGAACATCAGCGACAGCT
TTGAGATGACTTACCGATTCTGTTCCCAGTTCTCCAAAGGAGTGTACGCCATCTTTGGATTTTATGAACG
AAGGACTGTCAACATGCTGACCTCCTTCTGTGGGGCCCTCCATGTGTGCTTCATCACTCCAAGTTTTCCC
GTTGACACATCCAATCAGTTTGTCTTCCAGCTGCGCCCGGAACTACAGGAAGCTCTCATTAGCATTATCG
ACCATTACAAGTGGCAGACTTTTGTCTACATTTATGATGCTGACCGGGCCTGTGAGTCTGAGAGAGT
CTTGGATACAGCCGCGAGAAGAACTGGCAGGTGACGGCTGTCAACATTCTAACAACCACGGAGGAAGGA
TACCGGATGCTCTTTCAGGACCTGGAGAAGAAAAGGAGAGGCTGGTGGTGGTGGACTGTGAATCAGAAC
GCCTCAACGCCATCCTGGGCCAGATTGTGAAGTAGAAAAGAACGGCATCGGGTACCACTACATCCTCGC
CAACCTGGGCTTCATGGACATTGACTTAAATAAGTTCAAGGAGAGTGGAGCCAATGTGACAGGTTTTCCAA
CTGGTGAACACTACAGACACGATCCCAGCCAGAATCATGCAGCAGTGGAGGACAAGTGCAGCTCGGGACC
ACACCAGGGTGGACTGGAAGAGGCCAAAGTACACTTCTGCTCTTACCTATGATGGTGTGAAGGTGATGGC
GGAGGCCCTCCAGAGCCTGCGGAGGCAGAGGATTGACATATCCCGCGGAGGGAATGCTGGGGACTGTCTG
GCTAACCCAGCTGTGCCCTGGGGCCAAGGGATCGACATCCAGAGAGCCCTGCAGCAGGTGCGCTTTGAAG
GTTTGACAGGAAATGTGCAGTTTAAACGAGAAAGGGCGCCGGACCAACTACACCCTCATGTGATCGAAAT
GAAGCATGATGGAATCCGCAAGATTGGTTACTGGAATGAAGATGATAAATTTGTCCCGCAGCCACGGAC
GCTCAGGCTGGAGGGACAACCTCAAGCGTCCAGAATAGAACCTACATCGTCACGACTATCCTCGAAGATC
CTTACGTGATGCTTAAAAAGAATGCCAACCAATTTGAAGGCAATGACCGCTATGAGGGCTACTGCGTGGA
ACTGGCTGCGGAGATCGCCAAGCACGTGGGCTATTCTACCGACTTGAGATTGTCAGCGACGGCAAATAC
GGAGCCCCGGATCCTGACACAAAGGCCTGGAATGGCATGGTGGGAGAGCTAGTCTATGGAAGAGCAGATG
TGGCGGTGGCCCCCTTGACCATAACCTTGGTCCGGGAGGAAGTCACTCGACTTCTCCAAGCCATTCATGAG
TTTGGAACTCCATTATGATTAAGAAGCCACAGAAGTCCAAGCCAGGTGCTTCTCCTTTCTTGACCCT
TTGGCCTACGAGATCTGGATGTGTATAGTGTTCCTACATTGGAGTGAGCGTCGCTCCTTCTCCTGGTCA
GCCGTTTCAGTCTTATGAATGGCACAGTGAAGAGTTTGAAGAAGGACGAGATCAGACAACCAAGTACCA

```



[View online »](#)

GTCAAATGAGTTTGGCATATTCAACAGCCTGTGGTTCTCGCTGGGGCCTTCATGCAGCAAGGATGTGAC
 ATTTCCCCAGGTCCCTGTCTGGACGCATCGTCGGCGGTGTCTGGTGGTCTTCACTTTGATTATCATCT
 CCTCATACACAGCCAACCTGGCTGCCTTCTGACTGTGAAAGGATGGTGTCTCCATCGAGAGTGCAGA
 GGACCTGGCAAAGCAGACGGAAATTGCTTATGGGACATTGGAAGCAGGATCCACTAAGGAGTTCTCAGG
 AGGTCTAAAATCGCTGTGTTTGAGAAGATGTGGACATACATGAAGTCTGCAGAACCGTCTGTGTTGTTT
 GGACCACAGAGGAGGGCATGATCAGAGTGAGAAAGTCTAAAGGCAAATATGCCTACCTCTGGAGTCCAC
 CATGAATGAGTACATTGAGCAACGCAAGCCCTGTGACACCATGAAAGTGGGAGGTAACCTGGATTCCAAA
 GGCTATGGCATTGCAACACCCAAGGGTCCGCCCTGAGAAATCCAGTAAACCTGGCAGTGTAAAACTGA
