

## OriGene Technologies, Inc.

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## Product datasheet for RC201856L2V

## Aquaporin 3 (AQP3) (NM\_004925) Human Tagged ORF Clone Lentiviral Particle

## **Product data:**

Product Type:	Lentiviral Particles
Product Name:	Aquaporin 3 (AQP3) (NM_004925) Human Tagged ORF Clone Lentiviral Particle
Symbol:	Aquaporin 3
Synonyms:	AQP-3; GIL
Mammalian Cell Selection:	None
Vector:	pLenti-C-mGFP (PS100071)
Tag:	mGFP
ACCN:	NM_004925
ORF Size:	876 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC201856).
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. <u>More info</u>
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 004925.3</u>
RefSeq Size:	1882 bp
RefSeq ORF:	879 bp
Locus ID:	360
UniProt ID:	<u>Q92482</u>
Cytogenetics:	9p13.3
Domains:	MIP
Protein Families:	Druggable Genome, Transmembrane



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	Aquaporin 3 (AQP3) (NM_004925) Human Tagged ORF Clone Lentiviral Particle – RC201856L2V
MW:	31.5 kDa
Gene Summary:	This gene encodes the water channel protein aquaporin 3. Aquaporins are a family of small integral membrane proteins related to the major intrinsic protein, also known as aquaporin 0. Aquaporin 3 is localized at the basal lateral membranes of collecting duct cells in the kidney. In addition to its water channel function, aquaporin 3 has been found to facilitate the transport of nonionic small solutes such as urea and glycerol, but to a smaller degree. It has been suggested that water channels can be functionally heterogeneous and possess water and solute permeation mechanisms. Alternative splicing of this gene results in multiple transcript variants encoding different isoforms. [provided by RefSeq, Dec 2015]

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