

## Gene Section

### Mini Review

## ACTB (Actin, beta)

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### Identity

**Other names:** Beta cytoskeletal actin; Beta actin

**HGNC (Hugo):** ACTB

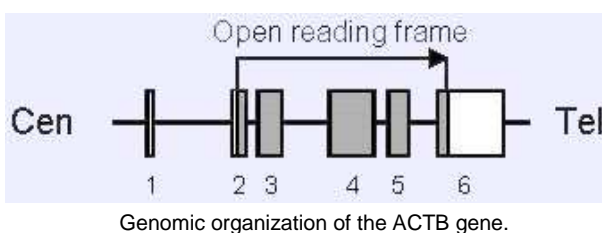
**Location:** 7p22

**Location (base pair):** Position 5340026-5343462 on the chromosome 7 genomic sequence.

#### Note

Six actin isoforms are known: two sarcomeric (a-skeletal and a-cardiac), two smooth muscle actins (a and g), and two non-muscles, cytoskeletal actins (b and g).

### DNA/RNA



#### Description

Six exons, spans approximately 3.4 kb of genomic DNA in the centromere-to-telomere orientation. The translation initiation codon ATG is located in exon 2, and the stop codon in exon 6.

#### Transcription

mRNA of approximately 1.8 kb.

#### Pseudogene

At least 19 processed, non-expressed, pseudogenes are dispersed throughout the genome.

### Protein

#### Description

The open reading frame encodes a 374 amino acid protein, with an estimated molecular weight of approximately 41.7 kDa.

#### Expression

Abundantly expressed in all mammalian and avian non-muscle cells.

#### Localisation

Cytoplasm.

#### Function

Component (together with cytoplasmic g actin) of the cytoskeletal microfilaments. Involved in the transport of chromosomes and organelles as well as in cell motility.

#### Homology

The ACTB proteins are evolutionary conserved. Mammalian cytoplasmic actins (actin g and b) are remarkably similar to each other, but differ in at least 25 residues from the muscle actins.

### Mutations

#### Somatic

ACTB is interrupted by the t(7;12)(p22;q13) detected in pericytoma with t(7;12).

### Implicated in

#### Disease

Pericytoma with t(7;12).

#### Prognosis

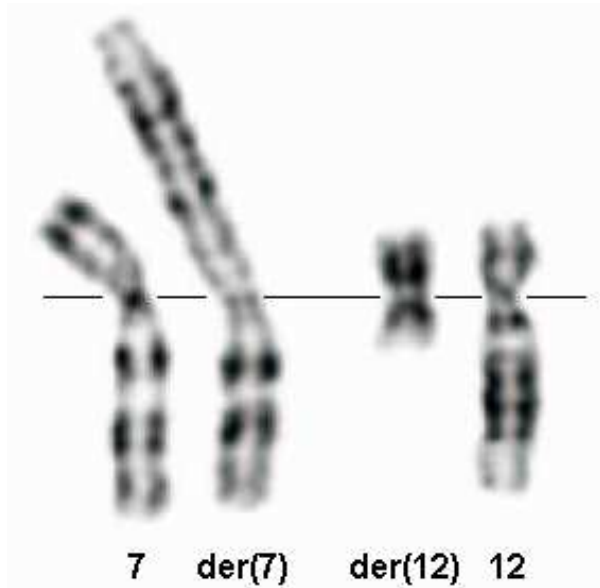
Benign or low-malignant.

**Cytogenetics**

t(7;12)(p22;q13).

**Hybrid/Mutated gene**

ACTB-GLI1 fusion gene. The breakpoints reported so far have been located to introns 1, 2 or 3 within the ACTB gene, and to introns 5 or 6 or to exon 7 within the GLI1 gene. Reciprocal GLI1-ACTB gene fusions have also been detected. The breakpoints have been located to introns 5 or 7 within the GLI1 gene, and to intron 3 of the ACTB gene.



Representative G-banded partial karyotype of the t(7;12)(p22;q13).

**Abnormal protein**

The ACTB-GLI1 fusion protein contains the N-terminal of ACTB and the C-terminal of GLI1, including the DNA-binding zinc finger motifs (encoded by exons 7-10) and transactivating motifs (exon 12).

**Oncogenesis**

It is suggested that the strong ACTB promoter causes an overexpression of GLI1 sequences important for transcriptional activation of downstream target genes, akin to the oncogenic mechanisms of the COL1A1-

PDGFB fusion gene detected in dermatofibrosarcoma protuberans.

**References**

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