

LOXL3 Antibody (C-term) Blocking peptide
Synthetic peptide
Catalog # BP12837b**Specification**

LOXL3 Antibody (C-term) Blocking peptide - Product InformationPrimary Accession [P58215](#)**LOXL3 Antibody (C-term) Blocking peptide - Additional Information****Gene ID** 84695**Other Names**

Lysyl oxidase homolog 3, 143-, Lysyl oxidase-like protein 3, LOXL3, LOXL

Format

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

Storage

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

Precautions

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

LOXL3 Antibody (C-term) Blocking peptide - Protein Information**Name** LOXL3 {ECO:0000303|PubMed:11386757, ECO:0000312|HGNC:HGNC:13869}**Function**

Protein-lysine 6-oxidase that mediates the oxidation of peptidyl lysine residues to allysine in target proteins (PubMed:17018530, PubMed:28065600). Catalyzes the post-translational oxidative deamination of peptidyl lysine residues in precursors of elastin and different types of collagens, a prerequisite in the formation of cross-links between collagens and elastin (PubMed:17018530). Required for somite boundary formation by catalyzing oxidation of fibronectin (FN1), enhancing integrin signaling in myofibers and their adhesion to the myotendinous junction (MTJ) (By similarity). Acts as a regulator of inflammatory response by inhibiting differentiation of naive CD4(+) T-cells into T-helper Th17 or regulatory T-cells (Treg): acts by interacting with STAT3 in the nucleus and catalyzing both deacetylation and oxidation of lysine residues on STAT3, leading to disrupt STAT3 dimerization and inhibit STAT3 transcription activity (PubMed:28065600). Oxidation of lysine residues to allysine on STAT3 preferentially takes place on lysine residues that are acetylated (PubMed:28065600). Also able to catalyze deacetylation of lysine residues on STAT3 (PubMed:28065600).

Cellular Location

Secreted, extracellular space {ECO:0000250|UniProtKB:Q9Z175}. Cytoplasm. Nucleus Note=It is unclear how LOXL3 is both intracellular (cytoplasmic and nuclear) and extracellular: it contains a clear signal sequence and is predicted to localize in the extracellular medium. However, the intracellular location is clearly reported and at least another protein of the family (LOXL2) also has intracellular and extracellular localization despite the presence of a signal sequence (PubMed:28065600). [Isoform 2]: Cytoplasm. Secreted, extracellular space

Tissue Location

Isoform 1: Predominantly detected in the heart, placenta, lung, and small intestine (PubMed:17018530). Isoform 2: Highly detected in the kidney, pancreas, spleen, and thymus, and is absent in lung (PubMed:17018530). In eye, present in all layers of corneas as well as in the limbus and conjunctiva (at protein level) (PubMed:26218558).

LOXL3 Antibody (C-term) Blocking peptide - Protocols

Provided below are standard protocols that you may find useful for product applications.

- [Blocking Peptides](#)

LOXL3 Antibody (C-term) Blocking peptide - Images**LOXL3 Antibody (C-term) Blocking peptide - Background**

This gene encodes a member of the lysyl oxidase gene family. The prototypic member of the family is essential to the biogenesis of connective tissue, encoding an extracellular copper-dependent amine oxidase that catalyses the first step in the formation of crosslinks in collagens and elastin. A highly conserved amino acid sequence at the C-terminus end appears to be sufficient for amine oxidase activity, suggesting that each family member may retain this function. The N-terminus is poorly conserved and may impart additional roles in developmental regulation, senescence, tumor suppression, cell growth control, and chemotaxis to each member of the family. Alternatively spliced transcript variants of this gene have been reported but their full-length nature has not been determined.

LOXL3 Antibody (C-term) Blocking peptide - References

Kim, Y., et al. Oncol. Rep. 22(4):799-804(2009) Sebban, S., et al. Virchows Arch. 454(1):71-79(2009) Akagawa, H., et al. Hum. Genet. 121 (3-4), 377-387 (2007) :Lee, J.E., et al. J. Biol. Chem. 281(49):37282-37290(2006) Peinado, H., et al. EMBO J. 24(19):3446-3458(2005)