

# **Product datasheet for PH304854**

### OriGene Technologies, Inc.

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## BRDG 1 (STAP1) (NM 012108) Human Mass Spec Standard

**Product data:** 

**Product Type:** Mass Spec Standards

**Description:** STAP1 MS Standard C13 and N15-labeled recombinant protein (NP\_036240)

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

or AA Sequence:

RC204854

Predicted MW: 34.3 kDa

>RC204854 protein sequence **Protein Sequence:** 

Red=Cloning site Green=Tags(s)

MMAKKPPKPAPRRIFQERLKITALPLYFEGFLLIKRSGYREYEHYWTELRGTTLFFYTDKKSIIYVDKLD IVDLTCLTEQNSTEKNCAKFTLVLPKEEVQLKTENTESGEEWRGFILTVTELSVPQNVSLLPGQVIKLHE VLEREKKRRIETEQSTSVEKEKEPTEDYVDVLNPMPACFYTVSRKEATEMLQKNPSLGNMILRPGSDSRN YSITIRQEIDIPRIKHYKVMSVGQNYTIELEKPVTLPNLFSVIDYFVKETRGNLRPFICSTDENTGQEPS

MEGRSEKLKKNPHIA

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:

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

NP 036240 RefSeq:

RefSeq Size: 1524 RefSeq ORF: 885

BRDG1; STAP-1 Synonyms:

Locus ID: 26228



#### BRDG 1 (STAP1) (NM\_012108) Human Mass Spec Standard - PH304854

UniProt ID: Q9ULZ2, A0A024RD91

**Cytogenetics:** 4q13.2

**Summary:** The protein encoded by this gene contains a proline-rich region, a pleckstrin homology (PH)

domain, and a region in the carboxy terminal half with similarity to the Src Homology 2 (SH2) domain. This protein is a substrate of tyrosine-protein kinase Tec, and its interaction with tyrosine-protein kinase Tec is phosphorylation-dependent. This protein is thought to participate in a positive feedback loop by upregulating the activity of tyrosine-protein kinase

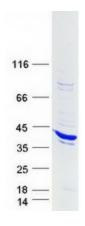
Tec. Variants of this gene have been associated with autosomal-dominant

hypercholesterolemia (ADH), which is characterized by elevated low-density lipoprotein cholesterol levels and in increased risk of coronary vascular disease. Alternative splicing

results in multiple transcript variants. [provided by RefSeq, Dec 2015]

**Protein Families:** Druggable Genome

# **Product images:**



Coomassie blue staining of purified STAP1 protein (Cat# [TP304854]). The protein was produced from HEK293T cells transfected with STAP1 cDNA clone (Cat# [RC204854]) using MegaTran 2.0 (Cat# [TT210002]).