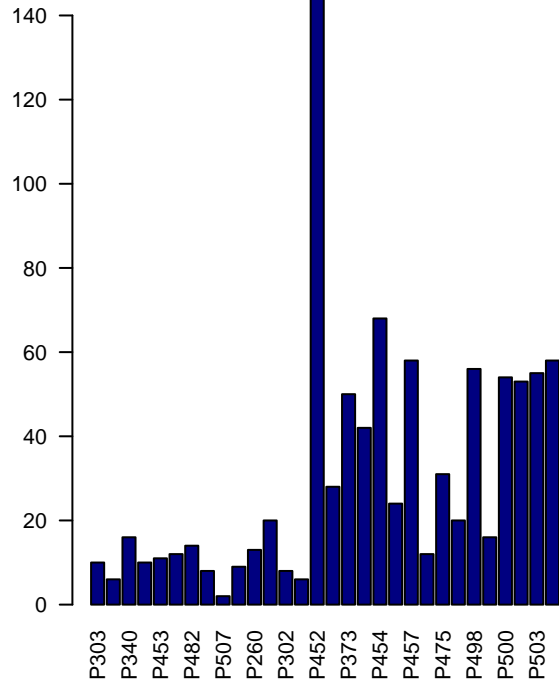
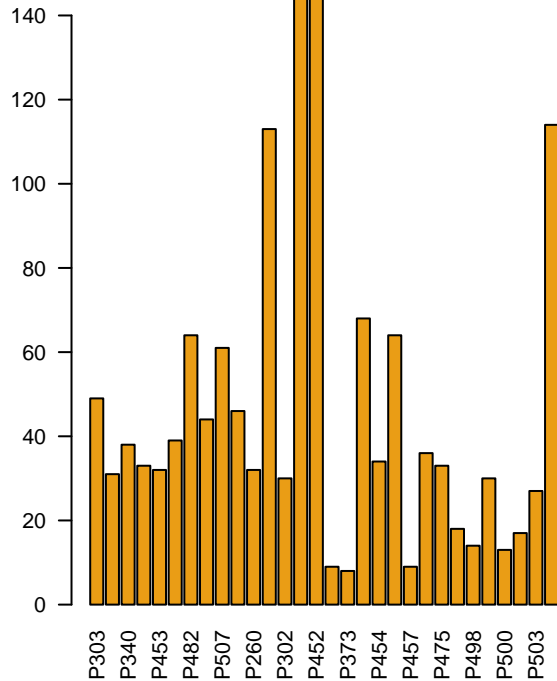


Number of clonal mutations



Number of shared mutations



Number of private mutations

