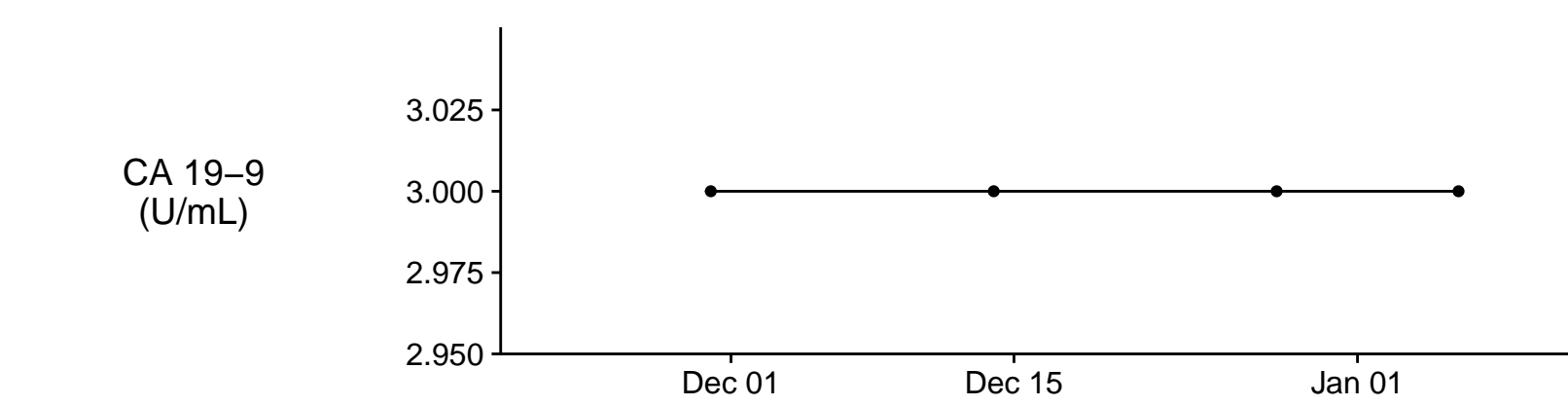
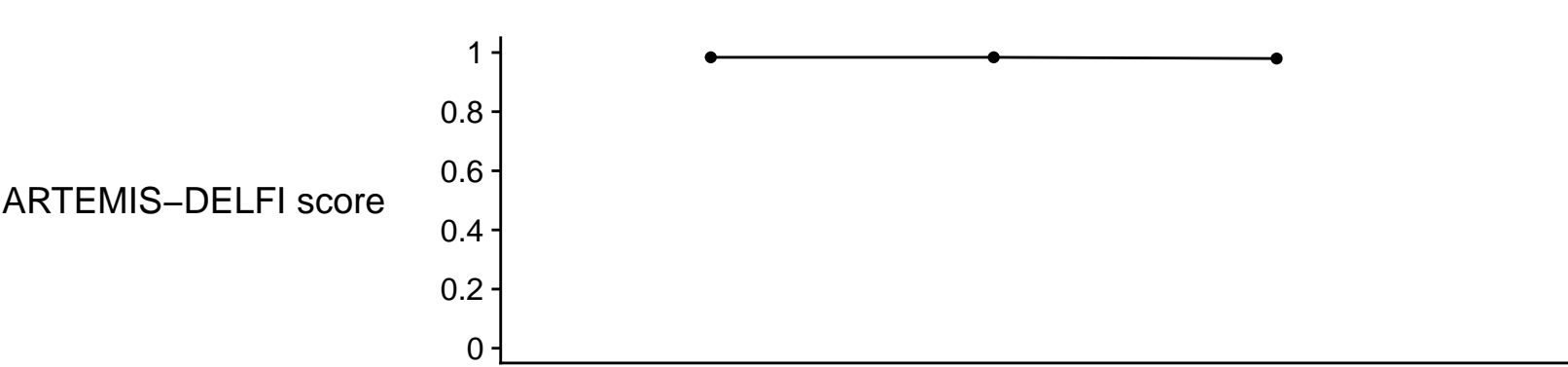
