

mutator  
fixation  
probability

available  
beneficial  
mutations = 1



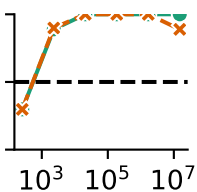
population size

available  
beneficial  
mutations = 2



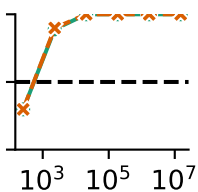
population size

available  
beneficial  
mutations = 3



population size

available  
beneficial  
mutations = 4



population size



well-mixed



2D demes