 ACGAGCAGGGGCTTTTGGACAAATTGAAAAACAATGGTGGTACGACAAGGGCGAGTGGCGCAGCGGGG
 AGGTGACTCCAAGGACAAGACCAGTGTCTGAGCCTGAGCAATGTGGCAGGCGTGTCTACATCCTGATT
 GGAGGGCTGGGATTGGCCATGCTGGTTGCCTTAATCGAGTTCTGCTACAAATCCCGTAGCGAGTGAAGC
 GGATGAAGGGTTTCTGTTGATTCCACAGCAATCCATCAATGAAGCCATACGGACATCGACCCTCCCCAG
 GAAACAGCGGGGAGGAGCCAGCGGAGGAAGTGGCAGTGGAGAGAATGGCAGAGTGGTCCAGCCAGGACTTC
 CCCAAGTCCATGCAATCCATTCCCTGCATGAGCCACAGTTCAGGGATGCCCTTGGGAGCCACAGGATTGT
 AACTGGAGCAGACAGGAAACCTTGGGGAGCAGGCTCAGGCTTCCCAGCCCCATCCCAAGCCCTTCAGT
 GCCAAAAACAAGAAATGAAACACCGCCTCCAACCACCACAACCATATGGAGGGCAATTCAGCCAATGTC
 CCTGAAGAATTAATAAAAAAAAAAAAAAAAAATCCACTAGACCACCATCCCTTTTGTGTTGGGATTTTTTTCA
 ACCTTTTCCATTTGCTGAGTGAAGATGGTCACTAATCTATGCTGCAATAACGGGAGAGATCCTGTCCAA
 GGAAGTCTAACGTCTATAAAAAATGGAGTCACTGGGATATCAATGAAGAAATCAAAGTGTAAATTTAAT
 TCAGTTGTTAATGTGCTTAGTGTGTGCATTTTTCTTACTATTAACCCATGGTTTGCAGGCTCTGTTC
 AACCTTTTCTTCTTCTATCCCCAGCCCCACCCACCCACCCCTTCCCATCACTTTTCAGTTTCA
 GATTGAAGACTCACGATTGGTTCACCTTACAAGCAAGGAAAGGAATATTGACCTTCAAAGTCACTCAGT
 GTTGGAGTCTCTATGAGGCCTTCTATCTTCTTGGCCCATGCCTTGGATGGAGACAGACACTACTGGAGAA
 GGGGGTGCCTTCCCAGAAAGAATACTACCTAAATGCTGGTTCATCAGAGAGTAACTGTGAAGAGCATC
 TCAGATAAGGGTTCTTAAAGGGGGGATGGGCAGAGTACAGTAGAGGAGGACTTGTAGGTTCAATTGTTT
 CCCGGAAGAAGCTTCTGCAATGCCAGGAGCATCCCACAGGCAATAGAGTACCAATATGGTAGACCTCT
 TTCACAGTCAATCTCAGCCCTGAAATATGTTTACTATGCTCAGGCCAAAGGCAATGTGGGGCCAA
 AGGGCAACTGGCCTTCTGACCCACTAGAGCCTTGCAAAGAGTGGCAGTCAAGGGGGTGCATACATGAT
 TCTTGCTTTTCTTTGCTTGTGTCTATACTGGATCCCATGGTGGAGTAGGACTCTCTGTGGAGCCTC
 TGGCATAAGAGGAAATGTTGCCGAGCTAAGAGCAAAAGGAAGGTGATAGGAATATGGAAGTCCATCC
 AAGCCCTGACACAGAGCCTTCTGTGCTGTTTCTGCTCCTAGGTTTGGAGACCAATATGAACTTGTGA
 GTTGTGTCATGACAGAGAACCGGGTTTTGTGTTAATAGAGTATTGTTTTAGCATGTGATCAGATCTTAA
 GAAAGAAAGAAAGAAAGACAGCAAAAGCCTGAATTTCTTTTGGAGAGTGGAGATGAGGTACATCAGGGGAC
 CTTAAGATGCCTGAGCCATCTTTAACTCTAACACAGGGATACTAACCTTTTCCCTGGCTAAGACAGA
 CTGAGGGCATTGCATTACAATGGCCAGAAACTACACCCATGAGTGCAACGACCAAGGTGAAGGACAGCCA
 GACAGGAAAGGTGACAAAAGCCACCCTGAGTGATACTAGGCTGCCTGAACCTTTGGGTGGGAGTGGAAAGT
 CCCTAGCACACAGCTTTGTGGAGACCTCCACTCTCACTCTTGGTGAACCTGACTTCCACTTCTTATGCAG
 CCACTCCCAGGTGCATGCTTCAGTGAAGTCCACATGCCACTAGCTTTTACCAATAACCATATTCTGGCTT
 ATTTCTACCCTCTGCTTCCATTCAATCTTCCCCTAGGGTTATCATTGGCAATGGACGAGCTGAAG
 ATTTTGGAGCCTATCCATGGTTCAGGGCAGTAGAACTGGCTTCCCTATGGATGCTTCCAGCGGACAACC
