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# **Gene Section**

**Mini Review** 

## LASP1 (LIM and SH3 protein)

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Published in Atlas Database: August 2005

Online updated version: http://AtlasGeneticsOncology.org/Genes/Lasp1ID203.html DOI: 10.4267/2042/38246

This article is an update of: Rio MC. LASP1 (LIM, actin binding and SH3 protein). Atlas Genet Cytogenet Oncol Haematol.2000;4(2):51.

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### Identity

Other names: MLN50, EVI149

HGNC (Hugo): LASP1

Location: 17q12-21

**Local order:** from centromere to telomere are: TRAF4 (alias MLN62/CART1), MLLT6 (alias AF17), LASP1, STARD3 (alias MLN64), ERBB2 (alias c-erbB2), and RARA.

## DNA/RNA

#### Description

LASP1 encompasses 51.65 kb on the genomic level and consists of 7 exons.

#### Transcription

3845 bp mRNA, 783 bp coding sequence.

#### **Protein**

#### Description

261 amino acids; 29 kDa. LASP1 encodes a member of a LIM (Lin-11, Isl-1 and Mec-3) protein subfamily and is characterized by a LIM motif (cysteine-rich LIM/double zinc finger motif) at the N-terminus, an SH3 domain (Src homology region 3) at the Cterminus, and two actin-binding domains in the core of the protein

#### Expression

Ubiquitous.

#### Localisation

Intracellular, cytoplasmic; associated with the F-actin rich cortical cytoskeleton.

#### Function

LASP1 plays an important role in the regulation of dynamic actin-based, cytoskeletal activities and cell motility. Agonist-dependent changes in LASP1 phosphorylation may also serve to regulate actinassociated ion transport activities, not only in the parietal cell but also in certain other F-actin-rich secretory epithelial cell types. Together, (LIM-) nebulette, Lasp-1, and zyxin may play an important role in the organization of focal adhesions.

#### Homology

LASP family of proteins: actin-binding repeats similar to those in LASP1 are also present in other nebulinrelated proteins such as NEBL (nebulette, 107 kD actin-binding Z-disk protein) and NRAP (nebulinrelated anchoring protein); NRAP also contains an Nterminal LIM domain and NEB (nebulin) a C-terminal SH3 domain, both of which are highly homologous to the respective domains of LASP1.

## Implicated in

#### t(11;17)(q23;q12) --> MLL-LASP1

#### Disease

Infant AML-M4; only one case described so far.

#### **Abnormal protein**

The MLL-LASP1 chimeric protein consists of the AThook DNA-binding domain and the methyltransferase motif including the CXXC zinc-finger domain of MLL and the SH3 domain of LASP1.

#### Breast carcinomas

#### Disease

17q11-q21 amplification is found in about 25% of primary breast carcinomas; simultaneous amplification

and overexpression of LASP1 and ERBB2.

#### Prognosis

Poor clinical outcome; increase risk of relapse.

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This article should be referenced as such:

Strehl S. LASP1 (LIM and SH3 protein). Atlas Genet Cytogenet Oncol Haematol. 2005; 9(4):310-311.