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Biomedicine : Cancer Research

Burtness, Barbara, Golemis, Erica A. (Eds.), Yale University School of Medicine, New Haven, CT, USA

Molecular Determinants of Head and Neck Cancer

- Summarize the unique pathobiology of head and neck cancers
- Addresses signaling pathways and individual proteins that are important in causing or supporting the aggressive nature of head and neck cancer
- Discusses the sequencing and epigenomic analysis of the head and neck cancer genome
- Focuses on a new understanding of stem cell function in head and neck prognosis and therapy resistance
- Discusses a shifting focus of care for HPV positive and HPV negative disease

Squamous cell cancers of the head and neck (SCCHN), also known as head and neck cancers (HNC) encompass malignancies of the oral cavity, larynx, nasopharynx and pharynx, and are diagnosed in over 500,000 patients worldwide each year, accounting for 5% of all malignancies. It is estimated that approximately 50,000 patients develop head and neck cancer annually in the United States, of whom approximately 50% succumb to this cancer. For most cases of SCCHN, treatment is multimodal, often combining surgery or irradiation with chemotherapy; even successfully treated patients frequently experience durable and severe side effects. Improving cure rates and reducing chronic morbidity are urgent clinical needs for head and neck cancer. However, in contrast to cancer types such as breast or prostate that have been much studied and have well-defined biology, until recently, relatively few researchers investigated the molecular basis of HNC, making it difficult to design targeted treatments with better efficacy and less debilitating side effects. This volume will provide an overview of the factors contributing to disease pathogenesis, including the recognition of discrete molecular subtypes with distinct etiology, prognosis, and treatment response. This volume will familiarize the reader with the critical signaling pathways and oncogenic drivers for HNC.

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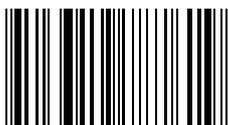
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