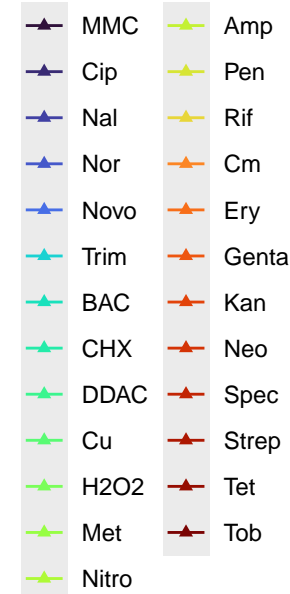
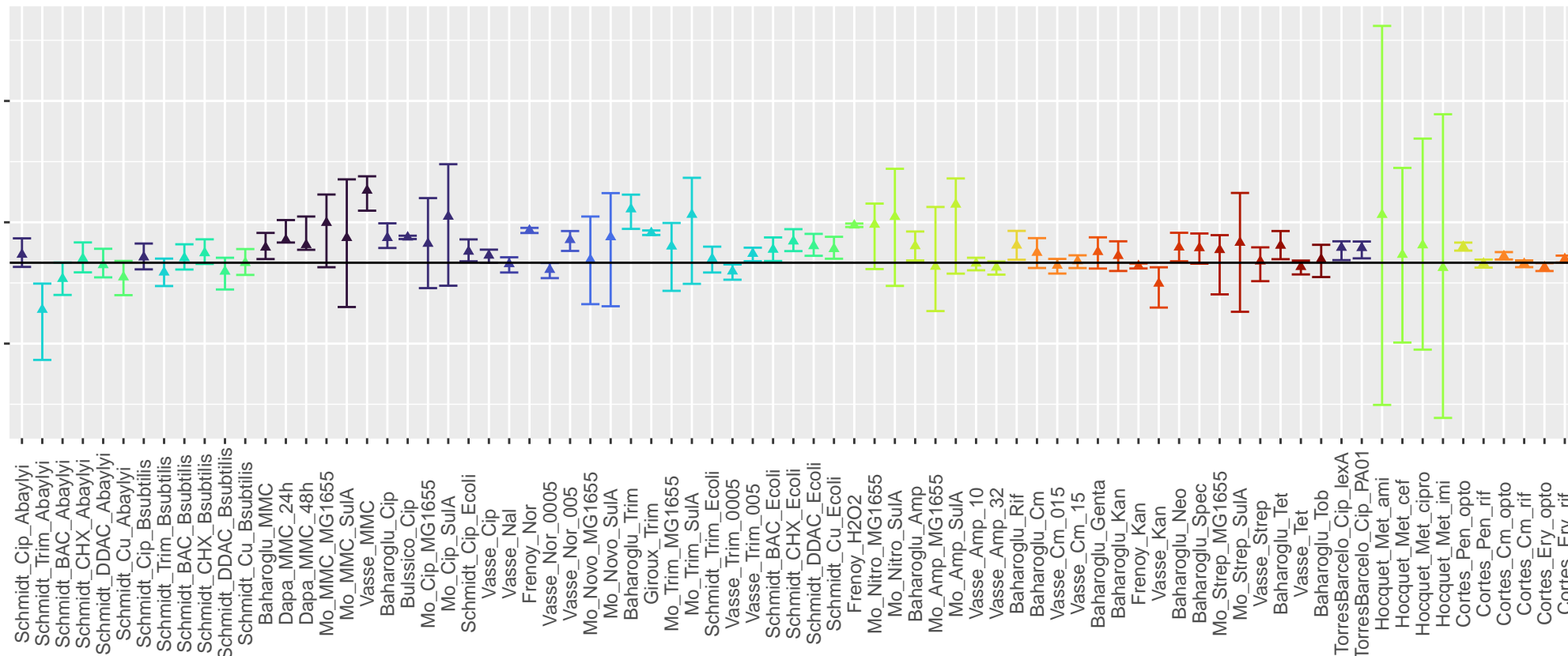


Estimated using model HOM0

Antimicrobial (abbr)



Fold-change population-wide mutation rate



Experiment ID