Author's response to reviews

Title: Whole-genome sequencing of matched primary and metastatic hepatocellular carcinomas

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Dear Editor,

Thank you very much for reviewing our manuscript.

As requested by the referee, we have validated major somatic mutations found in this study. Below please find our answer to your respectful comments:

Although these authors address some questions raised by this referee, they did not confirm the main findings such as some key somatic mutations in the revised version. Also, these authors state they add Figures 5 and 6; I can see figure legends but not figures in main text.

# To address the respectful comments by the reviewer, we selected 60 sites for sanger-seq. Finally, 66 somatic SNVs were covered with 60 site validatations including some main somatic SNVs. The results are attached as Supplementary Table 11.

# There was a technical error when uploading the files. We noticed that Figures 5 and 6 were not included. This time, we have included Figures 5 and 6.