

## **Product datasheet for PH301998**

## OriGene Technologies, Inc.

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## DBT (NM 001918) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** DBT MS Standard C13 and N15-labeled recombinant protein (NP\_001909)

Species: Human **HEK293 Expression Host:** 

**Expression cDNA Clone** 

RC201998

or AA Sequence: Predicted MW:

53.5 kDa

>RC201998 protein sequence **Protein Sequence:** 

Red=Cloning site Green=Tags(s)

MAAVRMLRTWSRNAGKLICVRYFQTCGNVHVLKPNYVCFFGYPSFKYSHPHHFLKTTAALRGQVVQFKLS DIGEGIREVTVKEWYVKEGDTVSQFDSICEVQSDKASVTITSRYDGVIKKLYYNLDDIAYVGKPLVDIET EALKDSEEDVVETPAVSHDEHTHQEIKGRKTLATPAVRRLAMENNIKLSEVVGSGKDGRILKEDILNYLE KQTGAILPPSPKVEIMPPPPKPKDMTVPILVSKPPVFTGKDKTEPIKGFQKAMVKTMSAALKIPHFGYCD EIDLTELVKLREELKPIAFARGIKLSFMPFFLKAASLGLLQFPILNASVDENCQNITYKASHNIGIAMDT EQGLIVPNVKNVQICSIFDIATELNRLQKLGSVGQLSTTDLTGGTFTLSNIGSIGGTFAKPVIMPPEVAI GALGSIKAIPRFNQKGEVYKAQIMNVSWSADHRVIDGATMSRFSNLWKSYLENPAFMLLDLK

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 001909

RefSeg Size: 10831 RefSeq ORF: 1446

BCATE2; BCKAD-E2; BCKADE2; BCKDH-E2; BCOADC-E2; E2; E2B Synonyms:





**Locus ID:** 1629

UniProt ID: P11182
Cytogenetics: 1p21.2

Summary: The branched-chain alpha-keto acid dehydrogenase complex (BCKD) is an inner-

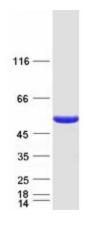
mitochondrial enzyme complex involved in the breakdown of the branched-chain amino acids isoleucine, leucine, and valine. The BCKD complex is thought to be composed of a core of 24 transacylase (E2) subunits, and associated decarboxylase (E1), dehydrogenase (E3), and regulatory subunits. This gene encodes the transacylase (E2) subunit. Mutations in this gene result in maple syrup urine disease, type 2. Alternatively spliced transcript variants have been described, but their biological validity has not been determined. [provided by RefSeq, Jul

2008]

**Protein Families:** Druggable Genome

**Protein Pathways:** Metabolic pathways, Valine, leucine and isoleucine degradation

## **Product images:**



Coomassie blue staining of purified DBT protein (Cat# [TP301998]). The protein was produced from HEK293T cells transfected with DBT cDNA clone (Cat# [RC201998]) using MegaTran 2.0 (Cat# [TT210002]).