

Product datasheet for RC204457L3V

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

KCNN4 (NM_002250) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: KCNN4 (NM_002250) Human Tagged ORF Clone Lentiviral Particle

Symbol: KCNN4

Synonyms: DHS2; hIKCa1; hKCa4; hSK4; IK; IK1; IKCA1; KCa3.1; KCA4; SK4

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-Myc-DDK-P2A-Puro (PS100092)

Tag: Myc-DDK
ACCN: NM_002250

ORF Size: 1281 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC204457).

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 002250.2

 RefSeq Size:
 2240 bp

 RefSeq ORF:
 1284 bp

 Locus ID:
 3783

 UniProt ID:
 015554

 Cytogenetics:
 19q13.31

Domains: SK_channel, CaMBD

Protein Families: Druggable Genome, Ion Channels: Potassium, Transmembrane





ORIGENE

MW: 47.7 kDa

Gene Summary: The protein encoded by this gene is part of a potentially heterotetrameric voltage-

independent potassium channel that is activated by intracellular calcium. Activation is followed by membrane hyperpolarization, which promotes calcium influx. The encoded protein may be part of the predominant calcium-activated potassium channel in T-

lymphocytes. This gene is similar to other KCNN family potassium channel genes, but it differs enough to possibly be considered as part of a new subfamily. [provided by RefSeq, Jul 2008]