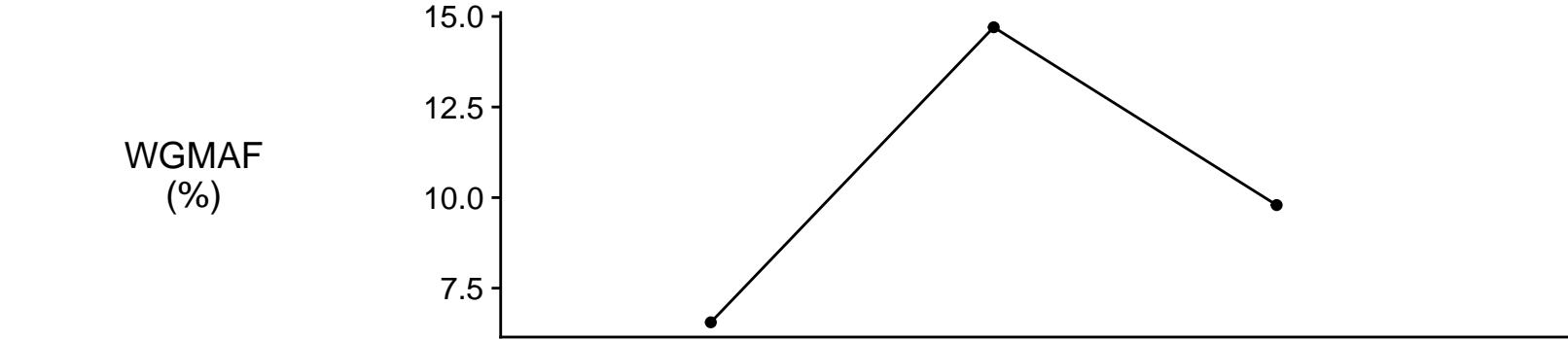
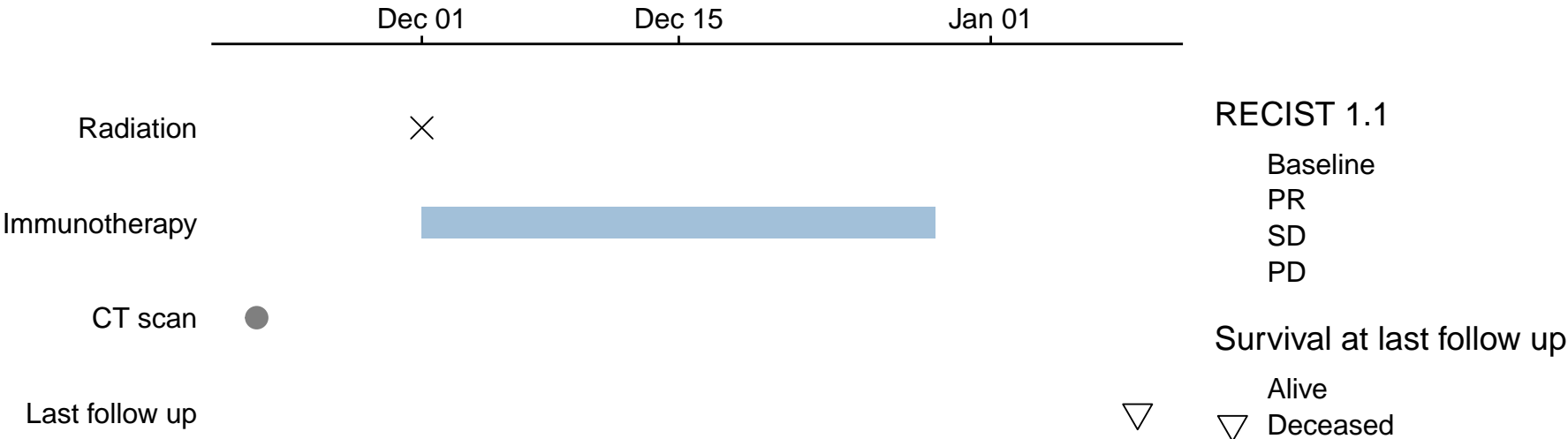
