

## Product datasheet for SC337021

### DGLUCY (NM\_001286470) Human Untagged Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	DGLUCY (NM_001286470) Human Untagged Clone
Tag:	Tag Free
Symbol:	DGLUCY
Synonyms:	C14orf159
Vector:	pCMV6-Entry (PS100001)
Fully Sequenced ORF:	>SC337021 representing NM_001286470. Blue=Insert sequence Red=Cloning site Green=Tag(s)

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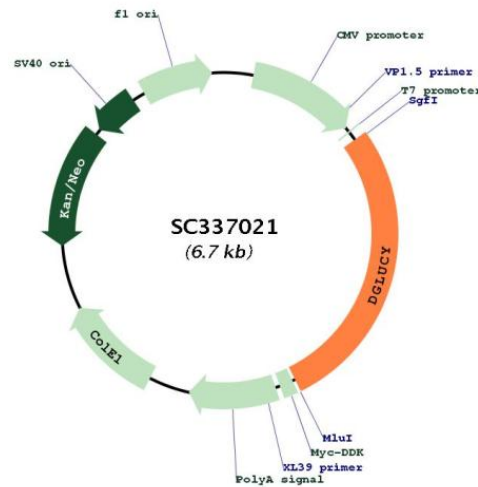
```



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Restriction Sites: SgfI-MluI

Plasmid Map:



ACCN: NM\_001286470

Insert Size: 1866 bp

**OTI Disclaimer:** Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).

**Components:** The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).

**Reconstitution Method:**

1. Centrifuge at 5,000xg for 5min.
2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.
3. Close the tube and incubate for 10 minutes at room temperature.
4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.
5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.

RefSeq: [NM\\_001286470.1](#)

RefSeq Size: 3476 bp

RefSeq ORF: 1866 bp

Locus ID: 80017

UniProt ID: [Q7Z3D6](#)

Cytogenetics: 14q32.11

MW: 66.9 kDa

**Gene Summary:**

D-glutamate cyclase that converts D-glutamate to 5-oxo-D-proline.[UniProtKB/Swiss-Prot Function]

Transcript Variant: This variant (6) differs in the 5' UTR and uses an alternate in-frame splice site, compared to variant 1. The encoded isoform (b) is longer than isoform a. Variants 4 and 6 encode the same isoform (b). Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.