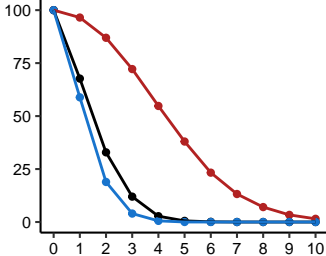
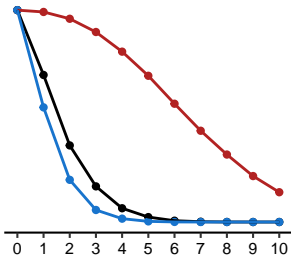


% of cells with mutations

500 input (179 mutations)



800 input (308 mutations)



# of mutations per cell