

# CPTC-FBN1-1 (CAB080202)

**Uniprot ID:** P35555

**Protein name:** FBN1\_HUMAN

**Full name:** Fibrillin-1

**Function:** [Fibrillin-1]: Structural component of the 10-12 nm diameter microfibrils of the extracellular matrix, which conveys both structural and regulatory properties to load-bearing connective tissues (PubMed:1860873, PubMed:15062093). Fibrillin-1-containing microfibrils provide long-term force bearing structural support (PubMed:27026396). In tissues such as the lung, blood vessels and skin, microfibrils form the periphery of the elastic fiber, acting as a scaffold for the deposition of elastin (PubMed:27026396). In addition, microfibrils can occur as elastin-independent networks in tissues such as the ciliary zonule, tendon, cornea and glomerulus where they provide tensile strength and have anchoring roles (PubMed:27026396). Fibrillin-1 also plays a key role in tissue homeostasis through specific interactions with growth factors, such as the bone morphogenetic proteins (BMPs), growth and differentiation factors (GDFs) and latent transforming growth factor-beta-binding proteins (LTBPs), cell-surface integrins and other extracellular matrix protein and proteoglycan components (PubMed:27026396). Regulates osteoblast maturation by controlling TGF-beta bioavailability and calibrating TGF-beta and BMP levels, respectively (By similarity). Negatively regulates osteoclastogenesis by binding and sequestering an osteoclast differentiation and activation factor TNFSF11 (PubMed:24039232). This leads to disruption of TNFSF11-induced Ca(2+) signaling and impairment of TNFSF11-mediated nuclear translocation and activation of transcription factor NFATC1 which regulates genes important for osteoclast differentiation and function (PubMed:24039232). Mediates cell adhesion via its binding to cell surface receptors integrins ITGAV:ITGB3 and ITGA5:ITGB1 (PubMed:12807887, PubMed:17158881). Binds heparin and this interaction has an important role in the assembly of microfibrils (PubMed:11461921). [Asprosin]: Adipokine secreted by white adipose tissue that plays an important regulatory role in the glucose metabolism of liver, muscle and pancreas (PubMed:27087445, PubMed:30853600). Hormone that targets the liver in response to fasting to increase plasma glucose levels (PubMed:27087445). Binds the olfactory receptor OR4M1 at the surface of hepatocytes and promotes hepatocyte glucose release by activating the protein kinase A activity in the liver, resulting in rapid glucose release into the circulation (PubMed:27087445, PubMed:31230984). May act as a regulator of adaptive thermogenesis by inhibiting browning and energy consumption, while increasing lipid deposition in white adipose tissue (By similarity). Also acts as an orexigenic hormone that increases appetite: crosses the blood brain barrier and exerts effects on the hypothalamus (By similarity). In the arcuate nucleus of the hypothalamus, asprosin directly activates orexigenic AgRP neurons and indirectly inhibits anorexigenic POMC neurons, resulting in appetite stimulation (By similarity). Activates orexigenic AgRP neurons via binding to the olfactory receptor OR4M1 (By similarity). May also play a role in sperm motility in testis via interaction with OR4M1 receptor (By similarity).

**Subcellular location:**

**Asprosin:**

Secreted (*experimental evidence*)

**NOTE:** Secreted by white adipose tissue and circulates in the plasma.

**Unnamed:**

Secreted (*experimental evidence*)

**NOTE:** Fibrillin-1 and Asprosin chains are still linked together during the secretion from cells, but are subsequently separated by furin (PubMed:24982166).

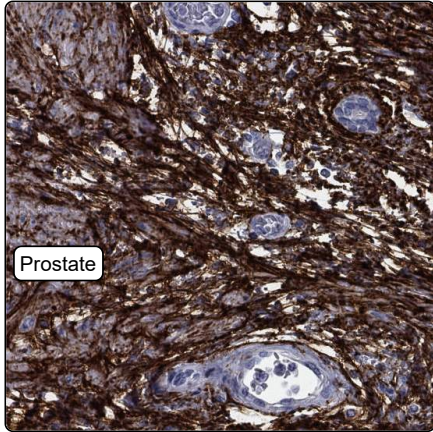
**Fibrillin-1:**

Secreted > Extracellular space > Extracellular matrix (*experimental evidence*)

**Protein existence:** Experimental evidence at protein level

**Comment:**

## Immunohistochemistry



<b>IHC protocol:</b>	HIER pH6, Dilution 1:800
<b>IHC test staining:</b>	Positivity in extracellular matrix.
<b>Literature conformance:</b>	Consistent with extensive gene/protein characterization data
<b>Literature significance:</b>	
<b>RNA similarity:</b>	High consistency between antibody staining and RNA expression data
<b>RNA tissue specificity:</b>	Tissue enhanced (adipose tissue,placenta)
<b>RNA tissue distribution:</b>	Detected in all
<b>IHC Sibling similarity:</b>	Other antibody shows similar IHC staining pattern
<b>Reliability score:</b>	Supported
<b>APE summary:</b>	Membranous and cytoplasmic expression in extracellular matrix, stromal cells, smooth muscle cells and decidual cells.
<b>APE explanatory sentences:</b>	High consistency between antibody staining and RNA expression data. Secreted protein, tissue location of RNA and protein is expected to differ. Antibody staining in cells/structures not annotated, view images.
<b>Orthogonal validation:</b>	No
<b>Independent validation:</b>	No
<b>IHC Annotation summary:</b>	Strong cytoplasmic positivity was observed in stromal cells, fibroblasts and extracellular matrix. All cancers were negative.