

## **Product datasheet for PH316974**

# OriGene Technologies, Inc.

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### ACSL6 (NM 015256) Human Mass Spec Standard

#### **Product data:**

**Product Type:** Mass Spec Standards

**Description:** ACSL6 MS Standard C13 and N15-labeled recombinant protein (NP 056071)

Species: Human
Expression Host: HEK293

**Expression cDNA Clone** 

RC216974

or AA Sequence: Predicted MW:

80.3 kDa

Protein Sequence: RC216974

Tag: C-Myc/DDK

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

Concentration:  $>0.05 \mu g/\mu L$  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

**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 056071

RefSeq Size: 3047 RefSeq ORF: 2166

Synonyms: ACS2; FACL6; LACS2; LACS5; LACS 6

**Locus ID:** 23305

UniProt ID: Q9UKU0, B4DFW3

**Cytogenetics:** 5q31.1



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**Summary:** The protein encoded by this gene catalyzes the formation of acyl-CoA from fatty acids, ATP,

and CoA, using magnesium as a cofactor. The encoded protein plays a major role in fatty acid metabolism in the brain. Translocations with the ETV6 gene are causes of myelodysplastic syndrome with basophilia, acute myelogenous leukemia with eosinophilia, and acute eosinophilic leukemia. Several transcript variants encoding different isoforms have been

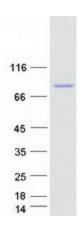
found for this gene.[provided by RefSeq, Apr 2011]

**Protein Families:** Druggable Genome, Transmembrane

**Protein Pathways:** Adipocytokine signaling pathway, Fatty acid metabolism, Metabolic pathways, PPAR signaling

pathway

## **Product images:**



Coomassie blue staining of purified ACSL6 protein (Cat# [TP316974]). The protein was produced from HEK293T cells transfected with ACSL6 cDNA clone (Cat# [RC216974]) using MegaTran 2.0 (Cat# [TT210002]).