

Gene Section

Mini Review

ADD3 (adducin 3)

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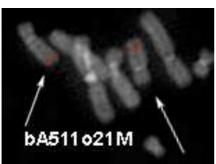
Identity

Other names: ADDL; Adducin-like; adducin 3 (gamma); gamma adducin (adducin-like protein 70)

HGNC (Hugo): ADD3 **Location:** 10q25.1-q25.2

Local order: MXI1 (MAX-interacting protein 1) is more telomeric. XPNPEP1 (X-prolylaminopeptidase 1)

is more telomeric.



Probe(s) - Courtesy Mariano Rocchi, Resources for Molecular Cytogenetics.

DNA/RNA

Description

15 exons spanning 129.52 Kb on 10q25.1-10q25.2. Transcription is from centromere to telomere.

Transcription

2 alternative transcripts:

Variant 1 includes an in-frame alternate coding exon and encodes a longer protein isoform (isoform a) compared to transcript variant 2.

Variant 2 lacks an in-frame alternate coding exon and exon 13 and encodes a shorter protein (isoform b) compared to transcript variant 1.

3 alternative transcripts corresponding to variant 1 and 2.



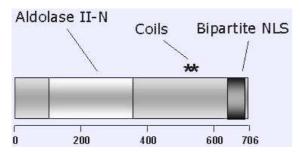
Genomic structure of ADD3. Black boxes indicate exons.

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Protein

Note

Adducin 3 (adducin-like).



Schematic representation of ADD3. AldolaseII-N: domain belonging to aldolase class I family, adducin subfamily N-terminal, Coils: coiled-coil domain, Bipartite NLS: Bipartite nuclear localization signal.

Description

membrane-cytoskeleton-associated protein that promotes the assembly of the spectrin-actin network binds to calmodulin. Adducins are heteromeric proteins composed of different subunits referred to as adducin alpha, beta and gamma encoded by distinct genes and belong to a family of membrane skeletal proteins involved in the assembly of spectrin-actin network in erythrocytes and at sites of cell-cell contact in epithelial tissues. Structurally, each subunit is comprised of two distinct domains. The amino-terminal region is protease resistant and globular in shape, while the carboxyterminal region is protease sensitive. The latter contains multiple phosphorylation sites for protein kinase C, the binding site for calmodulin, and is required for association with spectrin and actin. Alternatively spliced adducin gamma transcripts encoding different isoforms have been described. The functions of the different isoforms are not known.

Expression

Ubiquitous. Heart only expresses isoform 1 or a.

Localisation

Membrane skeleton.

Function

Adducins are membrane skeletal proteins involved in the assembly of spectrin-actin network in erythrocytes and at sites of cell-cell contact in epithelial tissues. Adducin is a heterodimeric cytoskeleton protein and consists of an [alpha]-subunit (Mr 103 kDa) and either

a [beta]- (Mr 97 kDa) or [gamma]-subunit (Mr 90kDa). Three genes (ADD1, ADD2, and ADD3, respectively) that map to different chromosomes encode these subunits. Adducin promotes the organization of the spectrin-actin lattice by favoring the spectrin-actin binding and controlling the rate of actin polymerization as an end-capping actin protein. Its function is calciumand calmodulin-dependent. It is phosphorylated by protein kinases A and C, tyrosine, and [rho]-kinases.10 It is a member of the myristoylated alanine-rich C kinase substrate protein family, which is involved in signal transduction, cell-to-cell contact formation, and cell migration. Non-silent polymorphisms resulting in subunits alpha and beta have been associated with the regulation of blood pressure in an animal model of hypertension.

Homology

It presents homology in various species. It also belongs to ADDUCIN protein family (3 members ADD1, ADD2 and ADD3).

Mutations

Germinal

It has been shown that the interaction of ADD1 and ADD3 gene variants in humans is statistically associated with variation in blood pressure, suggesting the presence of epistatic effects among these loci.

Somatic

t(10;11)(q25;p15) 5' NUP98-ADD3 3' fusion in a T-cell acute lymphoblastic leukemia with biphenotipic characteristics (T/myeloid).

Implicated in

t(10;11)(q25;p15)

Disease

T-cell acute lymphoblastic leukemia with biphenotipic characteristics (T/myeloid).

Prognosis

Bad.

Cytogenetics

t(10;11)(q25;p15) (cryptic).

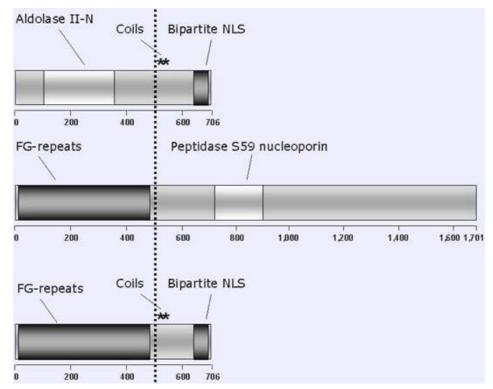
Hybrid/Mutated gene

5' NUP98 - ADD3 3'.

Abnormal protein

NUP98-ADD3.

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Schematic representation of the fusion NUP98-ADD3 consecuence of the t(10;11)(q25;p15) in a T-cell acute lymphoblastic leukemia with biphenotipic characteristics. From up to down: ADD3, NUP98 and the putative chimeric NUP98-ADD3 structure. FG-repeats, phenilalanine-glycine repeats; bipartite NLS, bipartite nuclear localization signal. Coiled coil domains on ADD3 and NUP98-ADD3 are indicated with asterisks.

References

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