

## Author's Response To Reviewer Comments

Close

Dear Editor

Thanks a lot for the positive feedback! Although you suggested not to carry out any further analyses, we decided to run several assemblers with k-mer sizes provided by rnaSPAdes (as suggested by Reviewer 3). Result for different assemblers are rather inconsistent (e.g. in some cases the default k is better, in some it is not). Even in cases when an assembly is improved, it still does not outperform rnaSPAdes.

We have added a paragraph regarding k-mer size selection in the beginning of the Analyses section. Detailed response to the Reviewer's comments are attached.

Best regards

Elena Bushmanova and Andrey Prjibelski

Close