

Product datasheet for PH311607

COG5 (NM_006348) Human Mass Spec Standard

Product data:

Product Type:	Mass Spec Standards
Description:	COG5 MS Standard C13 and N15-labeled recombinant protein (NP_006339)
Species:	Human
Expression Host:	HEK293
Expression cDNA Clone or AA Sequence:	RC211607
Predicted MW:	94.7 kDa
Protein Sequence:	>RC211607 representing NM_006348 Red=Cloning site Green=Tags(s)

MGWVGGRRRDSASPPGRSRSAAADDINPAPANMEGGGSVAVAGLGARGSGAAAATVRELLQDGCYSDFLN
EDFDVKTYTSQSIHQAVIAEQLAKLAQGISQLDRELHLQVVARHEDLLAQTGIESLEGVLQMMQTRIGA
LQGAVDRIKAKIVEPYNKIVARTAQLARLQVACDLLRRIIRILNLSKRLQGQLQGGGREITKAAQSLNEL
DYL SQGIDL SGIEVIENDLLFIARARLEVENQAKRLLEQGLETQNPQTQVGTALQVFYNLGLTKDITISVV
DGYCATLEENINSALDIKVLTPSQSAVRGGPGRSTMPPTGNTAALRASLWLNMEKLMDHIAVCGQVQH
LQKVLAKKRDVPVSHICFIEEIVKDGQPEIFYTFWNSVTQALSSQFHMATNSSMFLKQAFEGEYPKLLRLY
NDLWKRLQYQSQHIQGNFNASGTTDL YVDLQHMEDDAQDIFIPKKPDYDPEKALKDSLQPYEAAYLSKSL
SRLFDPINLVFPPGGRNPPSSDEL DGI IKTIASELNVAAVDTNL TLAVSKNVAKTIQLYSVKSEQLLSTQ
GDASQVIGPLTEGQRRNVAVVNSLYKHLQSVTKVVSSQSFPAAEQTIISALKATHALMENAVQPLLTS
VGDAIEAIIITMHQEDFSGSLSSSGKPDVPCSLYMKELQGFARVMSDYFKHFECDFVFNTEAIAQRA
VELFIRHASLIRPLGEGGKMLAADFQAQMEAVGPFRRVSDLGKSYRMLRSFRPLLQASEHVASSPAL
GDVIFPFSIIIQFLFTRAPAEKSPFQRAEWSHTRFSQWLDDHPSEKDRLLLIRGALEYVQSVRSREGKE
FAPVYPI MVQLLQKAMSALQ

TRTRPLEQKLI SEEDLAANDILDYKDDDDKV

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

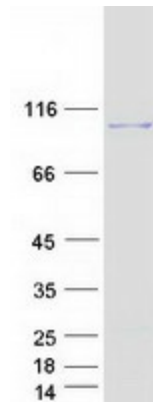


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RefSeq:	NP_006339
RefSeq Size:	3604
RefSeq ORF:	2580
Synonyms:	CDG2I; GOLTC1; GTC90
Locus ID:	10466
UniProt ID:	Q9UP83
Cytogenetics:	7q22.3

Summary: The protein encoded by this gene is one of eight proteins (Cog1-8) which form a Golgi-localized complex (COG) required for normal Golgi morphology and function. The encoded protein is organized with conserved oligomeric Golgi complex components 6, 7 and 8 into a sub-complex referred to as lobe B. Alternative splicing results in multiple transcript variants. Mutations in this gene result in congenital disorder of glycosylation type 2I.[provided by RefSeq, Jan 2011]

Product images:



Coomassie blue staining of purified COG5 protein (Cat# [TP311607]). The protein was produced from HEK293T cells transfected with COG5 cDNA clone (Cat# [RC211607]) using MegaTran 2.0 (Cat# [TT210002]).