

Supplementary Appendix B: 10,000 vs. 1,000 replications

2017-11-01

```
## Warning in bind_rows(x, .id): binding character and factor vector,  
## coercing into character vector
```

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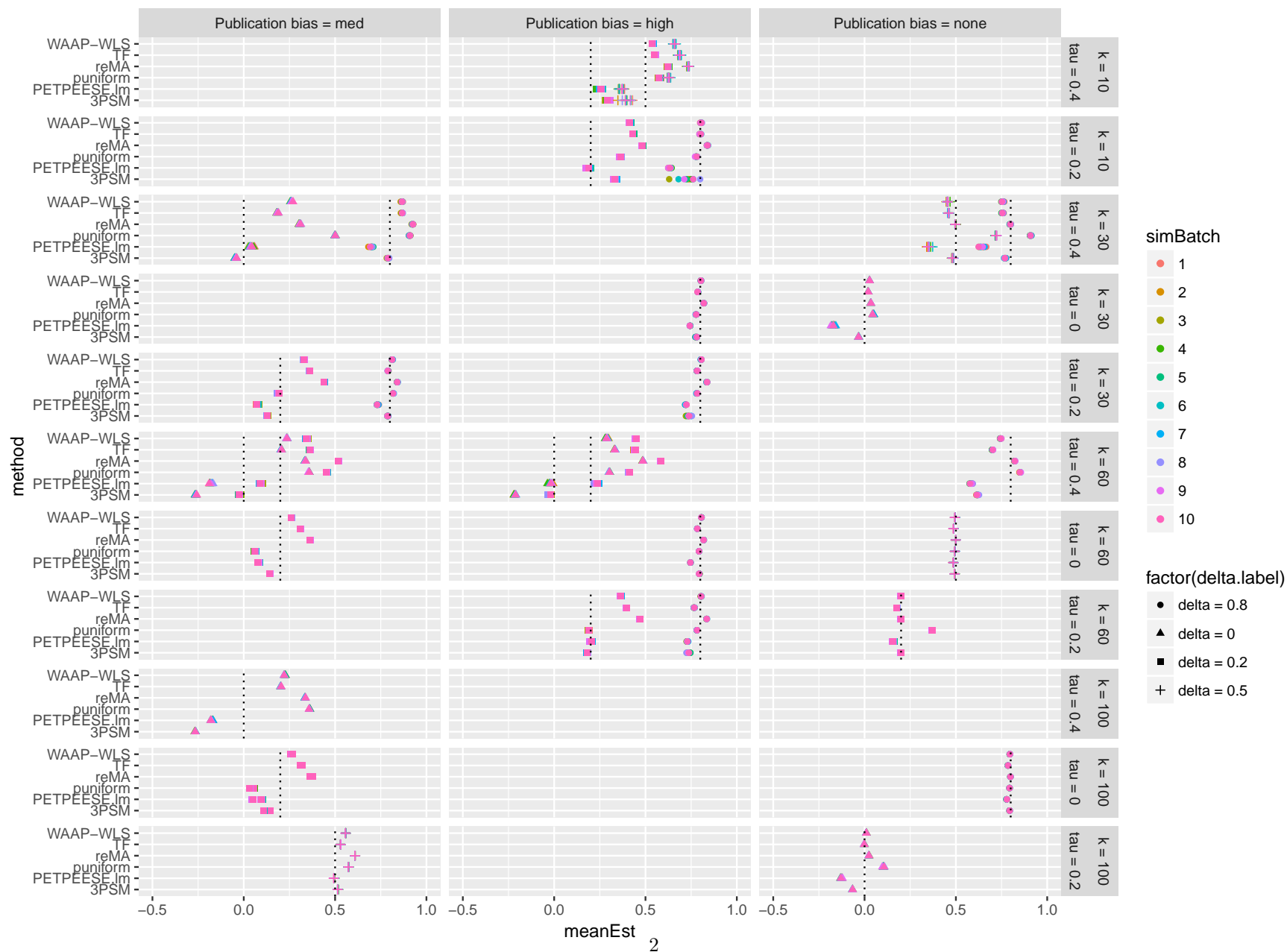
This is Online Appendix B for:

Carter, E. C., Schönbrodt, F. D., Hilgard, J., & Gervais, W. (2017). *Correcting for bias in psychology: A comparison of meta-analytic methods*. Retrieved from <https://osf.io/rf3ys/>.

Method

Simulations were run with 10,000 replications for a selection of conditions. Then, these 10,000 runs were divided in 10 batches of 1000 simulations each. The following plots show points for each of the 10 batches. One can see that they are virtually identical, as the points of the different batches overlap.

Variance in ES estimates between simulation batches



Variance in RMSE between simulation batches

