

# **Product datasheet for PH314027**

## OriGene Technologies, Inc.

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#### TBCE (NM 001079515) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** TBCE MS Standard C13 and N15-labeled recombinant protein (NP\_001072983)

Species: Human **HEK293 Expression Host: Expression cDNA Clone** 

or AA Sequence:

RC214027

Predicted MW: 59.3 kDa

>RC214027 protein sequence **Protein Sequence:** 

Red=Cloning site Green=Tags(s)

MSDTLTADVIGRRVEVNGEHATVRFAGVVPPVAGPWLGVEWDNPERGKHDGSHEGTVYFKCRHPTGGSFI RPNKVNFGTDFLTAIKNRYVLEDGPEEDRKEQIVTIGNKPVETIGFDSIMKQQSQLSKLQEVSLRNCAVS CAGEKGGVAEACPNIRKVDLSKNLLSSWDEVIHIADQLRHLEVLNVSENKLKFPSGSVLTGTLSVLKVLV LNQTGITWAEVLRCVAGCPGLEELYLESNNIFISERPTDVLQTVKLLDLSSNQLIDENQLYLIAHLPRLE QLILSDTGISSLHFPDAGIGCKTSMFPSLKYLVVNDNQISQWSFFNELEKLPSLRALSCLRNPLTKEDKE AETARLLIIASIGQLKTLNKCEILPEERRRAELDYRKAFGNEWKQAGGHKDPEKNRLSEEFLTAHPRYOF LCLKYGAPEDWELKTQQPLMLKNQLLTLKIKYPHQLDQKVLEKQLPGSMTIQKVKGLLSRLLKVPVSDLL

LSYESPKKPGREIELENDLKSLQFYSVENGDCLLVRW

TRTRPLEQKLISEEDLAANDILDYKDDDDKV

C-Myc/DDK Tag:

**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

Store at -80°C. Avoid repeated freeze-thaw cycles. Storage:

Stable for 3 months from receipt of products under proper storage and handling conditions. Stability:

RefSeq: NP 001072983

RefSeq Size: 2025 RefSeq ORF: 1581



### TBCE (NM\_001079515) Human Mass Spec Standard - PH314027

**Synonyms:** HRD; KCS; KCS1; pac2; PEAMO

 Locus ID:
 6905

 UniProt ID:
 Q15813

 Cytogenetics:
 1q42.3

Summary: Cofactor E is one of four proteins (cofactors A, D, E, and C) involved in the pathway leading to

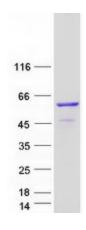
correctly folded beta-tubulin from folding intermediates. Cofactors A and D are believed to play a role in capturing and stabilizing beta-tubulin intermediates in a quasi-native confirmation. Cofactor E binds to the cofactor D/beta-tubulin complex; interaction with cofactor C then causes the release of beta-tubulin polypeptides that are committed to the

native state. Two transcript variants encoding the same protein have been found for this

gene. [provided by RefSeq, Jul 2008]

**Protein Families:** Druggable Genome

## **Product images:**



Coomassie blue staining of purified TBCE protein (Cat# [TP314027]). The protein was produced from HEK293T cells transfected with TBCE cDNA clone (Cat# [RC214027]) using MegaTran 2.0 (Cat# [TT210002]).

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