

250bp loci

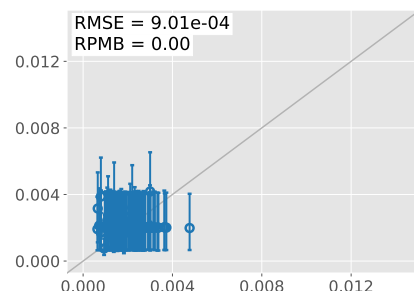
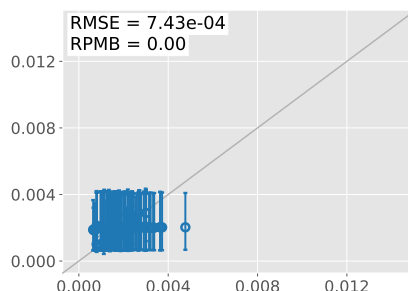
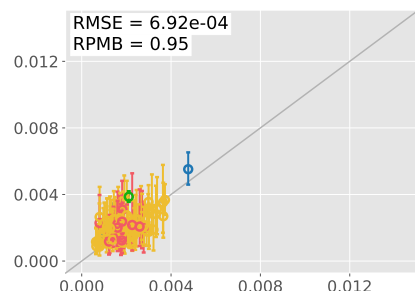
StarBEAST2

ecoevolity

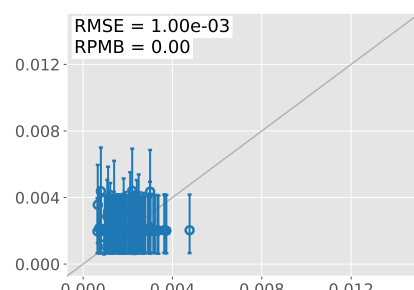
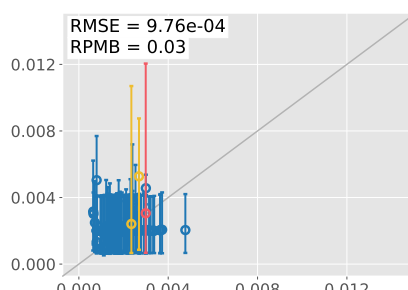
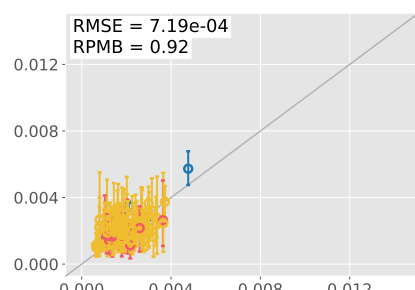
Estimated ancestral $N_e\mu$

all sites

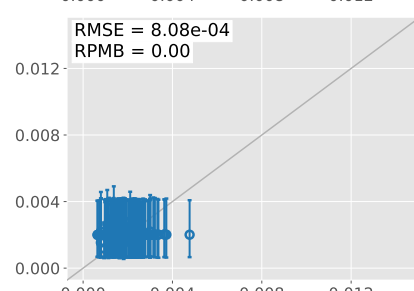
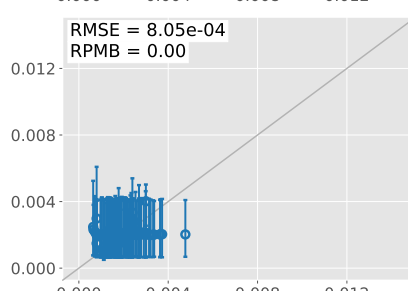
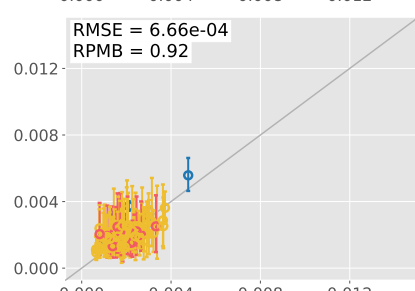
SNPs



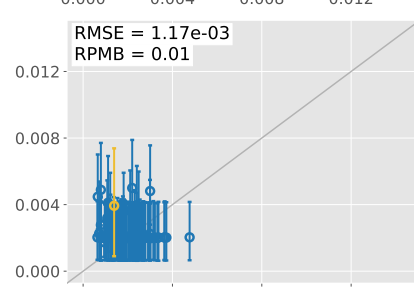
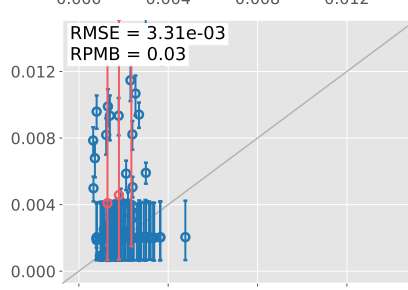
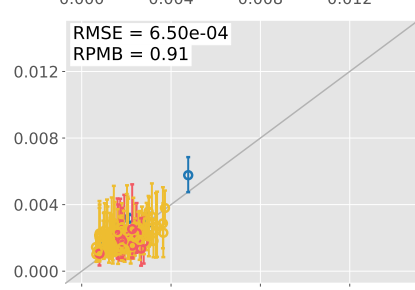
No errors



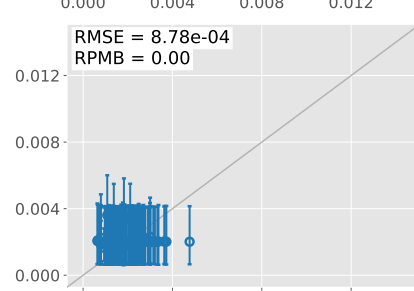
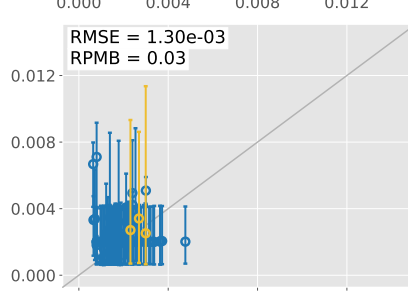
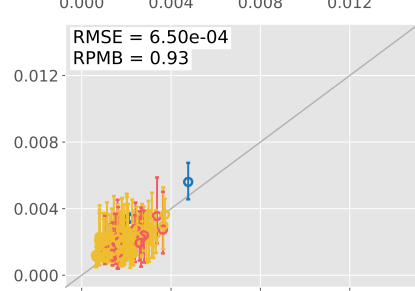
20% singleton errors



20% het errors



40% singleton errors



40% het errors

True ancestral $N_e\mu$