

# **Product datasheet for PH325667**

### OriGene Technologies, Inc.

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## BAAT (NM 001127610) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** BAAT MS Standard C13 and N15-labeled recombinant protein (NP 001121082)

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

or AA Sequence:

RC225667

Predicted MW:

46.3 kDa

>RC225667 protein sequence **Protein Sequence:** 

Red=Cloning site Green=Tags(s)

MIQLTATPVSALVDEPVHIQATGLIPFQMVSFQASLEDENGDMFYSQAHYRANEFGEVDLNHASSLGGDY MGVHPMGLFWSLKPEKLLTRLLKRDVMNRPFQVQVKLYDLELIVNNKVASAPKASLTLERWYVAPGVTRI KVREGRLRGALFLPPGEGLFPGVIDLFGGLGGLLEFRASLLASRGFASLALAYHNYEDLPRKPEVTDLEY FEEAANFLLRHPKVFGSGVGVVSVCQGVQIGLSMAIYLKQVTATVLINGTNFPFGIPQVYHGQIHQPLPH SAQLISTNALGLLELYRTFETTQVGASQYLFPIEEAQGQFLFIVGEGDKTINSKAHAEQAIGQLKRHGKN NWTLLSYPGAGHLIEPPYSPLCCASTTHDLRLHWGGEVIPHAAAQEHAWKEIQRFLRKHLIPDVTSQL

**TRTRPL**EQKLISEEDLAANDILDYKDDDDK**V** 

Tag: C-Myc/DDK

**Purity:** > 80% as determined by SDS-PAGE and Coomassie blue staining

>0.05 µg/µL as determined by microplate BCA method **Concentration:** 

**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 001121082

RefSeg Size: 3377 RefSeq ORF: 1254

Synonyms: BACAT; BACD1; BAT; HCHO

Locus ID: 570



### BAAT (NM\_001127610) Human Mass Spec Standard - PH325667

 UniProt ID:
 Q14032

 Cytogenetics:
 9q31.1

Summary: The protein encoded by this gene is a liver enzyme that catalyzes the transfer of C24 bile

acids from the acyl-CoA thioester to either glycine or taurine, the second step in the

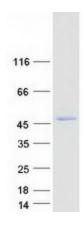
formation of bile acid-amino acid conjugates. The bile acid conjugates then act as a detergent in the gastrointestinal tract, which enhances lipid and fat-soluble vitamin absorption. Defects in this gene are a cause of familial hypercholanemia (FHCA). Two transcript variants encoding

the same protein have been found for this gene. [provided by RefSeq, Jul 2008]

**Protein Pathways:** Biosynthesis of unsaturated fatty acids, Metabolic pathways, Primary bile acid biosynthesis,

Taurine and hypotaurine metabolism

## **Product images:**



Coomassie blue staining of purified BAAT protein (Cat# [TP325667]). The protein was produced from HEK293T cells transfected with BAAT cDNA clone (Cat# [RC225667]) using MegaTran 2.0 (Cat# [TT210002]).