

January 01, 2014

Dear Editor,

The manuscript has been improved according to the suggestions of reviewers' comments

Title: Epigenetics in hepatocellular carcinoma: Updates and future therapy perspectives

Author: Li Ma, Mei-Sze Chua, Ourania Andrisani, Samuel So

Name of Journal: World Journal of Hepatology

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Answer to reviewer No. 1

Comments 1-4, typographical and grammatical errors have been corrected. 5. Statement of "What is transcribed by RNA polymerase..." have been corrected, 6. Brief description of Methylation450 BeadChip is included.

Answer to reviewer No. 2

1. Please discuss why overexpression of Ezh2 is found in advanced HCC.

Explanation of EZH2 overexpression is discussed in references 35 and 36.

2. The Authors state that miR-26 expression is correlated to the response to interferon. Is there any evidence for this?

As discussed in reference 62, interferon response were evaluated in HCC patients.

Answer to reviewer No. 3

Comments 1: Reference of "Exploring genome-wide DNA methylation profiles altered in hepatocellular carcinoma using Infinium HumanMethylation 450 BeadChips" is included.

Comments 2: All the references are updated. Abbreviation of HDACi are indicated appropriately in the revised manuscript

  
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