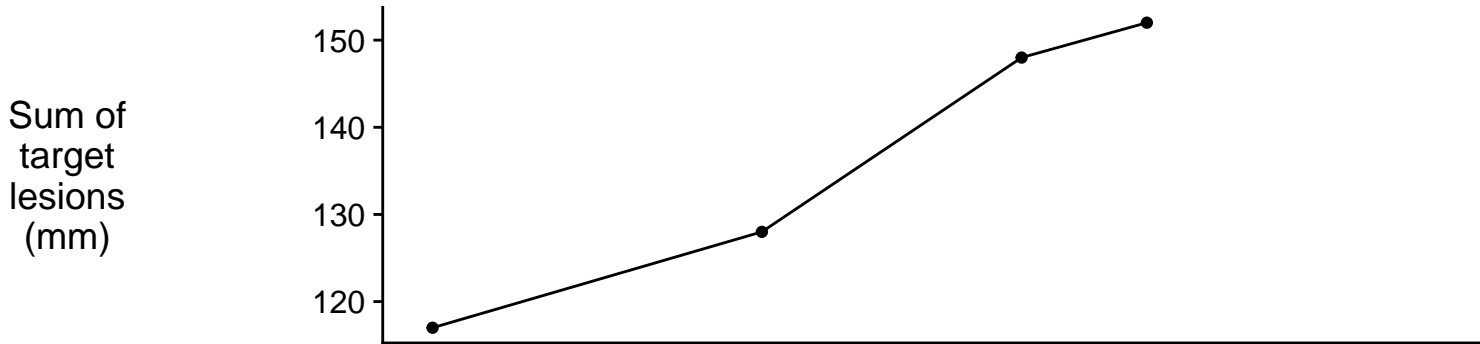
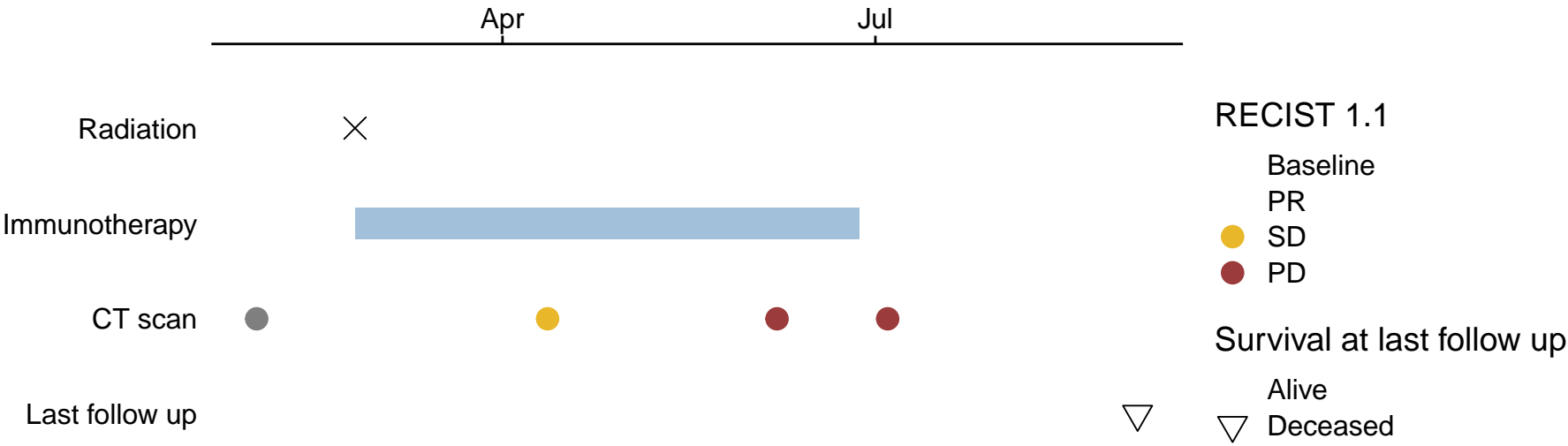


