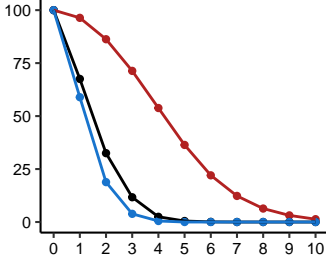
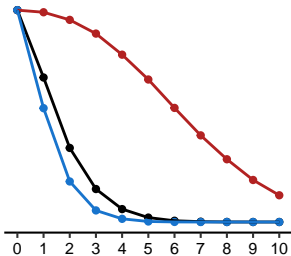


% of cells with mutations

500 input (175 mutations)



800 input (305 mutations)



# of mutations per cell