

## Product datasheet for **SC332397**

### Giantin (GOLGB1) (NM\_001256488) Human Untagged Clone

#### Product data:

**Product Type:** Expression Plasmids  
**Product Name:** Giantin (GOLGB1) (NM\_001256488) Human Untagged Clone  
**Tag:** Tag Free  
**Symbol:** GOLGB1  
**Synonyms:** GCP; GCP372; GOLIM1  
**Vector:** pCMV6-Entry (PS10001)  
**Fully Sequenced ORF:** >SC332397 representing NM\_001256488.  
Blue=Insert sequence Red=Cloning site Green=Tag(s)

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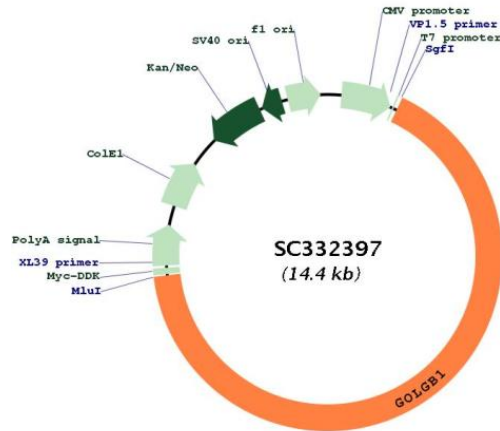
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```

**Restriction Sites:**

Sgfl-MluI

**Plasmid Map:**



**ACCN:**

NM\_001256488

**Insert Size:**

9555 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\\_001256488.1](#)

**RefSeq Size:**

11133 bp

**RefSeq ORF:**

9555 bp

<b>Locus ID:</b>	2804
<b>UniProt ID:</b>	<a href="#">Q14789</a>
<b>Cytogenetics:</b>	3q13.33
<b>Protein Families:</b>	Transcription Factors, Transmembrane
<b>MW:</b>	367.4 kDa
<b>Gene Summary:</b>	<p>May participate in forming intercisternal cross-bridges of the Golgi complex.[UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (4) uses an alternate splice site at its 5' end and initiates translation at a downstream start codon, compared to variant 1. It also lacks an internal in-frame exon and uses an alternate in-frame splice site in the 3' coding region compared to variant 1. The encoded isoform (4) is shorter than isoform 1.</p>