CGPLPA765
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 4494, 1785

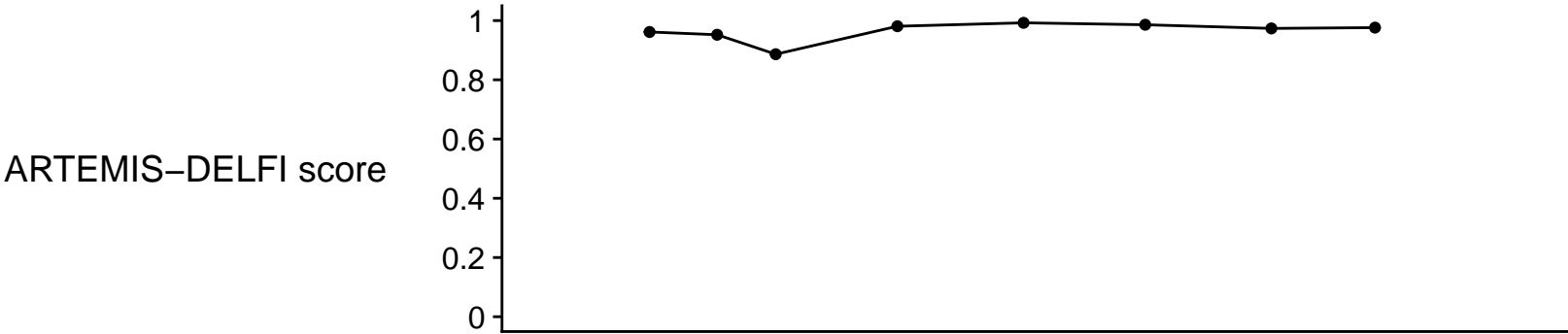


CGPLPA770
WGS data: Tumor (No), Normal (Yes), Plasma (8 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured

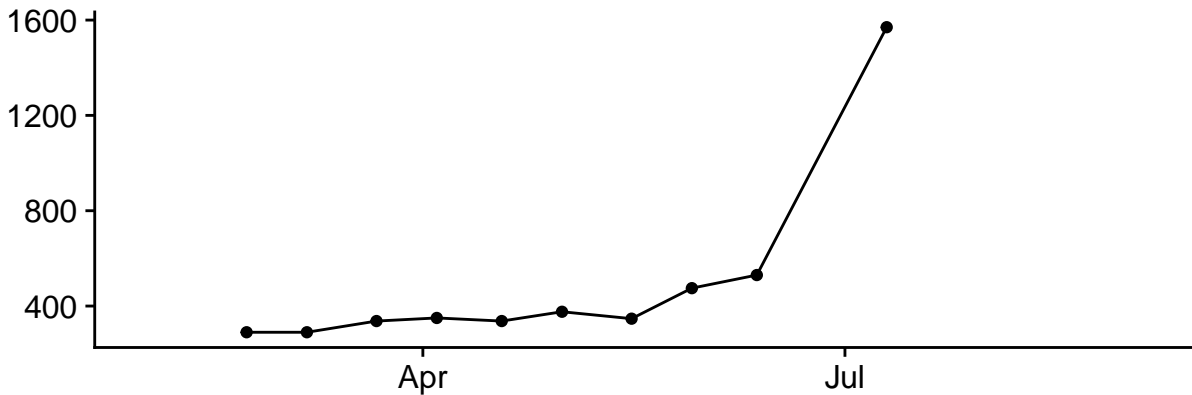


WGMAF (%)

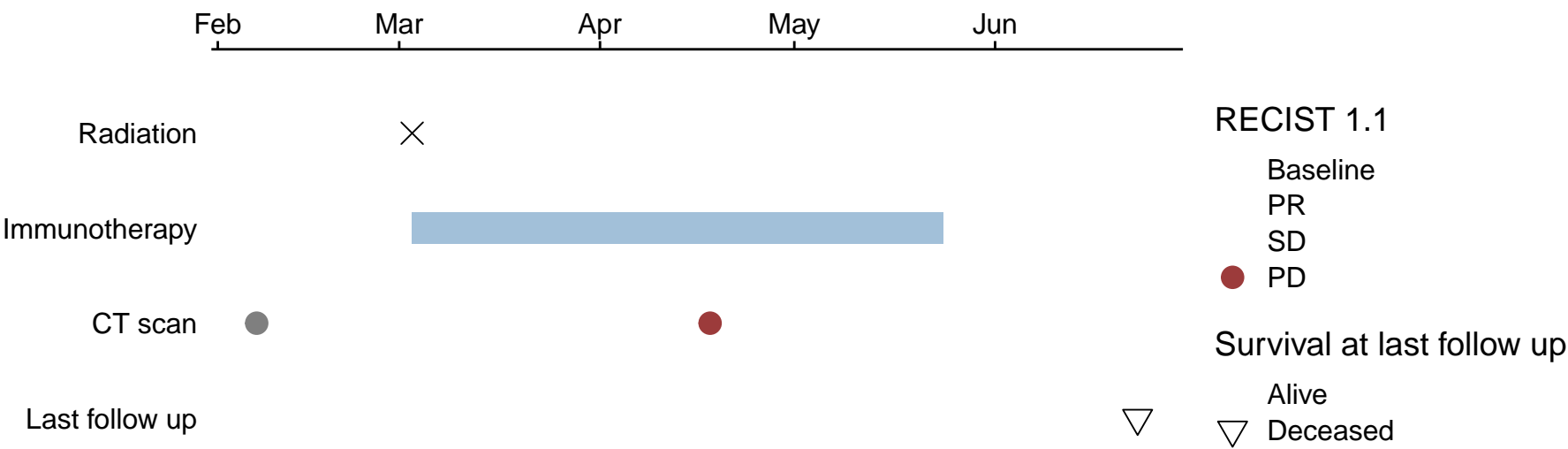
Targeted MAF (%)



