



## A Brief Note on Cancer Genetics

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### DESCRIPTION

Cancer is a hereditary disease, and tumor cells differ from normal progenitor cells due to genetic changes that affect growth regulatory genes. There are two classes of such oncogenes: oncogenes that function as positive growth regulators and tumor suppressor genes that act as negative growth regulators. Oncogenes are mainly conserved among various organisms and are active in transmitting growth signals from the periphery of cells to the cell nucleus. These signaling functions can be disrupted by many genetic changes. The result of altered growth signals is often cancer, tumor suppressor genes have an inhibitory effect on cell proliferation lost by gene inactivation or deletion. In cells transformed with the DNA virus, it is abolished by neutralization of the tumor suppressor protein by the viral gene product. Tumor suppressor genes were first recognized in hereditary cancers. Defects in tumor suppressors transmitted to the germline can lead to increased tumor incidence in offspring. However, tumor suppressors also play an important role in non-hereditary cancers.

Cancer genetics is a scientific field that studies genes and signaling pathways that promote the development of cancer. Cancer geneticists use several approaches, including analysis of the genome of cancer patients and their tumors, to identify oncogenes. These studies will be conducted in combination with experiments in *in vitro* and *in vivo* models to elucidate the mechanisms that promote tumorigenesis. Here's how to use these approaches to better understand how cancer develops. The goal of cancer genetics is to catalog and understand the genetic changes that contribute to the development of cancer. Oncogeneists are genes that, when mutated, contribute to the development of cancer by giving them the ability to promote the growth of cancer cells and bypass the cell cycles and cell death checkpoints that normally control their growth. We aim to identify oncogenes. Cancer geneticists are also interested in the

networks and pathways that contribute to tumor development, and how cancer genes work together to promote tumor development.

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### Diagnosis of familial cancer syndromes

The primary care physician is first to apply cancer genetics to clinical care when there is the suspicion of a familial cancer-prone syndrome. It could now no longer be viable to be expecting which mutations reason what familial cancers primarily based totally on expertise of the gene function. The diagnoses depend on pattern recognition and linkage to the known mutations. The popularity of a familial sample of disorder may also bring about in addition research of probably affected sufferers and families. The aim might be powerful screening and control recommendations.

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