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Gene Section

Mini Review

LASP1 (LIM and SH3 protein)

Sabine Strehl

Children's Cancer Research Institute, Kinderspitalgasse 6, A-1090 Vienna, Austria (SS)

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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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