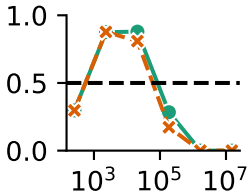
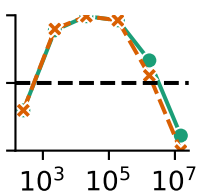


hypermutator
fixation
probability

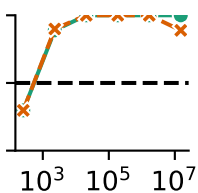
available
beneficial
mutations = 1



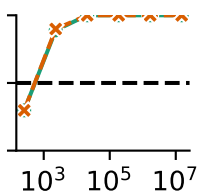
available
beneficial
mutations = 2



available
beneficial
mutations = 3



available
beneficial
mutations = 4



well-mixed



2D demes