

Gene Section

Short Communication

APC (adenomatous polyposis coli)

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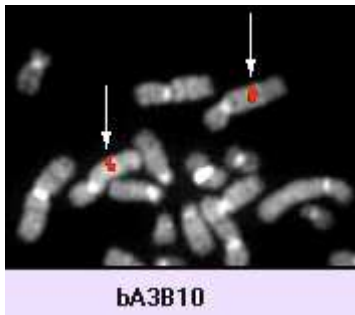
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Identity

Location: 5q21



APC (5q21) - Courtesy Mariano Rocchi, Resources for Molecular Cytogenetics. Laboratories willing to validate the probes are welcome: contact_rocchi@biologia.uniba.it.

DNA/RNA

Description

15 exons (with a particularly large 15th exon).

Transcription

9.0 kb mRNA; 8538 bp open reading frame.

Protein

Description

2843 amino acids; 300 kDa.

Function

Tumour suppressor gene; the APC normal gene product interacts with the adherens junction proteins α and β -catenin suggesting it is involved in cell adhesion; since β -catenin activates transcription by forming complexes with members of the Tcf-Lef family, it is supposed that APC, by complexing β -catenin, may also inhibit the transcription pathway regulated by the β -catenin/Tcf complex.

Homology

Limited functional homology to known proteins.

Mutations

Germinal

In familial adenomatous polyposis.

Somatic

The APC gene is mutated in about 50% of sporadic colorectal tumours; the great majority of APC gene mutations are frameshifts of nonsense mutations resulting in premature stop codons.

Implicated in

Colorectal cancer

Disease

Adenocarcinoma; the third most frequent cancer in the world.

Prognosis

5 yr survival rate is around 50%.

Cytogenetics

There are two types, according to the ploidy: 1-aneuploid tumours showing numerous allelic losses, in particular loss of heterozygosity on chromosomes 17 and 18, and 2- diploid tumours without frequent allelic losses.

Oncogenesis

A number of genes (oncogenes and tumour suppressor genes) are known to be implicated in tumour progression in colorectal cancers; other genes are: p53, Ki-ras, and mismatch repair genes (MMR genes).

Familial adenomatous polyposis

Disease

Autosomal dominant cancer prone disease.

Prognosis

Colorectal cancer is the first cause of death in this disease.

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