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These studies have successfully identified numerous mutations, gene expression differences, protein abundance differences, epigenetic mutations, and metabolite concentrations to be linked with gastric cancer heterogeneity and staging, which significantly improve our understanding of gastric cancer. Systems biology approaches aim to the transcendence of individual genes/proteins and to the integration of biological system that taking account into the intrinsic interactions.

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In this article, the authors discuss the molecular biology of esophageal and gastric cancer with a focus on esophageal adenocarcinoma. They review data from The Cancer Genome Atlas project and advances in the molecular stratification and classification of esophageal carcinoma and gastric cancer. They also summarize advances in microRNA, molecular staging, gene expression profiling, tumor microenvironment, and detection of circulating tumor DNA.

The Molecular Biologic Basis of Esophageal and Gastric Cancers

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The Biology Of Gastric Cancers

A research team led by Professor Wang Wen-ching of the Institute of Molecular and Cellular Biology, National Tsing Hua University (NTHU) has exploited the power of big data to identify two key biomarkers contributing to the proliferation and metastasis of gastric (stomach) cancer. Using the US Food and

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Molecular Biology of Gastric Cancer—PubMed

A research team led by Professor Wang Wen-ching of the Institute of Molecular and Cellular Biology, National Tsing Hua University (NTHU) has exploited the power of big data to identify two key biomarkers contributing to the proliferation and metastasis of gastric (stomach) cancer. Using the US Food and Drug Administration (FDA)-approved drugs, the team conducted targeted therapy which can ...