

## **Anti-BAAT** antibody



**Description** Unconjugated Rabbit polyclonal to BAAT

Model STJ190610

**Host** Rabbit

**Reactivity** Human

**Applications** ELISA, WB

**Gene ID** <u>570</u>

Gene Symbol BAAT

**Dilution range** WB 1:500-2000 ELISA 1:5000-20000

**Specificity** BAAT Polyclonal Antibody detects endogenous levels of protein.

**Tissue Specificity** Expressed in liver, gallbladder mucosa and pancreas.

**Purification** BAAT antibody was affinity-purified from rabbit antiserum by affinity-

chromatography using epitope-specific immunogen.

**Note** For Research Use Only (RUO).

Protein Name

Bile acid-CoA:amino acid N-acyltransferase BACAT BAT Glycine N-

choloyltransferase Long-chain fatty-acyl-CoA hydrolase

Molecular Weight 45 kDa

**Clonality** Polyclonal

**Conjugation** Unconjugated

**Isotype** IgG

**Formulation** Liquid form in PBS containing 50% glycerol, and 0.02% sodium azide.

**Concentration** 1 mg/ml

**Storage Instruction** Store at -20°C, and avoid repeat freeze-thaw cycles.

Database Links <u>HGNC:932OMIM:602938</u>

Alternative Names Bile acid-CoA:amino acid N-acyltransferase BACAT BAT Glycine N-

choloyltransferase Long-chain fatty-acyl-CoA hydrolase

**Function** Involved in bile acid metabolism. In liver hepatocytes catalyzes the second

step in the conjugation of C24 bile acids (choloneates) to glycine and taurine before excretion into bile canaliculi. The major components of bile are cholic acid and chenodeoxycholic acid. In a first step the bile acids are converted to an acyl-CoA thioester, either in peroxisomes (primary bile acids deriving from

the cholesterol pathway), or cytoplasmic at the endoplasmic reticulum

(secondary bile acids). May catalyze the conjugation of primary or secondary bile acids, or both. The conjugation increases the detergent properties of bile

acids in the intestine, which facilitates lipid and fat-soluble vitamin

absorption. In turn, bile acids are deconjugated by bacteria in the intestine and are recycled back to the liver for reconjugation (secondary bile acids). May also act as an acyl-CoA thioesterase that regulates intracellular levels of free fatty acids. In vitro, catalyzes the hydrolysis of long- and very long-chain saturated acyl-CoAs to the free fatty acid and coenzyme A (CoASH), and

conjugates glycine to these acyl-CoAs.

**Cellular Localization** Cytoplasm

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