Author's response to reviews

Title: Using a practical molecular capsular serotype prediction strategy to investigate Streptococcus pneumoniae serotype distribution and antimicrobial resistance in Chinese local hospitalized children

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Reply to the editor’ comments

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Dr Catherine Olino
BMC Pediatrics
Journal Editorial Office BioMed Central

Dear Dr Olino,

RE: MS: 1670154452176925 - Using a practical molecular capsular serotype prediction strategy to investigate *Streptococcus pneumoniae* serotype distribution and antimicrobial resistance in Chinese local hospitalized children

Thank you for the opportunity to revise our manuscript. We are pleased to address the editor’s comments point-by-point as below.

**Question 1.** Please move the section on ethical approval and consent from the end of the paper to the methods section.

**Answer:** Thanks for your suggestion. We have moved it to the methods section.

**Question 2.** The author information section (currently titled author details) is not compulsory. If you would like to include it, rather than stating the author affiliations, please format it according to the following guideline: http://bmcpediatr.biomedcentral.com/submission-guidelines/research-article

**Answer:** Thanks, we have simplified authors information section in title page.

**Question 3.** We encourage authors whose supporting data are available in an open access repository to include an Availability of supporting data section in their manuscript, before the Competing interests and Authors contributions. The section should state the name of the repository in which your data is deposited and include a link to the dataset DOI. If all the supporting data are included as additional files the section should state this. If your manuscript has any supporting sequence data, microarray data, or proteomic data this must be deposited in the appropriate repository and a link to the dataset should be included in the Availability of supporting data section.

**Answer:** Thanks for your suggestion. We have submitted all supplementary files to LabArchives, state the dataset DOI and link information in our manuscript. In addition, all our *cpsB* sequence GenBank sequences (accession numbers KT164777-KT164783) have been released to the public.
Thank you for taking the time to consider this manuscript for publication.

Yours sincerely,

Professor Qiyi Zeng