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Cited by: 11 Author: Xiaoxiao Yao, Guangqiang You, Chen Zh...

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Background Cervical cancer (CC) is a malignant tumor found in the lowermost part of the womb. Evolving studies on CC have reported that circRNA plays a crucial role in CC **progression**. In this study, we investigated the main function of a novel circRNA, circ_0084927, and its regulatory network in CC development. Methods qRT-PCR was applied to evaluate the expression of circ_0084927, miR-1179 ...

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Fig. 5 miR-1179 directly inhibited CDK2 mRNA expression by binding to its 3UTR. a The potential binding site between miR-1179 and CDK2 was predicted by TargetScan Human 7.2. b The potential binding between miR-1179 and the 3UTR of CDK2 gene was validated by the luciferase reporter gene assay. CDK2 mutant or CDK2 wild-type plasmids containing the fluorescence group and miR-1179 were co ...

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Basic Study

CircRNA_0084927 promoted colorectal cancer progression by regulating the miRNA-20b-3p/glutathione S-transferase mu 5 axis

CircRNA_0084927 promoted CRC progression

Feng Liu, Xiao-Li Xiao, Yu-Jing Liu, Ruo-Hui Xu, Wen-jun Zhou, Han-Chen Xu, Ai-Guang Zhao, Yang-Xian Xu, Yan-Qi Dang, Guang Ji

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Accumulating evidence has shown that long non-coding RNAs (lncRNAs) play significant roles in the development and progression of many types of cancer including colorectal cancer. RP11-400N13.3 is a novel lncRNA discovered recently and its biological function and underlying mechanism in colorectal cancer remain elusive. This study aimed to reveal the relationship between RP11-400N13.3 and ...

Author: Hongju Yang, Qian Li, Yanrui Wu, Jian... Publish Year: 2020

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