

Product datasheet for RC215676L4V

OriGene Technologies, Inc.

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PANK2 (NM_153638) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: PANK2 (NM_153638) Human Tagged ORF Clone Lentiviral Particle

Symbol: PANK2

Synonyms: C20orf48; HARP; HSS; NBIA1; PKAN

Mammalian Cell

Selection:

Puromycin

Vector: pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_153638 **ORF Size:** 1710 bp

ORF Nucleotide

Sequence:
OTI Disclaimer:

The ORF insert of this clone is exactly the same as(RC215676).

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 153638.2, NP 705902.2

 RefSeq Size:
 2280 bp

 RefSeq ORF:
 1713 bp

 Locus ID:
 80025

 UniProt ID:
 Q9BZ23

 Cytogenetics:
 20p13

Protein Families: Druggable Genome

Protein Pathways: Metabolic pathways, Pantothenate and CoA biosynthesis





ORIGENE

MW: 59.1 kDa

Gene Summary:

This gene encodes a protein belonging to the pantothenate kinase family and is the only member of that family to be expressed in mitochondria. Pantothenate kinase is a key regulatory enzyme in the biosynthesis of coenzyme A (CoA) in bacteria and mammalian cells. It catalyzes the first committed step in the universal biosynthetic pathway leading to CoA and is itself subject to regulation through feedback inhibition by acyl CoA species. Mutations in this gene are associated with HARP syndrome and pantothenate kinase-associated neurodegeneration (PKAN), formerly Hallervorden-Spatz syndrome. Alternative splicing, involving the use of alternate first exons, results in multiple transcripts encoding different isoforms. [provided by RefSeq, Jul 2008]