

250bp loci

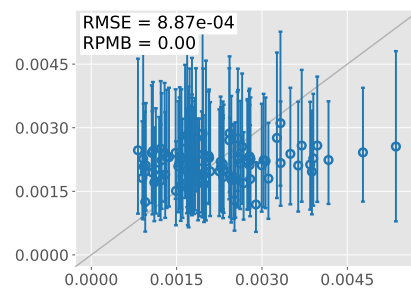
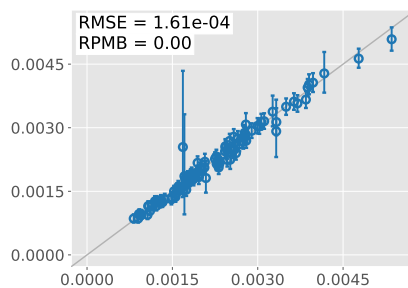
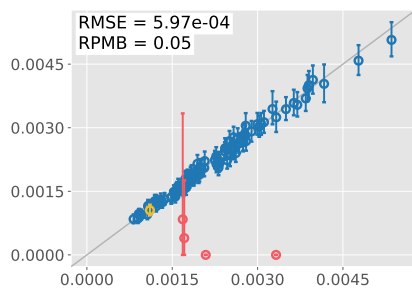
StarBEAST2

ecoevoly

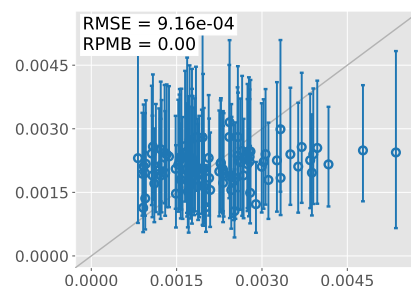
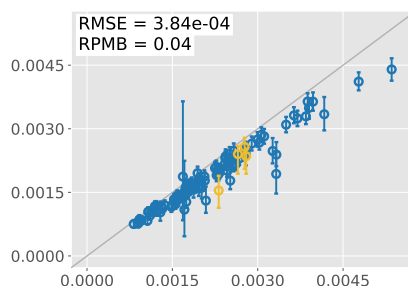
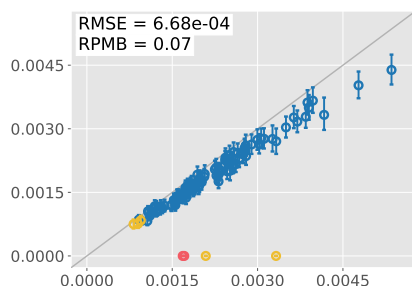
Estimated descendant $N_e\mu$

all sites

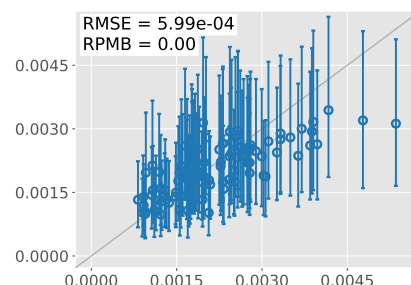
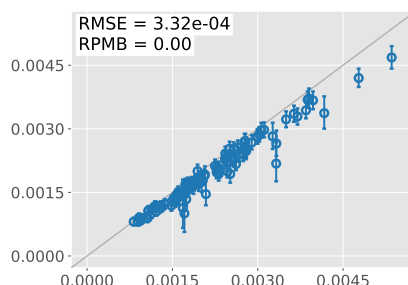
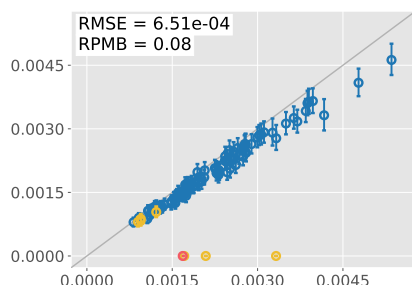
SNPs



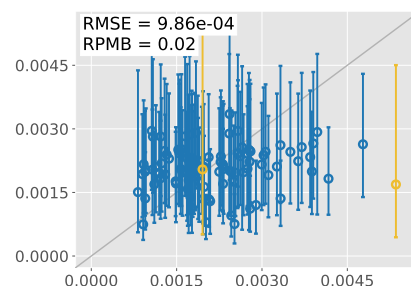
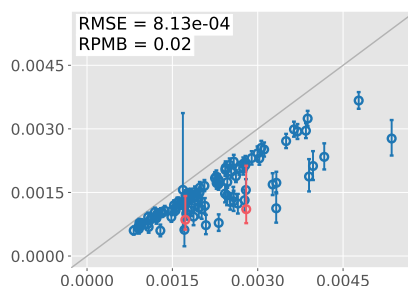
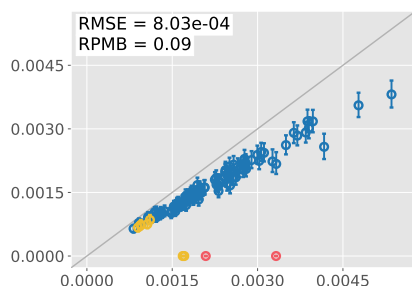
No errors



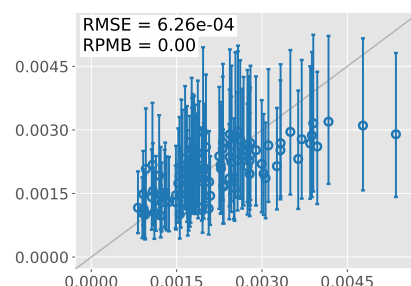
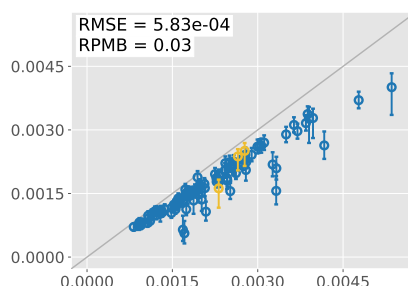
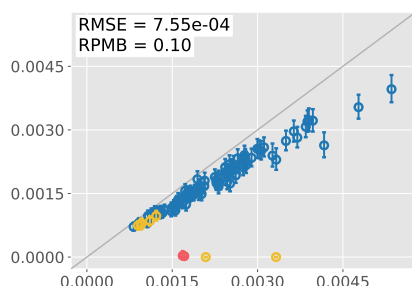
20% singleton errors



20% het errors



40% singleton errors



40% het errors

True descendant $N_e\mu$