

mutator
fixation
probability

available
beneficial
mutations = 1



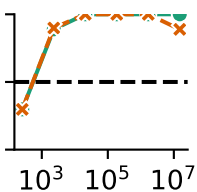
population size

available
beneficial
mutations = 2



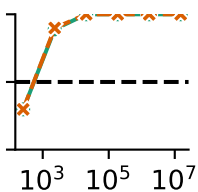
population size

available
beneficial
mutations = 3



population size

available
beneficial
mutations = 4



population size



well-mixed



2D demes