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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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~~The Molecular Biologic Basis of Esophageal and Gastric Cancers~~

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A research team led by Professor Wang Wen-ching of the

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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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