Author's response to reviews

Title: Paired design study by Real-time PCR: MiR-378 is one of the most possible early diagnose biomarker of human colorectal cancer.

Authors:

peng juan (dielianhuayaya@163.com)
xie zhengyong (pwkxiezy@163.com)
cheng liyang (chliyang2008@sina.com)
zhang yuxin (zhangyuxin-53825@163.com)
yu hongping (yhp268@163.com)
chen juyong (294821766@qq.com)
kang huixin (1367706921@qq.com)

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1. Although the content wasn’t fresh, but the statistical methods such as conditional logistic regression to analyze the same content was new. Also, the results and conclusion is valuable.
2. Yes, it’s TaqManHuman MicroRNA Array.
3. Yes, Normal mucosa usually from the edge of the site by at least 5 cm. Now, we think it’s better to exclude the surgery details for normal mucosa.
4. Yes, our department have the spss 16.0 software.
5. We have modified the table.
6. In multivariate regression analysis, we always to explore each microRNA is significant or not in the analysis. Then, select these significant microRNAs to do the last regression analysis. This variable selection process is called a series of selection process in the paper.
7. Yes, there is no mistake. As the figure 2 is only to describe the delta ct in both groups while the statistical test between groups is based the 2- Δ Δ ct value. Therefore, figure 2 is correct.
8. Well, ROC method is more visible presented with the figure, we have already added it in the paper.
9. The study determines we should use the previous results in different stage of miR-378*. Yes, to detect the miR-378* in early stage is an ideal result and it’s relies on more evident proof to make this conclusion.
10. Well, It’s better exclude.