

# Product datasheet for TR517752

## Golga2 Mouse shRNA Plasmid (Locus ID 99412)

## **Product data:**

#### 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

Product Type:	shRNA Plasmids
Product Name:	Golga2 Mouse shRNA Plasmid (Locus ID 99412)
Locus ID:	99412
Synonyms:	AW555139; GM130; mKIAA4150
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	Golga2 - Mouse, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 99412). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	BC011407, NM 001080968, NM 133852, NM 133852.1, NM 133852.2, NM 001080968.1, BC024099, NM 001362695, NM 001362696, NM 001362697, NM 001362698
UniProt ID:	<u>Q921M4</u>



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### **ORIGENE** Golga2 Mouse shRNA Plasmid (Locus ID 99412) – TR517752

Summary:	Peripheral membrane component of the cis-Golgi stack that acts as a membrane skeleton
	that maintains the structure of the Golgi apparatus, and as a vesicle thether that facilitates
	vesicle fusion to the Golgi membrane (PubMed:28028212). Required for normal protein
	transport from the endoplasmic reticulum to the Golgi apparatus and the cell membrane
	(PubMed:28028212). Together with p115/USO1 and STX5, involved in vesicle tethering and
	fusion at the cis-Golgi membrane to maintain the stacked and inter-connected structure of
	the Golgi apparatus. Plays a central role in mitotic Golgi disassembly: phosphorylation at Ser-
	37 by CDK1 at the onset of mitosis inhibits the interaction with p115/USO1, preventing
	tethering of COPI vesicles and thereby inhibiting transport through the Golgi apparatus
	during mitosis. Also plays a key role in spindle pole assembly and centrosome organization
	(By similarity). Promotes the mitotic spindle pole assembly by activating the spindle assembly
	factor TPX2 to nucleate microtubules around the Golgi and capture them to couple mitotic
	membranes to the spindle: upon phosphorylation at the onset of mitosis, GOLGA2 interacts
	with importin-alpha via the nuclear localization signal region, leading to recruit importin-alpha
	to the Golgi membranes and liberate the spindle assembly factor TPX2 from importin-alpha.
	TPX2 then activates AURKA kinase and stimulates local microtubule nucleation. Upon filament
	assembly, nascent microtubules are further captured by GOLGA2, thus linking Golgi
	membranes to the spindle (By similarity). Regulates the meiotic spindle pole assembly,
	probably via the same mechanism (PubMed:21552007). Also regulates the centrosome
	organization (By similarity). Also required for the Golgi ribbon formation and glycosylation of
	membrane and secretory proteins (By similarity).[UniProtKB/Swiss-Prot Function]
shRNA Design:	These shRNA constructs were designed against multiple splice variants at this gene locus. To
	be certain that your variant of interest is targeted, please contact techsupport@origene.com.
	If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u> .

Performance Guaranteed: OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must be used in comparison with the target-specific shRNA transfected samples.

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).

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