

Product datasheet for PH304770

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

TTC35 (EMC2) (NM_014673) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

Description: TTC35 MS Standard C13 and N15-labeled recombinant protein (NP_055488)

Species:HumanExpression Host:HEK293

Expression cDNA Clone

RC204770

or AA Sequence: Predicted MW:

34.8 kDa

Protein Sequence: >RC204770 protein sequence

Red=Cloning site Green=Tags(s)

MAKVSELYDVTWEEMRDKMRKWREENSRNSEQIVEVGEELINEYASKLGDDIWIIYEQVMIAALDYGRDD LALFCLQELRRQFPGSHRVKRLTGMRFEAMERYDDAIQLYDRILQEDPTNTAARKRKIAIRKAQGKNVEA IRELNEYLEQFVGDQEAWHELAELYINEHDYAKAAFCLEELMMTNPHNHLYCQQYAEVKYTQGGLENLEL SRKYFAQALKLNNRNMRALFGLYMSASHIASNPKASAKTKKDNMKYASWAASQINRAYQFAGRSKKETKY

SLKAVEDMLETLQITQS

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 055488

RefSeq Size: 1268 RefSeq ORF: 891

Synonyms: KIAA0103; TTC35

Locus ID: 9694



TTC35 (EMC2) (NM_014673) Human Mass Spec Standard - PH304770

UniProt ID: Q15006

Cytogenetics: 8q23.1

Summary: Part of the endoplasmic reticulum membrane protein complex (EMC) that enables the

energy-independent insertion into endoplasmic reticulum membranes of newly synthesized

membrane proteins (PubMed:30415835, PubMed:29809151, PubMed:29242231, PubMed:32459176, PubMed:32439656). Preferentially accommodates proteins with

transmembrane domains that are weakly hydrophobic or contain destabilizing features such

as charged and aromatic residues (PubMed:30415835, PubMed:29809151,

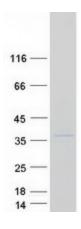
PubMed:29242231). Involved in the cotranslational insertion of multi-pass membrane proteins in which stop-transfer membrane-anchor sequences become ER membrane spanning helices (PubMed:30415835, PubMed:29809151). It is also required for the post-translational insertion of tail-anchored/TA proteins in endoplasmic reticulum membranes (PubMed:29809151, PubMed:29242231). By mediating the proper cotranslational insertion of N-terminal transmembrane domains in an N-exo topology, with translocated N-terminus in the lumen of the ER, controls the topology of multi-pass membrane proteins like the G protein-coupled receptors (PubMed:30415835). By regulating the insertion of various proteins in membranes, it is indirectly involved in many collular processes (Probable)

in membranes, it is indirectly involved in many cellular processes (Probable).

[UniProtKB/Swiss-Prot Function]

Protein Families: Protease

Product images:



Coomassie blue staining of purified EMC2 protein (Cat# [TP304770]). The protein was produced from HEK293T cells transfected with EMC2 cDNA clone (Cat# [RC204770]) using MegaTran 2.0 (Cat# [TT210002]).