

Dear editor,

Title: Bioinformatics analysis of aberrantly methylated-differentially expressed genes and pathways in hepatocellular carcinoma

MS Number: 38793

Our data are from microarrays downloaded from Gene Expression Omnibus (GEO, <https://www.ncbi.nlm.nih.gov/geo/>), which are not related human tissues or animals. So, we do not need comments including "Institutional animal care and use committee statement", "Institutional review board statement", and "The ARRIVE Guidelines".

Thank you very much.

Best Regards.

Name: Xuemei Wang

Signature:

Xuemei Wang 2018.3.14