

Product datasheet for RC201701L4V

OriGene Technologies, Inc.

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p38 (CRK) (NM_016823) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: p38 (CRK) (NM_016823) Human Tagged ORF Clone Lentiviral Particle

Symbol: p38

Synonyms: CRKII; p38

Mammalian Cell Puromycin

Selection:

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

Tag: mGFP

ACCN: NM_016823

ORF Size: 912 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC201701).

OTI Disclaimer:

Sequence:

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 016823.2

 RefSeq Size:
 3225 bp

 RefSeq ORF:
 915 bp

 Locus ID:
 1398

 UniProt ID:
 P46108

 Cytogenetics:
 17p13.3

 Domains:
 SH2. SH3

Protein Families: Druggable Genome, Transcription Factors





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Protein Pathways: Chemokine signaling pathway, Chronic myeloid leukemia, ErbB signaling pathway, Fc gamma

R-mediated phagocytosis, Focal adhesion, Insulin signaling pathway, MAPK signaling pathway, Neurotrophin signaling pathway, Pathways in cancer, Regulation of actin cytoskeleton, Renal

cell carcinoma

MW: 33.8 kDa

Gene Summary: This gene encodes a member of an adapter protein family that binds to several tyrosine-

phosphorylated proteins. The product of this gene has several SH2 and SH3 domains (src-homology domains) and is involved in several signaling pathways, recruiting cytoplasmic proteins in the vicinity of tyrosine kinase through SH2-phosphotyrosine interaction. The N-terminal SH2 domain of this protein functions as a positive regulator of transformation whereas the C-terminal SH3 domain functions as a negative regulator of transformation. Two alternative transcripts encoding different isoforms with distinct biological activity have been

described. [provided by RefSeq, Jul 2008]