

Product datasheet for RC211209L3V

OriGene Technologies, Inc.

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TAS2R38 (NM_176817) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: TAS2R38 (NM_176817) Human Tagged ORF Clone Lentiviral Particle

Symbol: TAS2R38

Synonyms: PTC; T2R38; T2R61; THIOT

Mammalian Cell

Selection:

ORF Size:

Puromycin

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

999 bp

Tag: Myc-DDK

ACCN: NM_176817

ORF Nucleotide

Sequence:

The ORF insert of this clone is exactly the same as(RC211209).

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 176817.2

 RefSeq Size:
 1143 bp

 RefSeq ORF:
 1002 bp

 Locus ID:
 5726

 UniProt ID:
 P59533

 Cytogenetics:
 7q34

Protein Families: Druggable Genome, Transmembrane

Protein Pathways: Taste transduction





ORIGENE

MW: 37.9 kDa

Gene Summary:

This gene encodes a seven-transmembrane G protein-coupled receptor that controls the ability to taste glucosinolates, a family of bitter-tasting compounds found in plants of the Brassica sp. Synthetic compounds phenylthiocarbamide (PTC) and 6-n-propylthiouracil (PROP) have been identified as ligands for this receptor and have been used to test the genetic diversity of this gene. Although several allelic forms of this gene have been identified worldwide, there are two predominant common forms (taster and non-taster) found outside of Africa. These alleles differ at three nucleotide positions resulting in amino acid changes in the protein (A49P, A262V, and V296I) with the amino acid combination PAV identifying the taster variant (and AVI identifying the non-taster variant). [provided by RefSeq, Oct 2009]