

## Product datasheet for **RC210005L4V**

### AGL (NM\_000642) Human Tagged ORF Clone Lentiviral Particle

#### Product data:

|                           |  |
|---------------------------|--|
| Product Type:             | Lentiviral Particles   |
| Product Name:             | AGL (NM_000642) Human Tagged ORF Clone Lentiviral Particle   |
| Symbol:                   | AGL  |
| Synonyms:                 | GDE  |
| Mammalian Cell Selection: | Puromycin  |
| Vector:                   | pLenti-C-mGFP-P2A-Puro (PS100093)  |
| Tag:                      | mGFP   |
| ACCN:                     | NM_000642  |
| ORF Size:                 | 4596 bp  |
| ORF Nucleotide Sequence:  | The ORF insert of this clone is exactly the same as(RC210005).   |
| 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. <a href="#">More info</a> |
| 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:                   | <a href="#">NM_000642.2</a>  |
| RefSeq Size:              | 7371 bp  |
| RefSeq ORF:               | 4599 bp  |
| Locus ID:                 | 178  |
| UniProt ID:               | <a href="#">P35573</a>   |
| Cytogenetics:             | 1p21.2   |
| Protein Families:         | Druggable Genome   |
| Protein Pathways:         | Metabolic pathways, Starch and sucrose metabolism  |



[View online »](#)

**MW:** 174.6 kDa

**Gene Summary:** This gene encodes the glycogen debrancher enzyme which is involved in glycogen degradation. This enzyme has two independent catalytic activities which occur at different sites on the protein: a 4-alpha-glucotransferase activity and a amylo-1,6-glucosidase activity. Mutations in this gene are associated with glycogen storage disease although a wide range of enzymatic and clinical variability occurs which may be due to tissue-specific alternative splicing. Alternatively spliced transcripts encoding different isoforms have been described. [provided by RefSeq, Jul 2008]