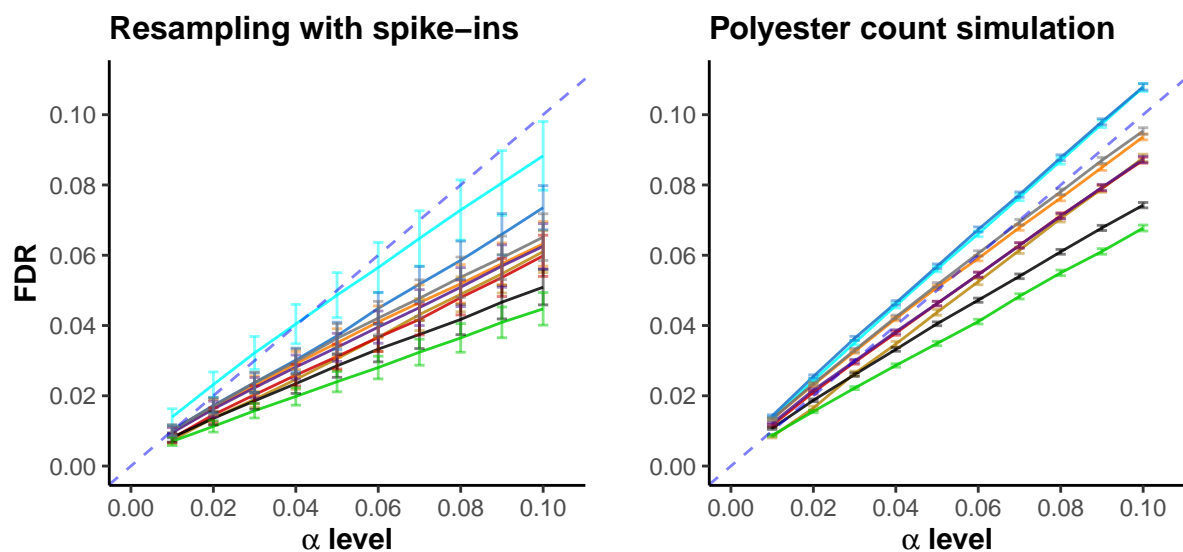
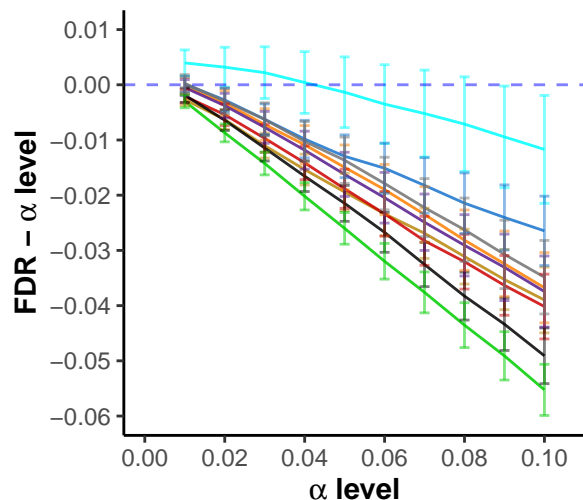


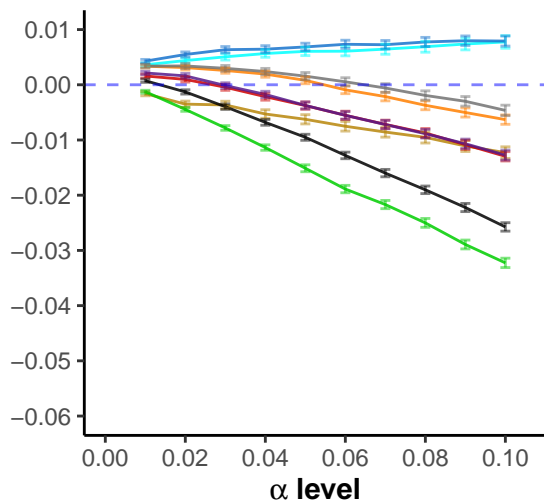
# A FDR control in RNA-seq *in silico* experiments



## B Resampling with spike-ins



## B Polyester count simulation



### Method

- adapt-glm
- adapt-gmm
- ashq
- bh
- bl
- fdrreg-t
- ihw
- lfdr
- qvalue

## C FDR across simulation settings ( $\alpha = 0.05$ )

