

LOXL2 Antibody

Rabbit mAb

Catalog # AP91535

Product Information

Application	WB, IF, FC, ICC
Primary Accession	Q9Y4K0
Reactivity	Rat, Human, Mouse
Clonality	Monoclonal
Other Names	LOR2; LOX L2; LOXL2; Lysyl oxidase homolog 2; Lysyl oxidase like 2; WS9 14;
Isotype	Rabbit IgG
Host	Rabbit
Calculated MW	86725

Additional Information

Dilution	WB 1:500~1:2000 ICC/IF 1:50~1:200 FC 1:50
Purification	Affinity-chromatography
Immunogen	A synthesized peptide derived from human LOXL2
Description	Mediates the post-translational oxidative deamination of lysine residues on target proteins leading to the formation of deaminated lysine (allysine). When secreted in extracellular matrix, promotes cross-linking of extracellular matrix proteins by mediating oxidative deamination of peptidyl lysine residues in precursors to fibrous collagen and elastin.
Storage Condition and Buffer	Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid freeze / thaw cycle.

Protein Information

Name	LOXL2
Function	Mediates the post-translational oxidative deamination of lysine residues on target proteins leading to the formation of deaminated lysine (allysine) (PubMed:27735137). Acts as a transcription corepressor and specifically mediates deamination of trimethylated 'Lys-4' of histone H3 (H3K4me3), a specific tag for epigenetic transcriptional activation (PubMed:27735137). Shows no activity against histone H3 when it is trimethylated on 'Lys-9' (H3K9me3) or 'Lys-27' (H3K27me3) or when 'Lys-4' is monomethylated (H3K4me1) or dimethylated (H3K4me2) (PubMed:27735137). Also mediates deamination of methylated TAF10, a member of the transcription factor IID (TFIID) complex, which induces release of TAF10 from promoters, leading to inhibition of TFIID-dependent transcription (PubMed:25959397). LOXL2-mediated deamination of TAF10 results in transcriptional repression of genes required for embryonic stem cell pluripotency including POU5F1/OCT4, NANOG, KLF4 and SOX2 (By similarity). Involved in epithelial to mesenchymal

transition (EMT) via interaction with SNAI1 and participates in repression of E-cadherin CDH1, probably by mediating deamination of histone H3 (PubMed:[16096638](#), PubMed:[24414204](#), PubMed:[27735137](#)). During EMT, involved with SNAI1 in negatively regulating pericentromeric heterochromatin transcription (PubMed:[24239292](#)). SNAI1 recruits LOXL2 to pericentromeric regions to oxidize histone H3 and repress transcription which leads to release of heterochromatin component CBX5/HP1A, enabling chromatin reorganization and acquisition of mesenchymal traits (PubMed:[24239292](#)). Interacts with the endoplasmic reticulum protein HSPA5 which activates the IRE1-XBP1 pathway of the unfolded protein response, leading to expression of several transcription factors involved in EMT and subsequent EMT induction (PubMed:[28332555](#)). Involved in E-cadherin repression following hypoxia, a hallmark of EMT believed to amplify tumor aggressiveness, suggesting that it may play a role in tumor progression (PubMed:[20026874](#)). When secreted into the extracellular matrix, promotes cross-linking of extracellular matrix proteins by mediating oxidative deamination of peptidyl lysine residues in precursors to fibrous collagen and elastin (PubMed:[20306300](#)). Acts as a regulator of sprouting angiogenesis, probably via collagen IV scaffolding (PubMed:[21835952](#)). Acts as a regulator of chondrocyte differentiation, probably by regulating expression of factors that control chondrocyte differentiation (By similarity).

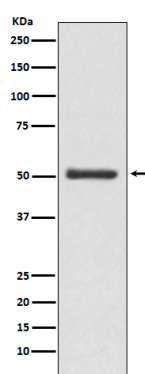
Cellular Location

Secreted, extracellular space, extracellular matrix, basement membrane. Nucleus. Chromosome. Endoplasmic reticulum. Note=Associated with chromatin (PubMed:[27735137](#)). It is unclear how LOXL2 is nuclear as it contains a signal sequence and has been shown to be secreted (PubMed:[23319596](#)) However, a number of reports confirm its intracellular location and its key role in transcription regulation (PubMed:[22204712](#), PubMed:[22483618](#)).

Tissue Location

Expressed in many tissues (PubMed:[10212285](#)). Highest expression in reproductive tissues, placenta, uterus and prostate (PubMed:[10212285](#)). In esophageal epithelium, expressed in the basal, prickle and granular cell layers (PubMed:[22204712](#)). Up-regulated in a number of cancers cells and tissues.

Images



Western blot analysis of LOXL2 expression in MCF7 cell lysate.

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