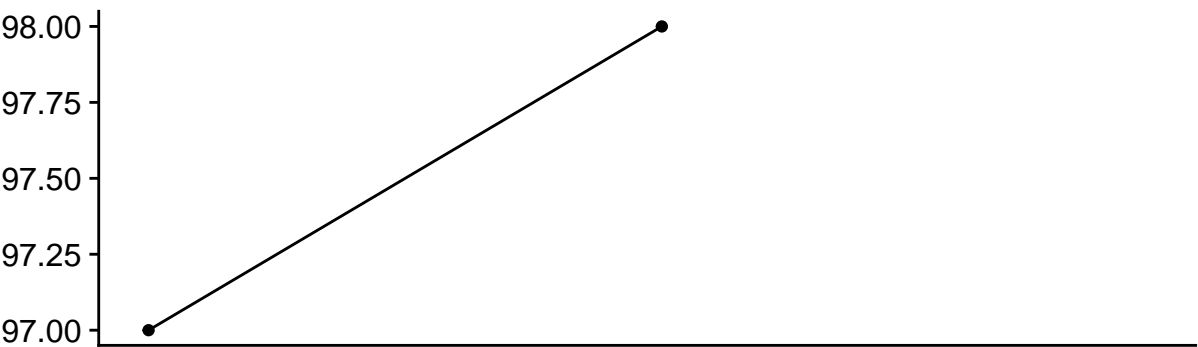
CA 19-9 (U/mL)



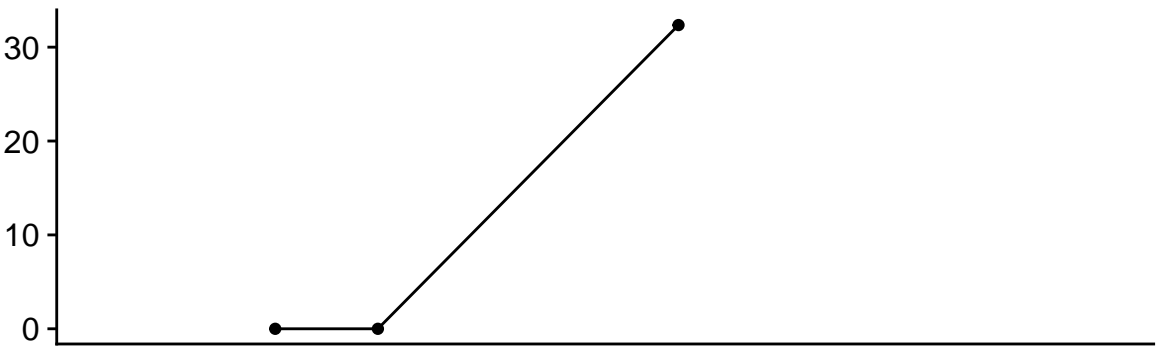
CGPLPA771
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 454, 51



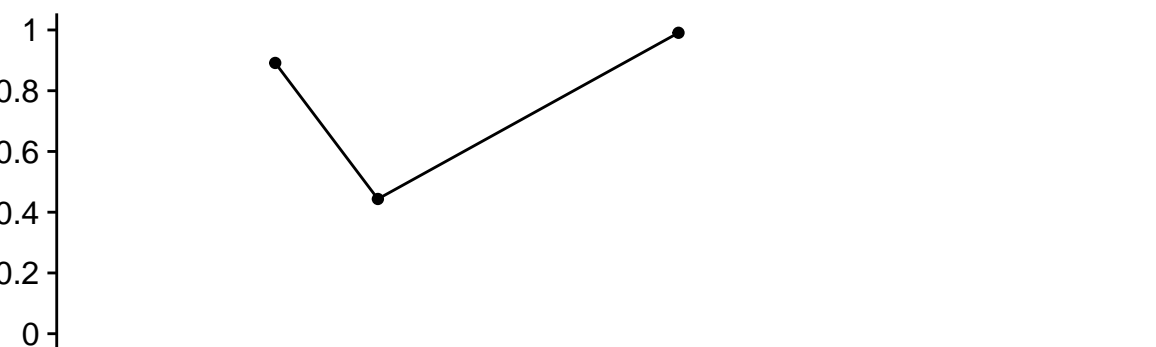
Sum of target lesions (mm)



