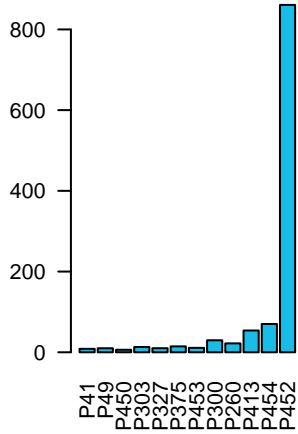
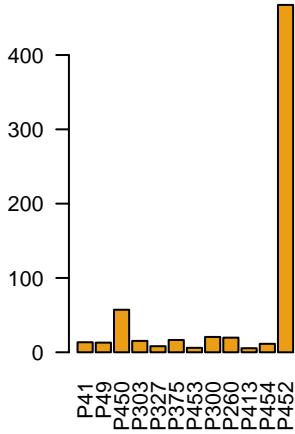


Number of tumor-wide mutations



Number of shared mutations



Number of private mutations

