

Product datasheet for RC223075L2V

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

FUT8 (NM_178155) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: FUT8 (NM 178155) Human Tagged ORF Clone Lentiviral Particle

Symbol: FUT8

Synonyms: CDGF; CDGF1

Mammalian Cell None

Selection:

Vector:

pLenti-C-mGFP (PS100071)

Tag: mGFP

ACCN: NM_178155 **ORF Size:** 1725 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC223075).

•

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 178155.1

 RefSeq Size:
 3775 bp

 RefSeq ORF:
 1728 bp

 Locus ID:
 2530

 UniProt ID:
 Q9BYC5

Cytogenetics: 14q23.3

Protein Families: Transmembrane

Protein Pathways: Keratan sulfate biosynthesis, Metabolic pathways, N-Glycan biosynthesis





FUT8 (NM_178155) Human Tagged ORF Clone Lentiviral Particle - RC223075L2V

MW: 66.3 kDa

Gene Summary:

This gene encodes an enzyme belonging to the family of fucosyltransferases. The product of this gene catalyzes the transfer of fucose from GDP-fucose to N-linked type complex glycopeptides. This enzyme is distinct from other fucosyltransferases which catalyze alpha1-2, alpha1-3, and alpha1-4 fucose addition. The expression of this gene may contribute to the malignancy of cancer cells and to their invasive and metastatic capabilities. Alternative splicing results in multiple transcript variants. [provided by RefSeq, May 2011]