

Product datasheet for PH321387

GOLGA3 (NM_005895) Human Mass Spec Standard

Product data:

Product Type:	Mass Spec Standards
Description:	GOLGA3 MS Standard C13 and N15-labeled recombinant protein (NP_005886)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC221387
Predicted MW:	167.2 kDa
Protein Sequence:	>RC221387 representing NM_005895 Red=Cloning site Green=Tags(s)

MDGASAEQDGLQEDRSHSGPSSSLPEAPLKPPGPLVPPDQDQKQVQCAEVNRASSTEGESPDGPGQGGLCQNG
PTPPFPDPPSSLDPTTSPVGPDA SPGVAGFHDNLRKSSQGTSAEGSVRKEALQSLRLSLPMQETQLCSTDS
PLPLEKEEQVRLQARKWLEEQKQYRVKRQQRSSQPATKTRLFSTLDPELMLNPNENLPRASTLAMTKEY
SFLRTSVPRGPKVGSGLPAHPREKTKSSKIRSLADYRTEDSNAGNSGGNVPAPDSTKGSCLKQRSSA
ASVVSEISLSPDTRDRENTSLAGDSVSEVDGNDSDSSSYSSASTRGTYGILSKTVGTQDTPYMVNGQEI
PADTLGQFSPSIKDVLAQAAAAHQDQGEVNGEVRSRRDSCSSVLESSAAETQEEMLVKKEKMRLEGQ
LEALSLEASQALKEKAEALQAQLAALSTKLQAQVECSHSSQQRQDSLSEVDTLKQSCWDLERAMTDLQNM
LEAKNASLASSNNDLQVAEEQYQRLMAKVEDMQRSMLSKDNTVHDLRQQMTALQSQQLQVQLERTTLTTSK
LKASQAEISSLQSVRQWYQQQLALAQAARVRLQGEMAHIQVGMQTAQAGLLEHLKLENVLSQQLTETQHR
SMKEKGRIAAQLQGIADMLDQEA AFMQIQEAKTMVEEDLQRRLEEFEGEGERERLQRMADSAASLEQLEQ
VKLTLQRDQQLQEQEHLDMKQLTLTQEALQSREQLDALQTHYDELQARLGELQGEAASREDTICL
LQNEKIILEAALQAAKSGKEELDRGARRLEEGTEETSETLEKLEELA IKSQVVEHLQEQETAALKKQMQK
IKEQFLQKVMVEAYRRDATSKDQLISELKATRRLDSELKELRQELMQVHGEKRTAEAE LSRHREVAQ
VRQHMADLEGLHLSAQKERDEMETHLQSLQFDKEQMVAVTEANEALKKQIEELQEQEARKAITEQKQKMR
LGSDLTSAQKEMKTKHKAYENAVGILSRRLQEALAAKEAADAELGQLRAQGGSSDSSLALHERIQALEAE
LQAVSHSKTLLKELQEVIALTSQEL EESREKVLLEDELEQESRGFRKKIKRLEESNKKLALAELEHEK GK
LTGLGQSNAALREHNSILETALAKREADLVQLNLQVQAVLQRKEEEDRQMKHLVQALQASLEKEKEKVN
LKEQVAAAKVEAGHNRHFKAASLELSEVKKELQAKEHLVQKLQAEADDLQIREGKHSQEIAQFQAEALAE
ARAQLQLLQKQLDEQLSKQPVGQEMENLQWEVDQKEREIQSLKQQLDLTEQQGRKELEGLQQLLQNVKS
ELEMAQEDLSMTQKDFMLQAKVSELKNNMKTLLQQNQQLKLDLRRGAATRKEPKGEASSSNPATPIKI
PDCVPVPSLLELLRPPPAVSKEPLKNLNSCLQQLKQEMDSLQRQMEHALTVHESLSSWTPLEPATASP
VPPGGHAGPRGDPQRHSQSRASKEGPGE

TRTRPLEQKLI SEEDLAANDILDYKDDDDKV

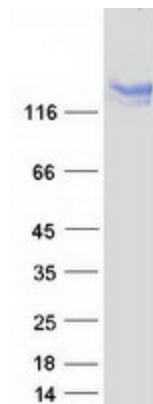
Tag:	C-Myc/DDK
Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining



View online >

Concentration:	>0.05 µg/µL as determined by microplate BCA method
Labeling Method:	Labeled with [U- ¹³ C ₆ , ¹⁵ N ₄]-L-Arginine and [U- ¹³ C ₆ , ¹⁵ N ₂]-L-Lysine
Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3
Storage:	Store at -80°C. Avoid repeated freeze-thaw cycles.
Stability:	Stable for 3 months from receipt of products under proper storage and handling conditions.
RefSeq:	NP_005886
RefSeq Size:	6956
RefSeq ORF:	4494
Synonyms:	GCP170; MEA-2
Locus ID:	2802
UniProt ID:	Q08378
Cytogenetics:	12q24.33
Summary:	The Golgi apparatus, which participates in glycosylation and transport of proteins and lipids in the secretory pathway, consists of a series of stacked cisternae (flattened membrane sacs). Interactions between the Golgi and microtubules are thought to be important for the reorganization of the Golgi after it fragments during mitosis. This gene encodes a member of the golgin family of proteins which are localized to the Golgi. Its encoded protein has been postulated to play a role in nuclear transport and Golgi apparatus localization. Several alternatively spliced transcript variants that encode different protein isoforms have been described for this gene. [provided by RefSeq, Feb 2010]

Product images:



Coomassie blue staining of purified GOLGA3 protein (Cat# [TP321387]). The protein was produced from HEK293T cells transfected with GOLGA3 cDNA clone (Cat# [RC221387]) using MegaTran 2.0 (Cat# [TT210002]).