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Basic Study

Identification of the circRNA-miRNA-mRNA regulatory network and its prognostic effect in colorectal cancer

Yin TF *et al.* CircRNA-miRNA-mRNA network in CRC

Teng-Fei Yin, ¹¹ Dong-Yan Zhao, Yuan-Chen Zhou, Qian-Qian Wang, Shu-Kun Yao

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Feb 15, 2020 · 1. Introduction. **Cancer** is the first or second leading cause of death in many countries with rapidly increasing incidence and mortality rates (Bray et al., 2018). **Colorectal cancer** (CRC) is a global disease with an incidence rate of 10.2% and a mortality rate of 9.2% (Bray et al., 2018). After being diagnosed with CRC, patients have a 65% 5-year survival rate (Siegel et al., 2017).

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Identification of the circRNA-miRNA-mRNA regulatory network of Hsp90 inhibitor-induced cell death in colorectal cancer by integrated analysis.

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Identification of the circRNA-miRNA-mRNA regulatory ...

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The circRNA-miRNA-mRNA **regulatory network** of Hsp90 inhibitor-induced cell death in **colorectal cancer** was constructed. This **regulatory network** extends the underlying mechanism of Hsp90 and improves our understanding of Hsp90 inhibitors as potential targeted therapeutic agents.

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Feb 15, 2020 · The circRNA-miRNA-mRNA **regulatory network** of Hsp90 inhibitor-induced cell death in **colorectal cancer** was constructed. This **regulatory network** extends the underlying mechanism of Hsp90 and improves our understanding of Hsp90 inhibitors as potential targeted therapeutic agents.

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Author: Kaisheng Liu, Yaomin Guo, Kai Zheng, ...

Publish Year: 2020

Identification of circRNA-miRNA-mRNA regulatory network in ...

<https://cancerci.biomedcentral.com/articles/10.1186/s12935-019-0905-z> ▾

Jul 16, 2019 · Gastric **cancer** (GC) is one of the most common malignancies and the second leading cause of **cancer**-related deaths worldwide [1, 2]. Despite advances in surgical techniques and combined chemotherapy strategies, the 5-year overall survival (OS) of GC remains poor []. The lack of improvement in OS is largely due to a low early diagnostic rate and a high frequency of ...

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Identification of circRNA-miRNA-mRNA regulatory network in ...

<https://europepmc.org/article/PMC/PMC6636116> ▾

Jul 16, 2019 · We also constructed a **circRNA-miRNA-hub gene** subnetwork based on **regulatory** modules identified in the **circRNA-miRNA-mRNA network**. Numerous studies have shown that expression of circRNA is dysregulated in GC and that this dysregulation is associated with pathogenesis and **prognosis**, suggesting that circRNAs could be used as tumor-associated ...

Cited by: 24

Author: Yong-jun Guan, Jian-ying Ma, Wei Song, ...

Publish Year: 2019

Identification and analysis of circRNA-miRNA-mRNA ...

<https://digital-library.theiet.org/content/journals/10.1049/iet-syb.2020.0061> ▾

This study was to identify important circRNA-miRNA-mRNA (ceRNAs) **regulatory** mechanisms in hepatocellular carcinoma (HCC). The circRNA dataset GSE97332 and miRNA dataset GSE57555 were used for analyses. Functional enrichment analysis for miRNA and target gene was