

Product datasheet for RC205437L2V

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CAMK2G (NM_172169) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: CAMK2G (NM_172169) Human Tagged ORF Clone Lentiviral Particle

Symbol: CAMK2G

Synonyms: CAMK; CAMK-II; CAMKG; MRD59

Mammalian Cell

Selection:

None

Vector: pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_172169 **ORF Size:** 1581 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC205437).

Sequence:

OTI Disclaimer: The molecular sequence of this clone aligns with the gene accession number as a point of

reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeg: NM 172169.1

RefSeq Size:3731 bpRefSeq ORF:1584 bp

Locus ID: 818

 UniProt ID:
 Q13555

 Cytogenetics:
 10q22.2

Protein Families: Druggable Genome, Protein Kinase



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Protein Pathways: Calcium signaling pathway, ErbB signaling pathway, Glioma, GnRH signaling pathway, Long-

term potentiation, Melanogenesis, Neurotrophin signaling pathway, Olfactory transduction,

Oocyte meiosis, Wnt signaling pathway

MW: 59 kDa

Gene Summary: The product of this gene is one of the four subunits of an enzyme which belongs to the

serine/threonine protein kinase family, and to the Ca(2+)/calmodulin-dependent protein kinase subfamily. Calcium signaling is crucial for several aspects of plasticity at glutamatergic synapses. In mammalian cells the enzyme is composed of four different chains: alpha, beta, gamma, and delta. The product of this gene is a gamma chain. Many alternatively spliced transcripts encoding different isoforms have been described but the full-length nature of all

the variants has not been determined.[provided by RefSeq, Mar 2011]