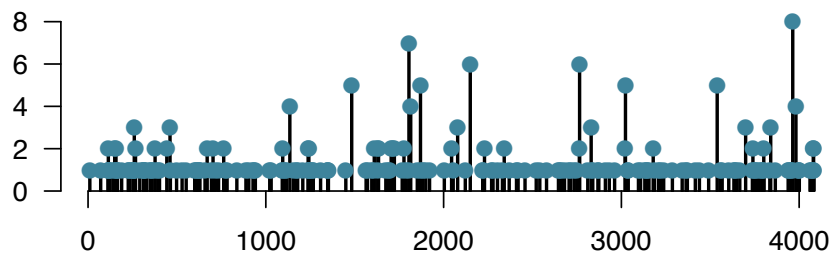
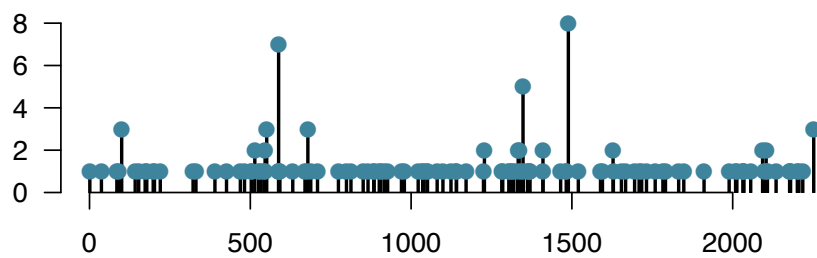


Number of specimens with mutation

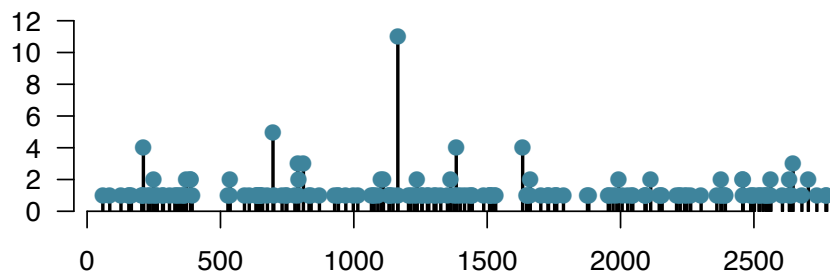
MSH6



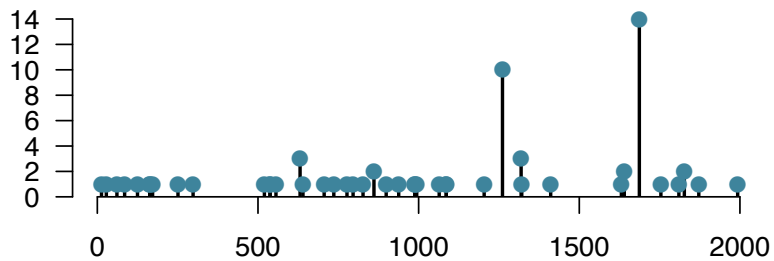
MLH1



MSH2



PMS2



POLE

