

## **Product datasheet for PH305570**

## OriGene Technologies, Inc.

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## CT45A5 (NM 001007551) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** CT45A5 MS Standard C13 and N15-labeled recombinant protein (NP\_001007552)

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

or AA Sequence:

RC205570

Predicted MW: 21.3 kDa

>RC205570 protein sequence **Protein Sequence:** 

Red=Cloning site Green=Tags(s)

MTDKTEKVAVDPETVFKRPRECDSPSYQKRQRMALLARKQGAGDSLIAGSAMSKEKKLMTGHAIPPSQLD SQIDDFTGFSKDRMMQKPGSNAPVGGNVTSSFSGDDLECRETASSPKSQREINADIKRKLVKELRCVGQK

YEKIFEMLEGVQGPTAVRKRFFESIIKEAARCMRRDFVKHLKKKLKRMI

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

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

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

RefSeq: NP 001007552

RefSeg Size: 1321 RefSeq ORF: 567

Synonyms: CT45-3; CT45-4; CT45-6; CT45.5; CT45A3; CT45A4; CT45A6; CT45A7; CT455

Locus ID: 441521

UniProt ID: Q6NSH3, P0DMU8, P0DMU7, P0DMV0





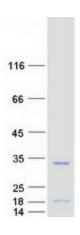
Cytogenetics:

Xq26.3

**Summary:** 

This gene represents one of a cluster of several similar genes located on the q arm of chromosome X. The genes in this cluster encode members of the cancer/testis (CT) family of antigens, and are distinct from other CT antigens. These antigens are thought to be novel therapeutic targets for human cancers. Alternative splicing results in multiple transcript variants. [provided by RefSeq, Apr 2014]

## **Product images:**



Coomassie blue staining of purified CT45A5 protein (Cat# [TP305570]). The protein was produced from HEK293T cells transfected with CT45A5 cDNA clone (Cat# [RC205570]) using MegaTran 2.0 (Cat# [TT210002]).