

## FABP6 Human, His

**Description:** FABP6 Human Recombinant produced in E.Coli is a single, non-glycosylated polypeptide chain containing 128 amino acids and having a molecular mass of 18 kDa. FABP6 is fused to His tag at N-terminus and purified by standard chromatography techniques.

**Catalog #:** PRPS-674

For research use only.

**Synonyms:** I-BABP, ILBP, I-15P, I-BAP, ILBP3, ILLBP, I-BABP, I-BALB, FABP-6, Gastrotropin, Ileal lipid-binding protein, Intestinal 15 kDa protein, Intestinal bile acid-binding protein, Fatty acid-binding protein 6, FABP6.

**Source:** Escherichia Coli.

**Physical Appearance:** Sterile Filtered colorless liquid formulation.

**Purity:** Greater than 95% as determined by SDS-PAGE.

**Formulation:**

FABP6 His-Tag is supplied in 20mM Tris HCL pH=8, 0.5mM DTT and 50% glycerol.

**Stability:**

Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. Please avoid freeze thaw cycles.

**Usage:**

NeoBiolab's products are furnished for LABORATORY RESEARCH USE ONLY. The product may not be used as drugs, agricultural or pesticidal products, food additives or household chemicals.

**Introduction:**

FABP6 also called ileal fatty acid binding protein, is part of the small family of highly conserved, cytoplasmic proteins that bind long-chain fatty acids and other hydrophobic ligands. FABP6 cytosolic protein binds bile acid. FABP6 plays a role in fatty acid uptake, transport, and metabolism. FABP6 stimulates gastric acid and pepsinogen secretion. Seems to be able to bind to bile salts and bilirubins. FABP6 expression is restricted in the small intestine to the ileum where it is involved in the enterohepatic circulation of bile acids. Alternate transcription promoters generate 2 transcript variants, encoding a 128 aa and a 177 aa residue protein. Human FABP6 isoform 2 contains 128 amino acid residues and is acetylated on Ala2. FABP6 binds together fatty acids and bile acids and is directly involved in fatty acid transport and metabolism.

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