

Product datasheet for RC202092L2V

OriGene Technologies, Inc.

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GJB2 (NM_004004) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: GJB2 (NM_004004) Human Tagged ORF Clone Lentiviral Particle

Symbol: GJB2

Synonyms: BAPS; CX26; DFNA3; DFNA3A; DFNB1; DFNB1A; HID; KID; NSRD1; PPK

Mammalian Cell

Selection:

None

Vector: pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_004004

ORF Size: 678 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC202092).

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 004004.3

 RefSeq Size:
 2347 bp

 RefSeq ORF:
 681 bp

 Locus ID:
 2706

 UniProt ID:
 P29033

 Cytogenetics:
 13q12.11

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

MW: 26.2 kDa







Gene Summary:

This gene encodes a member of the gap junction protein family. The gap junctions were first characterized by electron microscopy as regionally specialized structures on plasma membranes of contacting adherent cells. These structures were shown to consist of cell-to-cell channels that facilitate the transfer of ions and small molecules between cells. The gap junction proteins, also known as connexins, purified from fractions of enriched gap junctions from different tissues differ. According to sequence similarities at the nucleotide and amino acid levels, the gap junction proteins are divided into two categories, alpha and beta. Mutations in this gene are responsible for as much as 50% of pre-lingual, recessive deafness. [provided by RefSeq, Oct 2008]