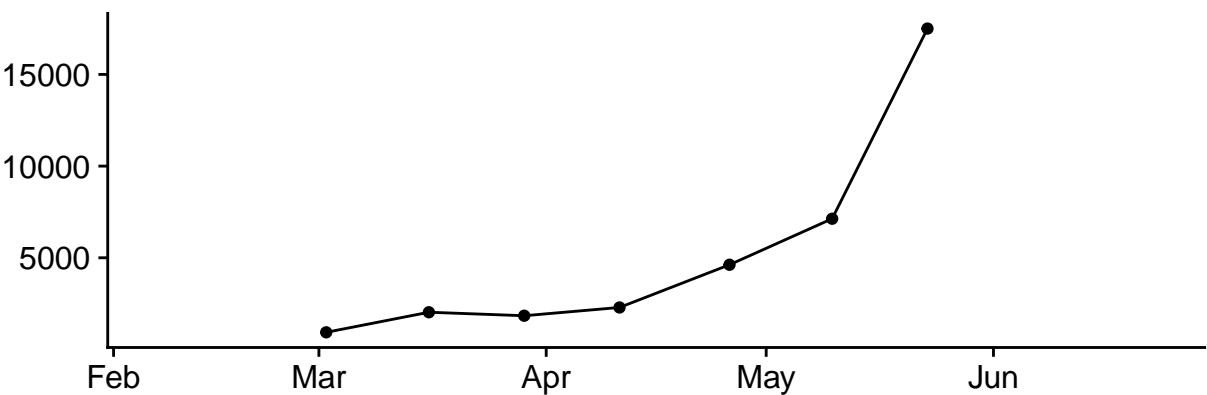
WGMAF (%)



