

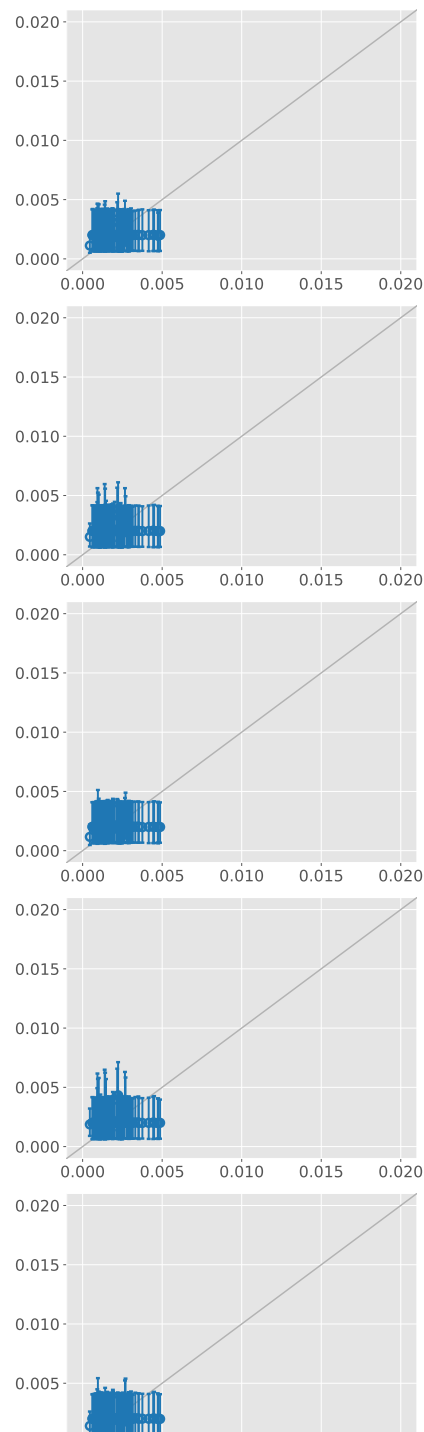
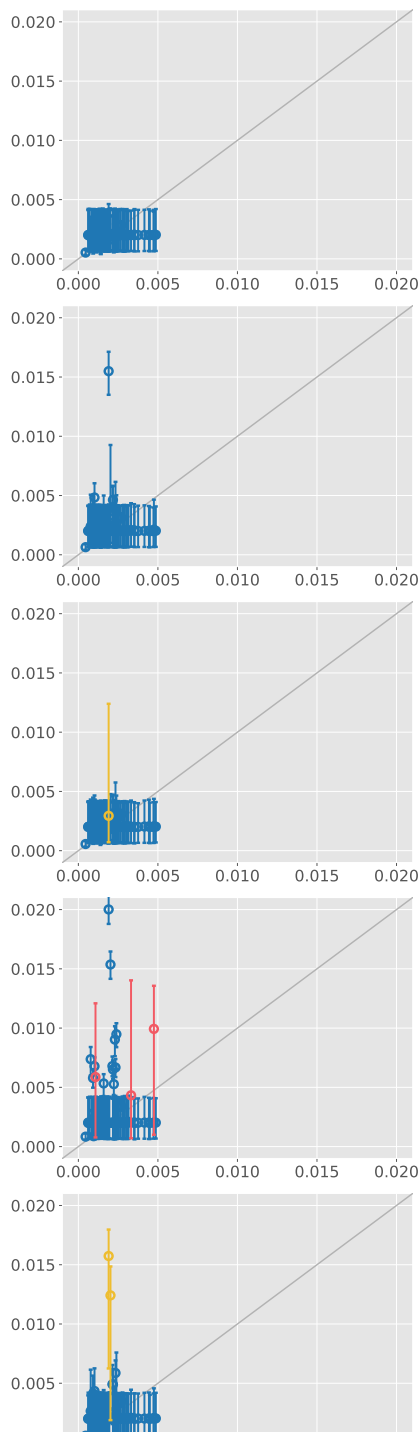
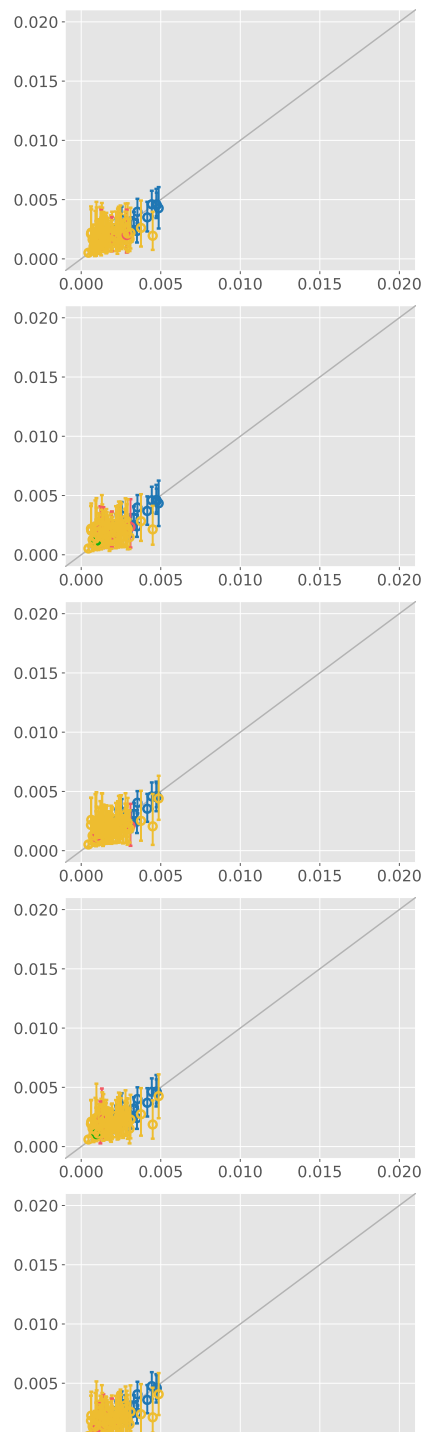
500bp loci

StarBEAST2

ecoevolity

all sites

SNPs

Estimated ancestral $N_e\mu$ 

No errors

20% singleton errors

20% het errors

40% singleton errors

40% het errors

True ancestral $N_e\mu$