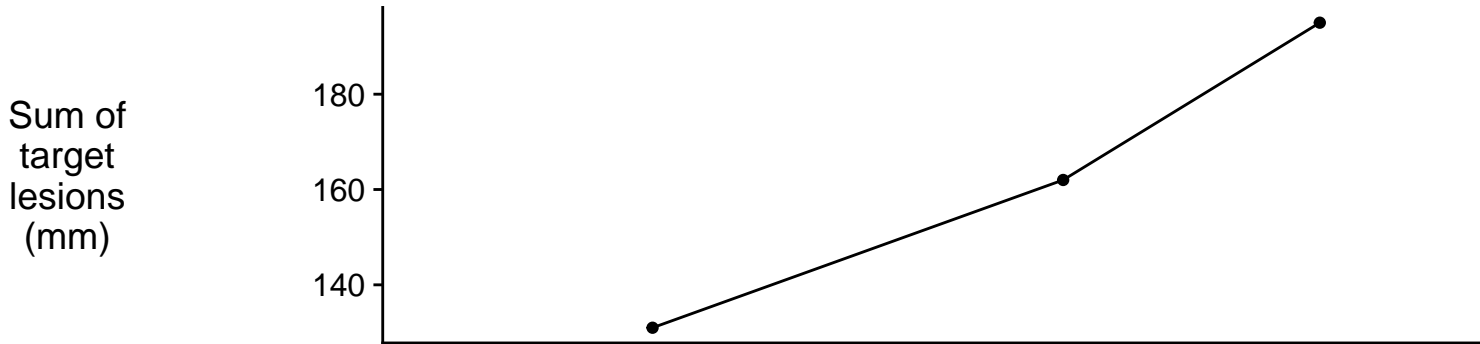
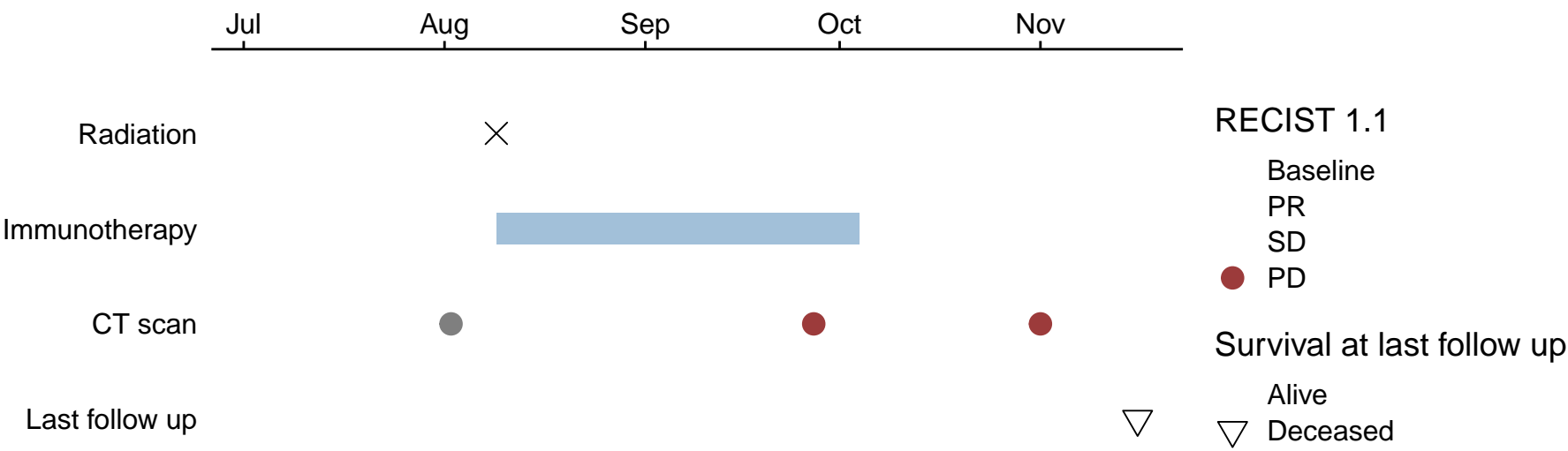
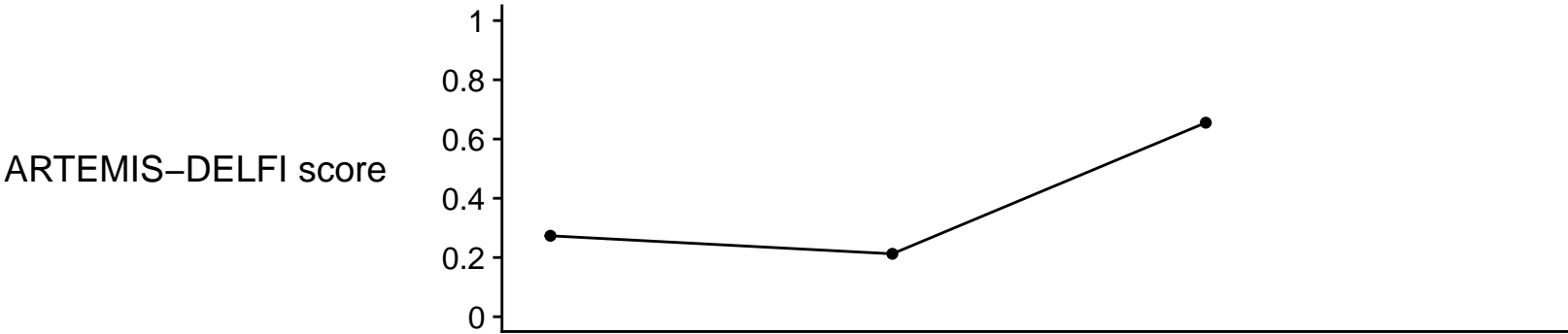


CGPLPA784  
WGS data: Tumor (No), Normal (No), Plasma (3 timepoints)  
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured



WGMAF (%)

Targeted MAF (%)



CA 19-9 (U/mL)

