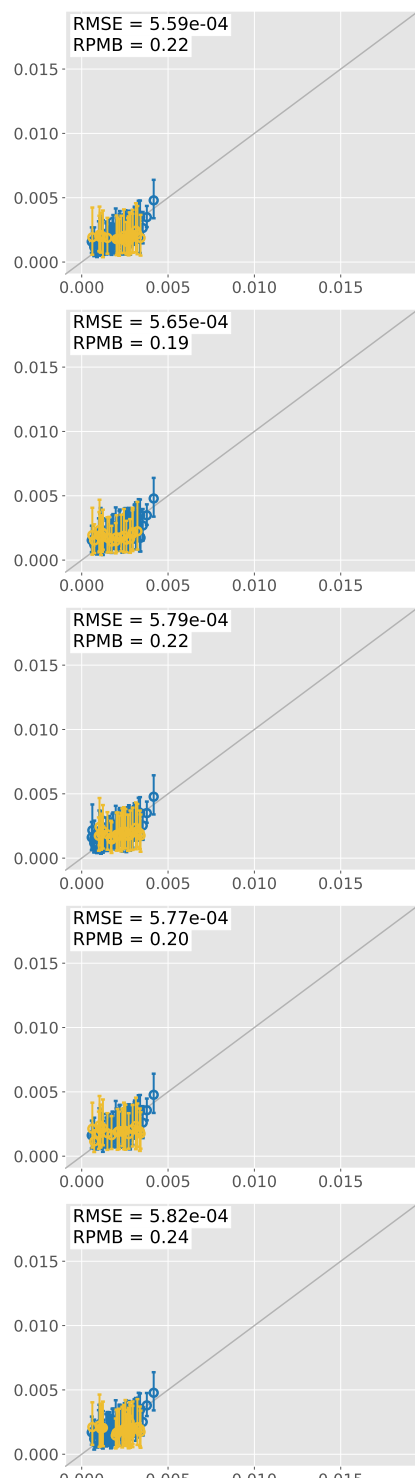


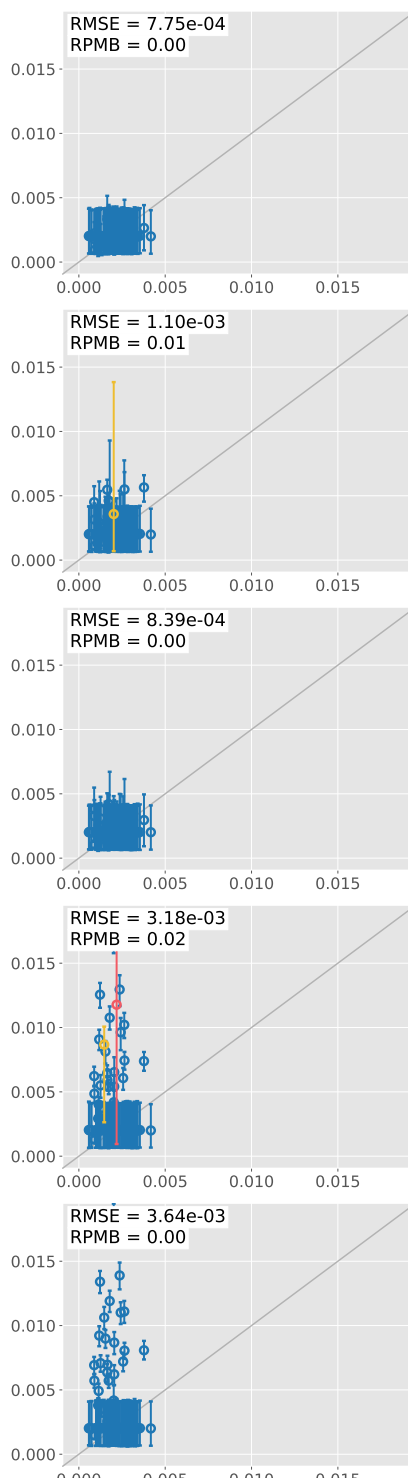
1000bp loci

StarBEAST2

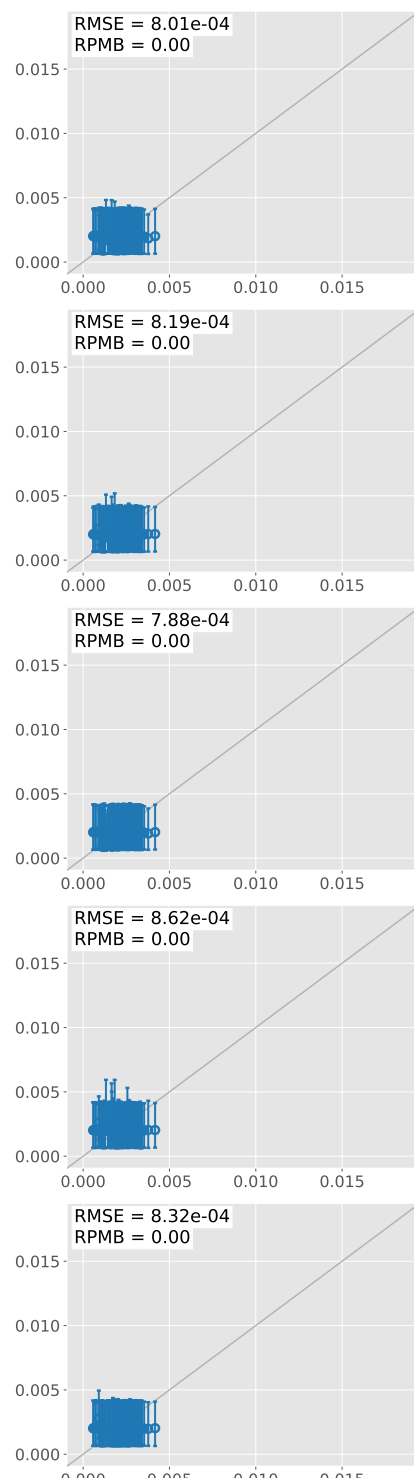
ecoevoity

Estimated ancestral $N_e\mu$ 

all sites



SNPs

True ancestral $N_e\mu$

No errors

20% singleton errors

20% het errors

40% singleton errors

40% het errors