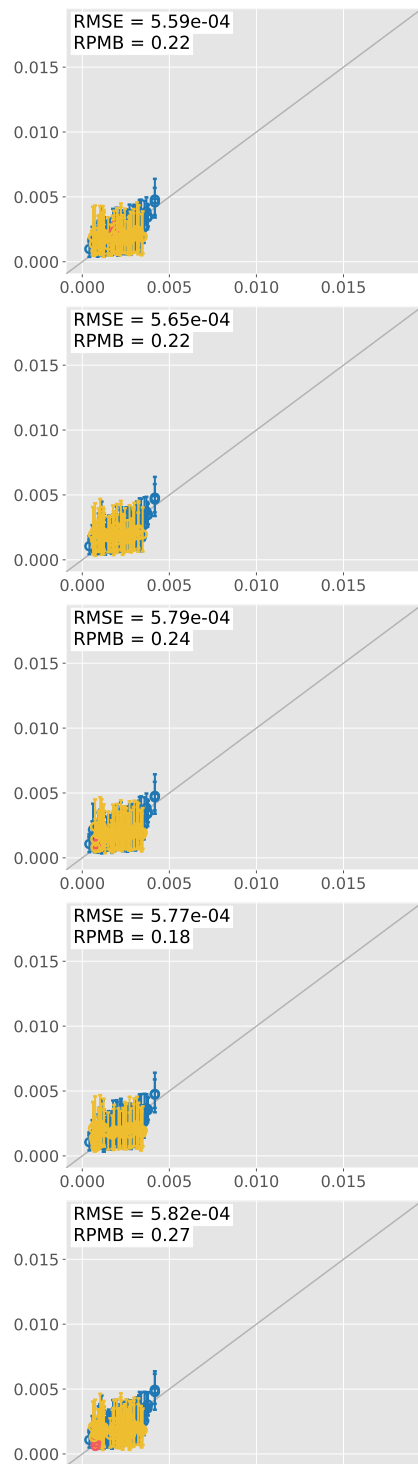


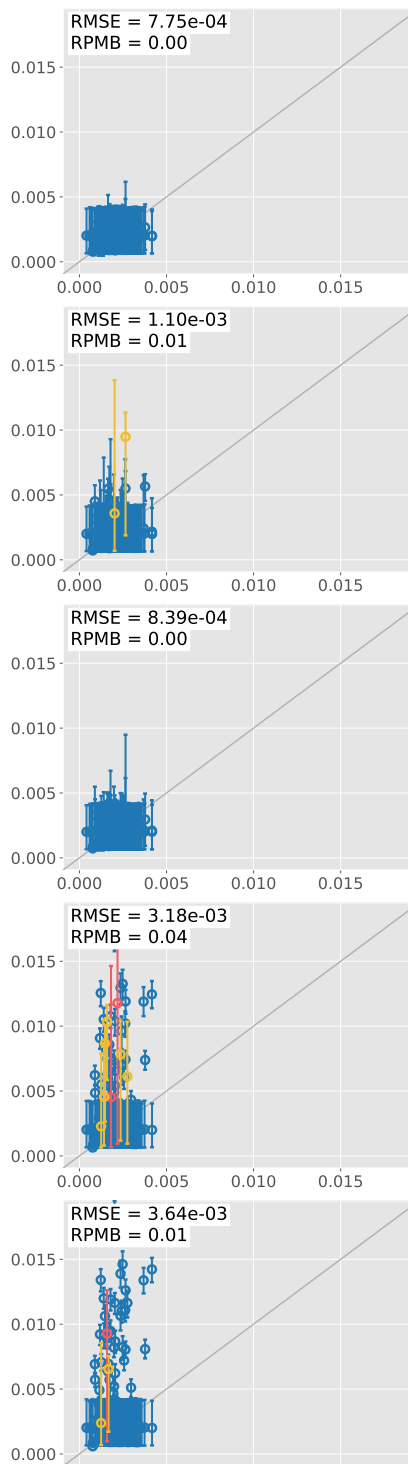
1000bp loci

StarBEAST2

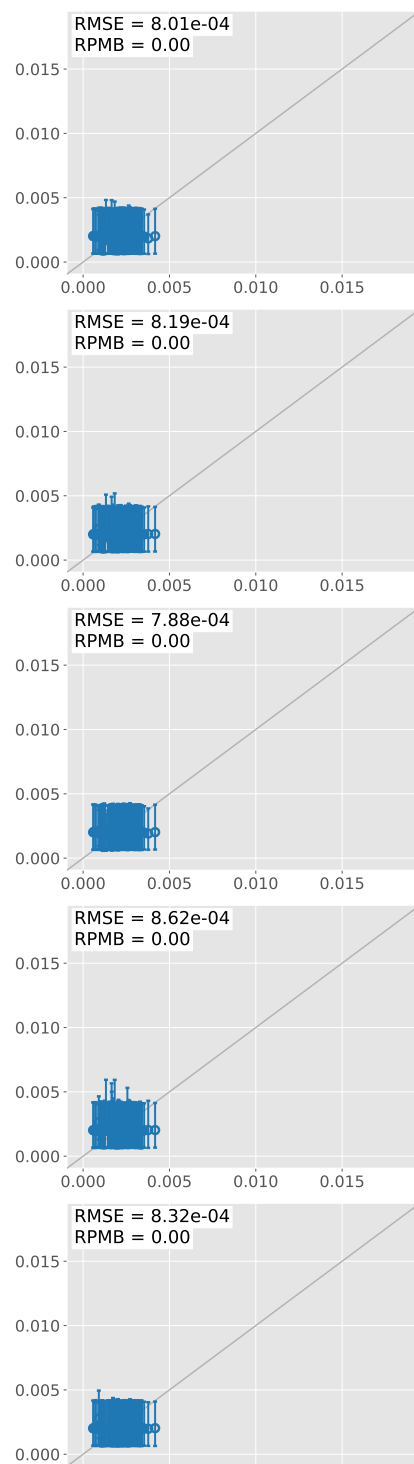
ecoevolity

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