

Product datasheet for RC216541L4V

OriGene Technologies, Inc.

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GABA A Receptor alpha 5 (GABRA5) (NM_000810) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: GABA A Receptor alpha 5 (GABRA5) (NM_000810) Human Tagged ORF Clone Lentiviral Particle

Symbol: GABA A Receptor alpha 5

Synonyms: DEE79; EIEE79

Mammalian Cell

Puromycin

Selection:

Vector:

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

Tag: mGFP

ACCN: NM_000810

ORF Size: 1386 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC216541).

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

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.

RefSeq: <u>NM 000810.2</u>

 RefSeq Size:
 2352 bp

 RefSeq ORF:
 1389 bp

 Locus ID:
 2558

 UniProt ID:
 P31644

 Cytogenetics:
 15q12

Domains: Neur_chan_memb, Neur_chan_LBD





GABA A Receptor alpha 5 (GABRA5) (NM_000810) Human Tagged ORF Clone Lentiviral Particle – RC216541L4V

Protein Families: Druggable Genome, Ion Channels: Cys-loop Receptors, Transmembrane

Protein Pathways: Neuroactive ligand-receptor interaction

MW: 52.1 kDa

Gene Summary: GABA is the major inhibitory neurotransmitter in the mammalian brain where it acts at GABA-

A receptors, which are ligand-gated chloride channels. Chloride conductance of these channels can be modulated by agents such as benzodiazepines that bind to the GABA-A receptor. At least 16 distinct subunits of GABA-A receptors have been identified. Transcript variants utilizing three different alternative non-coding first exons have been described.

[provided by RefSeq, Jul 2008]