

Gene Section

Short Communication

SS18L1 (synovial sarcoma translocation gene on chromosome 18-like 1)

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Identity

Other names: CREST; KIAA0693; MGC26711; SYT

homolog 1

HGNC (Hugo): SS18L1 **Location:** 20q13.3

Local order: chr20:60,152,245-60,190,935 (UCSC

latest release: May 2004).

DNA/RNA

Note

Member of the SS18 family.

Description

11 exons with similar splice sites as SS18.The promoter region lacks CAAT and TATA boxes but contains CpG islands, suggesting that SS18L1 is a housekeeping gene.

Pseudogene

SS18L2 (3p21).

Protein

Description

396 amino acids, 42990 Da. The SS18L1 protein, similarly to the SS18 protein, exhibits two domains: a SYT N-terminal homology domain found in a wide variety of species ranging from plants to humans and the QPGY domain at the COOH-terminal part, rich in glutamine, proline, glycine, and tyrosine. The QPGY domain of the SS18 protein may activate transcription when coupled to a DNA-binding domain.

Expression

Ubiquitous; with lowest levels in spleen.

Localisation

Nuclear?

Function

Calcium-responsive transactivator: CREST is a SYT related nuclear protein that interacts with CREB-binding protein (CBP) and is expressed in the developing brain.

Homology

SS18, SS18L2.

Implicated in

Disease

Synovial sarcoma.

Prognosis

Unknown.

Cytogenetics

t(X;20)(p11;q13.3).

Hybrid/Mutated gene

In the SS18L1/SSX1 transcript detected in the synovial sarcoma, the exon 10 of SS18L1, which corresponds to exon 10 of SS18, was fused to exon 6 of SSX1.

Abnormal protein

In the putative SS18L1/SSX1 chimeric protein, the last 8 amino acid residues of the SS18L1 protein are replaced by 78 amino acids from the COOH-terminal part of SSX1. By analogy with what is presumed to be the case for the SS18/SSX fusion protein,

SS18L1/SSX1 is likely to show an altered transcriptional pattern with the COOH-terminal SSX domain, redirecting the SS18L1 activation domain to new target promoters.

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