

# **Product datasheet for PH319271**

## OriGene Technologies, Inc.

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### TOM1L2 (NM\_001082968) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** TOM1L2 MS Standard C13 and N15-labeled recombinant protein (NP\_001076437)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC219271

or AA Sequence: Predicted MW:

55.4 kDa

Protein Sequence: >RC219271 representing NM\_001082968

Red=Cloning site Green=Tags(s)

MEFLLGNPFSTPVGQCLEKATDGSLQSEDWTLNMEICDIINETEEGPKDAIRALKKRLNGNRNYREVMLA LTVLETCVKNCGHRFHILVANRDFIDSVLVKIISPKNNPPTIVQDKVLALIQAWADAFRSSPDLTGVVHI YEELKRKGVEFPMADLDALSPIHTPQRSVPEVDPAATMPRSQSQQRTSAGSYSSPPPAPYSAPQAPALSV TGPITANSEQIARLRSELDVVRGNTKVMSEMLTEMVPGQEDSSDLELLQELNRTCRAMQQRIVELISRVS NEEVTEELLHVNDDLNNVFLRYERFERYRSGRSVQNASNGVLNEVTEDNLIDLGPGSPAVVSPMVGNTAP PSSLSSQLAGLDLGTESVSGTLSSLQQCNPRDGFDMFAQTRGNSLAEQRKTVTYEDPQAVGGLASALDNR KQSSEGIPVAQPSVMDDIEVWLRTDLKGDDLEEGVTSEEFDKFLEERAKAAEMVPDLPSPPMEAPAPASN

PSGRKKPERSEDALFAL

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

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- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-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 001076437

RefSeq Size: 5825 RefSeq ORF: 1521



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 Locus ID:
 146691

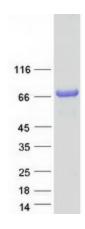
 UniProt ID:
 Q6ZVM7

 Cytogenetics:
 17p11.2

Summary: This gene belongs to a small gene family whose members have an N-terminal VHS domain

followed by a GAT domain; domains which typically participate in vesicular trafficking. The canonical protein encoded by this gene also has a C-terminal clathrin binding motif. This protein has been shown to interact with Tollip, clathrin and ubiquitin and is thought to play a role in endosomal sorting. This gene resides in the 3.7 Mb deletion of chromosome region 17p11.2 that is associated with Smith-Magenis syndrome. Alternative splicing results in multiple transcript variants encoding distinct proteins. [provided by RefSeq, Apr 2017]

### **Product images:**



Coomassie blue staining of purified TOM1L2 protein (Cat# [TP319271]). The protein was produced from HEK293T cells transfected with TOM1L2 cDNA clone (Cat# [RC219271]) using MegaTran 2.0 (Cat# [TT210002]).