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## Non-coding RNAs in hepatitis C-induced hepatocellular ...

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3848131>

Nov 28, 2013 · Core tip: Regulatory non-coding **RNAs** (ncRNAs), mainly microRNAs (miRNAs) and **long** non-coding **RNAs** (lncRNAs), are likely to play important roles in a variety of biological processes, including development and progression of **hepatitis C-induced hepatocellular carcinoma** (HCC). In this review, we focus on recent studies of ncRNAs, especially miRNAs and lncRNAs associated with **hepatitis C virus** ...

Cited by: 22

Author: Weihong Hou, Herbert L Bonkovsky

Publish Year: 2013

## Serum and exosome long non coding RNAs as potential ...

[jancer.org/v09p2631.htm](http://jancer.org/v09p2631.htm) ▾

**Serum and exosome long non coding RNAs** as **potential** biomarkers for **hepatocellular carcinoma**

Li Sun **1,2** , Yingying Su **2** , Xiangxiang Liu **1** , Mu Xu **1** , Xiaoxiang Chen **1** , Yefei Zhu **2** , Zhirui Guo **2** , Tingting Bai **2** , Lin Dong **1** , Chenchen Wei **2** , Xiaoxiao Cai **2** , Bangshun He **1** , Yuqin Pan **1** , Huiling Sun **1** , Shukui Wang **1**

Cited by: 13

Author: Li Sun, Yingying Su, Xiangxiang Liu, Mu X...

Publish Year: 2018

## Long non-coding RNA AOC4P suppresses hepatocellular ...

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4695122>

Sep 15, 2015 · Increasing evidence indicates that **long** non-coding **RNAs** (lncRNAs) regulate diverse cellular processes, including cell growth, differentiation, apoptosis, and cancer progression. However, the function of lncRNAs in the progression of **hepatocellular carcinoma** (HCC) remains largely unknown.

Cited by: 77

Author: Tong-Hong Wang, Yong-Shiang Lin, Ying ...

Publish Year: 2015

## Long non-coding RNAs and hepatocellular carcinoma (Review)

<https://www.spandidos-publications.com/mco/3/1/13> ▾

Recent advances in **next-generation sequencing** technology in transcriptome analysis have helped

## Match Overview



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**Name of Journal:** *World Journal of Gastrointestinal Oncology*

**Manuscript NO:** 47823

**Manuscript Type:** ORIGINAL ARTICLE

***Basic Study***

**Eight key long non-coding RNAs predict hepatitis virus positive  
hepatocellular carcinoma as the prognostic targets**

Huang ZL *et al.* Eight-lncRNA signature for virus HCC prognosis

Zi-Lin Huang, Wang Li, Qi-Feng Chen, Pei-Hong Wu, Lu-Jun Shen





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## Novel predictive and prognostic strategies of hepatitis B ...

<https://hrjournal.net/article/view/1716> ▾

Hepatitis B virus-DNA level and basal core promoter A1762T/G1764A mutation in liver tissue independently predict postoperative survival in hepatocellular carcinoma. Hepatology 2010;52:1922-33. ...

## Analysis of long noncoding RNA expression in ...

[www.ncbi.nlm.nih.gov/Journal List > J Transl Med > v.14; 2016](http://www.ncbi.nlm.nih.gov/JournalList/JTranslMed.v.14.2016)

Nov 28, 2016 · Keywords: Hepatocellular carcinoma, Long noncoding RNA, Hepatitis B virus, Hepatitis C virus, Hepatitis D virus Background Recent findings from genome tiling arrays and RNA sequencing have revealed the existence of a large number of RNAs that lack protein-coding capacity, which can be divided into two groups, namely, small non-coding RNAs (sncRNA) (<200 bp) and long non-coding RNAs (lncRNAs)...

Cited by: 12

Author: Quan Zhang, Quan Zhang, Kentaro Matsu...

Publish Year: 2016

## Alcohol and hepatitis virus-dysregulated lncRNAs as ...

[www.oncotarget.com/index.php?journal=oncotarget&page=article&op...](http://www.oncotarget.com/index.php?journal=oncotarget&page=article&op...) ▾

17. Iizuka N, Oka M, Yamada-Okabe H, Mori N, Tamesa T, Okada T, Takemoto N, Tangoku A, Hamada K, Nakayama H, Miyamoto T, Uchimura S, Hamamoto Y. Comparison of gene expression profiles between hepatitis B virus- and hepatitis C virus-infected hepatocellular carcinoma by oligonucleotide microarray data on the basis of a supervised learning method.

Cited by: 1

Author: Hao Zheng, Pinxue Li, James G. Kwok, A...

Publish Year: 2018

## Development Of A Three-Gene Prognostic Signature For ...

[jcancer.org/v09p1989.htm](http://jcancer.org/v09p1989.htm) ▾

Hepatitis B virus (HBV) is a major cause of hepatocellular carcinoma (HCC), however, it remains largely unknown about the specific gene prognostic signature of HBV-associated HCC. Using Robust Rank Aggreg (RRA) method to integrate seven whole genome expression datasets, we identified 82 up-regulated genes and 577 down-regulated genes in HBV-associated HCC patients.

## Non-coding RNAs in hepatitis B or C-associated ...

<https://www.sciencedirect.com/science/article/pii/S030438351200170X>

Some ncRNAs have diagnostic and prognostic potentials and may also serve as therapeutic targets for cancers. Hepatocellular carcinoma (HCC) is the third leading cause of worldwide cancer-related death. The major risk factor for HCC is chronic hepatitis B virus (HBV) and/or hepatitis C virus (HCV) infection. There are few effective biomarkers for early diagnosis and prognosis prediction of HCC.