 ACCATCTCTGTGCTCTCTCATTCTCTTCCACTGCCATAGCAAAGAAACATCCCAGAATCGACCTTCT
 TCCATTGATCCATGTCTCCCGTGCACCATGGGCATGGCATGTGAAGCAATGCTCTGCAGATGGATTAC
 CACCTTTTATTTCTGCCAGCTAACATCCTCCCAGCCTCTGCCTACTACTGTAGACTCCGGTGTGACCCA
 CCACTGGGGAGTTCACACCTAGAAAAGAGATGGAAGTAACTCTACTATCAGCTGCTTCAAAAACCTTTTG
 TCTACGGCCTGGGAAGATGTTCTCCAGCCCTGAGCCTATTGGTTGGATAGAGAGAGACTGTGGTGGAAA
 TGAGGGTCTTGTGGGAGGGTCCATGACTCCGGGAAGACGCTTCTCAACGAGGGATGTGTGGAAAGAA
 AACAGAAGTGAGAAAGAAGTTACCTTGTATTATGTATTTTACTACACTTTTCTTAAAAATGGAGCAGTG
 GAAAAACTCTGAAAGAGATTGACATTTTCTCAACAGGAATCCACTTAAACAGTTCTGGCTTTCATTAA
 ATTTTGTCTTTTGGTACCTGGGCCTTTTATTTAACATATATTTATTTTAACTCTTGGAAAGATGTGT
 GAGGGAAAAAATAACTCTCATTGATCAGACACTAAATATTTGGGGTAGGGATTCTGTCTTATTTTCA
 AATAGCCAGGGTTTTTCTTCAATTTTTTGGTATTTGCATAAAGTGAACAATATCACAGAATTAATCAC
 TGTGGACCATTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA

| | |
|-------------------------------|---|
| Restriction Sites: | Ascl-NotI |
| ACCN: | NM_008165 |
| Insert Size: | 2724 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: | <ol style="list-style-type: none">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: | BC056397 , AAH56397 |
| RefSeq Size: | 5434 bp |
| RefSeq ORF: | 2724 bp |
| Locus ID: | 14799 |
| Cytogenetics: | 11 34.51 cM |
| Gene Summary: | <p>Ionotropic glutamate receptor. L-glutamate acts as an excitatory neurotransmitter at many synapses in the central nervous system. Binding of the excitatory neurotransmitter L-glutamate induces a conformation change, leading to the opening of the cation channel, and thereby converts the chemical signal to an electrical impulse. The receptor then desensitizes rapidly and enters a transient inactive state, characterized by the presence of bound agonist. In the presence of CACNG4 or CACNG7 or CACNG8, shows resensitization which is characterized by a delayed accumulation of current flux upon continued application of glutamate.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (2) uses an alternate in-frame exon, compared to variant 1, resulting in a protein (isoform 2) with a distinct internal segment compared to isoform 1.</p> <p>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.</p> |