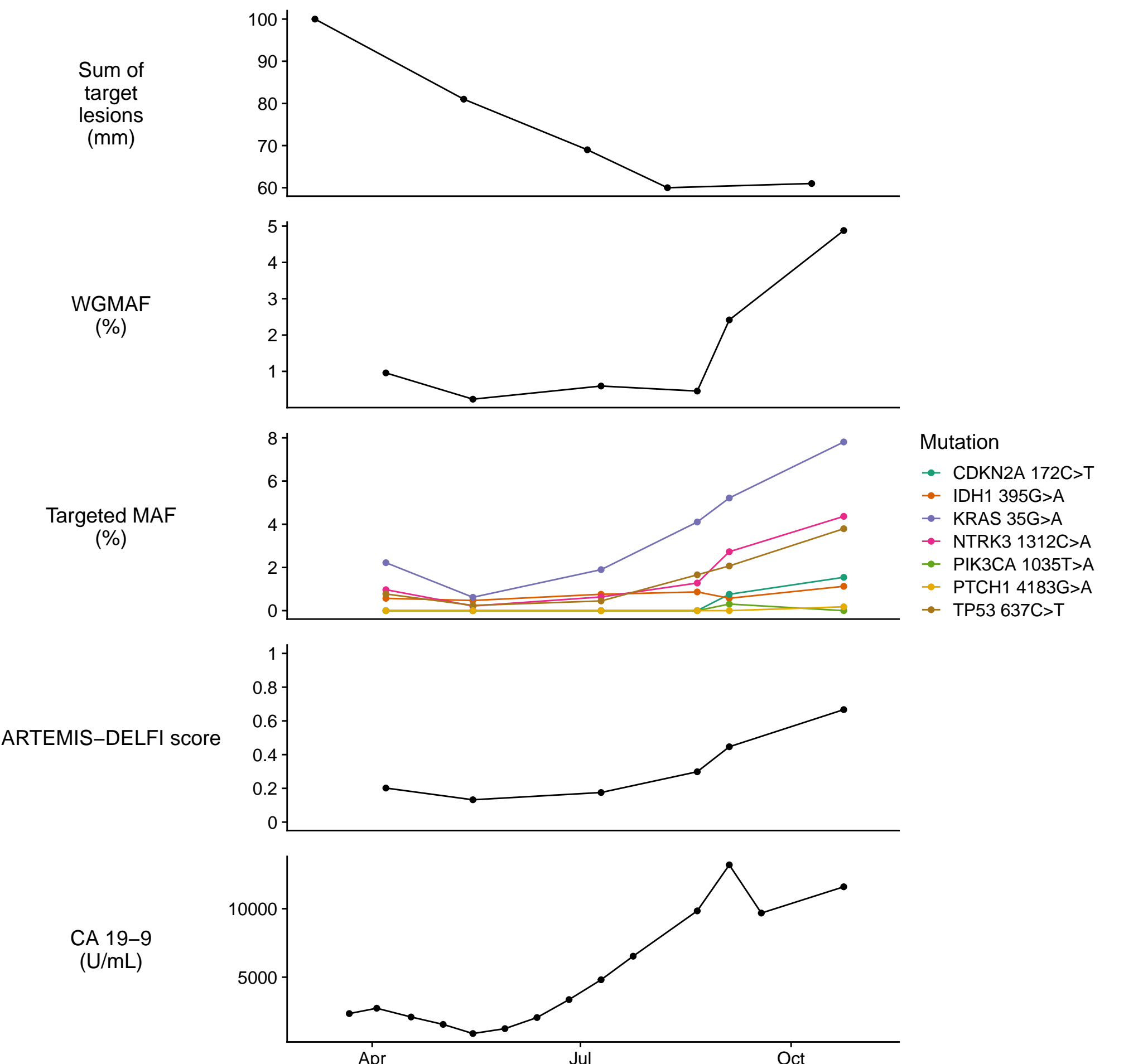
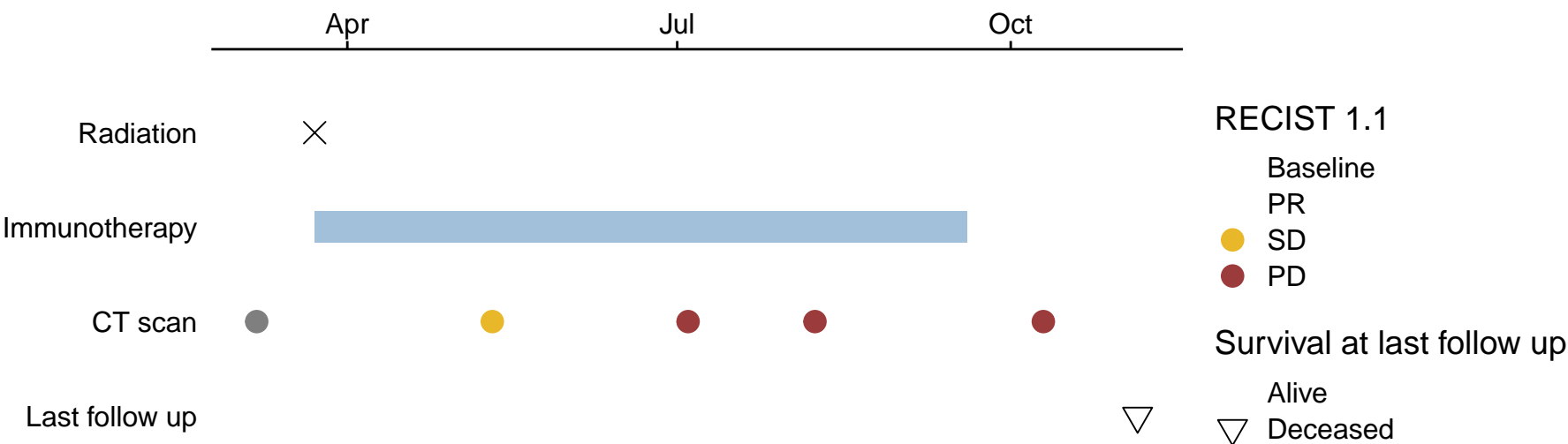
Targeted MAF (%)



