

## Re: SARS manuscript

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From: zlshi <zlshi@wh.iov.cn>  
To: lifang <lifang@umn.edu>  
Sent: April 25, 2018 9:39:52 PM CDT  
Received: April 25, 2018 9:40:03 PM CDT

Dear Fang,

I've read your manuscript and it's very good. Thank you!

There two modifications I would suggest:

- 1) the name of bat SL-CoVs in the figures. To be consistent with the description in other section, can you please modify the three representative bat strains as HKU3, WIV1 (identical as Rs3367) and RsSHC014 to replace the previously names? I think we need not to indicate the isolating years, because these viruses are very popular in bat populations through our longitudinal surveilances.
- 2) We have data to demonastrate the WIV1 (Yang et al., JVI, 2016) and RsSHC014 (data unpublished) can use both human and civet ACE2. Can you please write some comments on these data?

Thanks,

Best regards,

Zhengli,

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**From:** [Fang Li](#)  
**Date:** 2018-04-26 07:05  
**To:** [Zhengli Shi](#)  
**Subject:** SARS manuscript

Dear Zhengli,

Please see attached my portion of the SARS manuscript and also a figure. If you would like me to make changes to the text or figure, please free feel to let me know.

Thanks,