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Erratum

Macrophages activating chemokine (C-X-C motif) ligand 8/miR-17 cluster modulate hepatocellular carcinoma cell growth and metastasis: Am J Transl Res. 2017; 9(5): 2403-2411

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In this published article, there was a statistical error in the **Figure 5E**, **5F**, in which the shape of survival curve was not accurately presented.

In the previous calculation, due to the improper use of statistical methods (simply connecting the survival time of patients with curve segments instead of using Kaplan Meier algorithm), the survival plot in the original paper turned into a curly line, rather than a broken line that should have been accurately presented. Therefore, to correct this error, we conducted survival statistics again. Kaplan-Meier method is now used to analyze the survival curves and a log-rank test is used to compare the survival curves.

This error does not change the main finding of **Figure 5** and the scientific conclusions of this manuscript in any way. The authors sincerely apologize for this error. The revised **Figure 5** is presented as the following.

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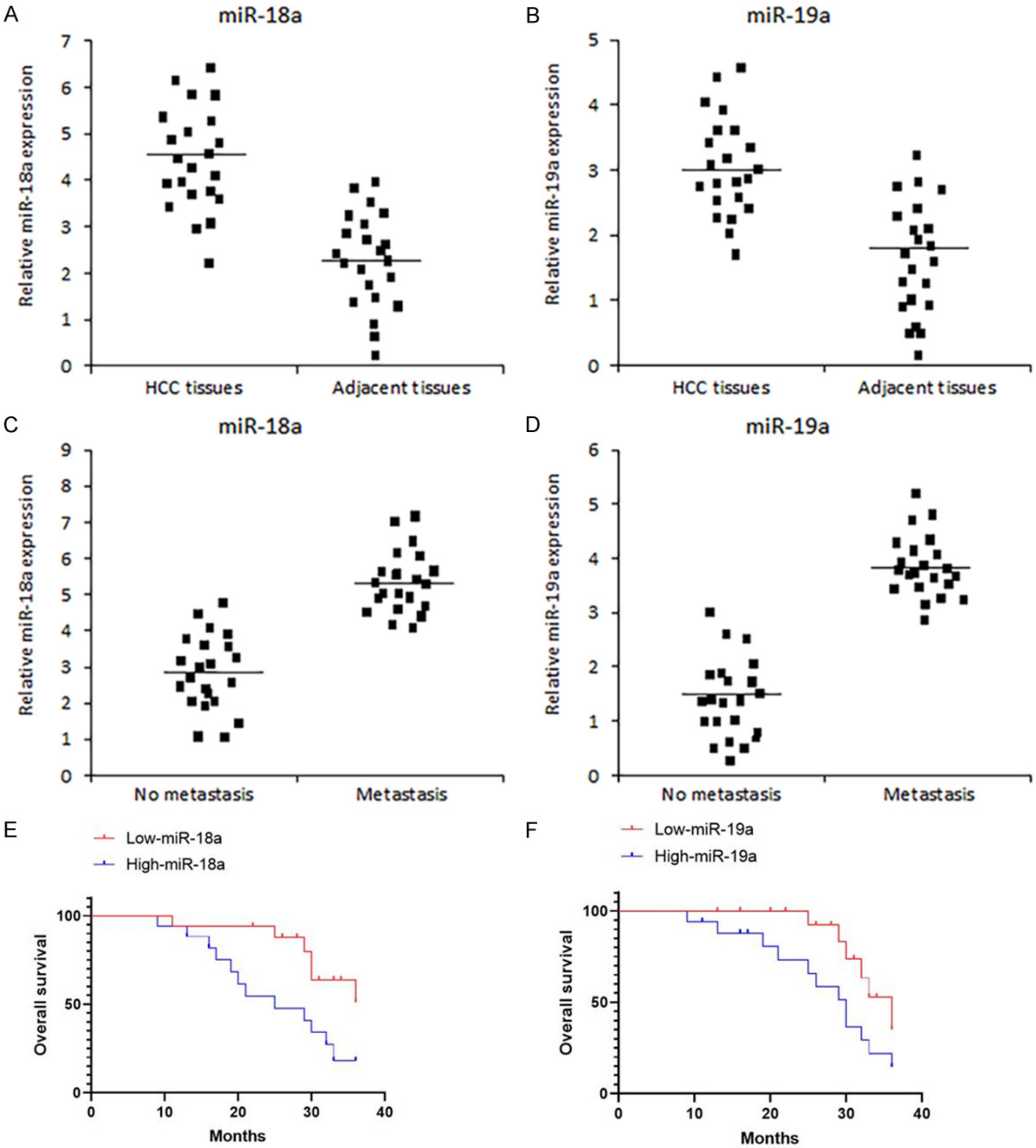

Figure 5. The expression of miR-18a and miR-19a is associated with HCC clinical features. A. B. miR-18a and miR-19b expression in HCC tissues examined by real time RT-PCR. C, D. Relationship between miR-18a or miR-19a expression and metastasis of HCC tissues. E, F. Relationship between miR-18a or miR-19a expression and life span in patients with HCC.