ARTEMIS-DELFI score

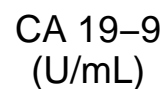
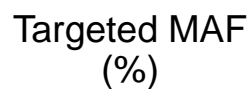
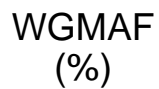
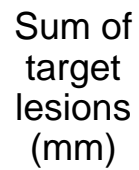


CGPLPA219
WGS data: Tumor (Yes), Normal (Yes), Plasma (6 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 2900, 1984

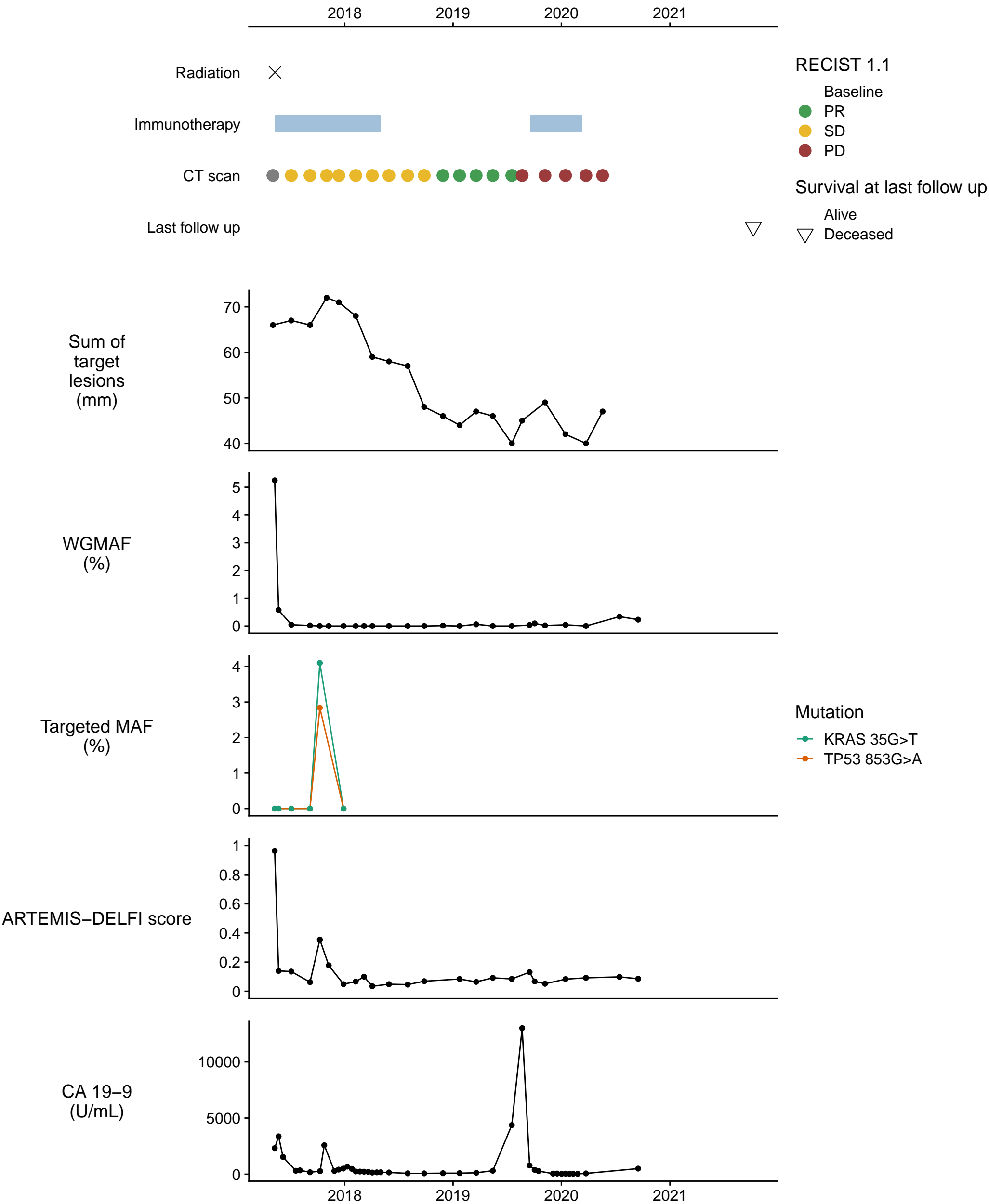


WGS data: Tumor (Yes), Normal (Yes), Plasma (15 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 60061, 44737

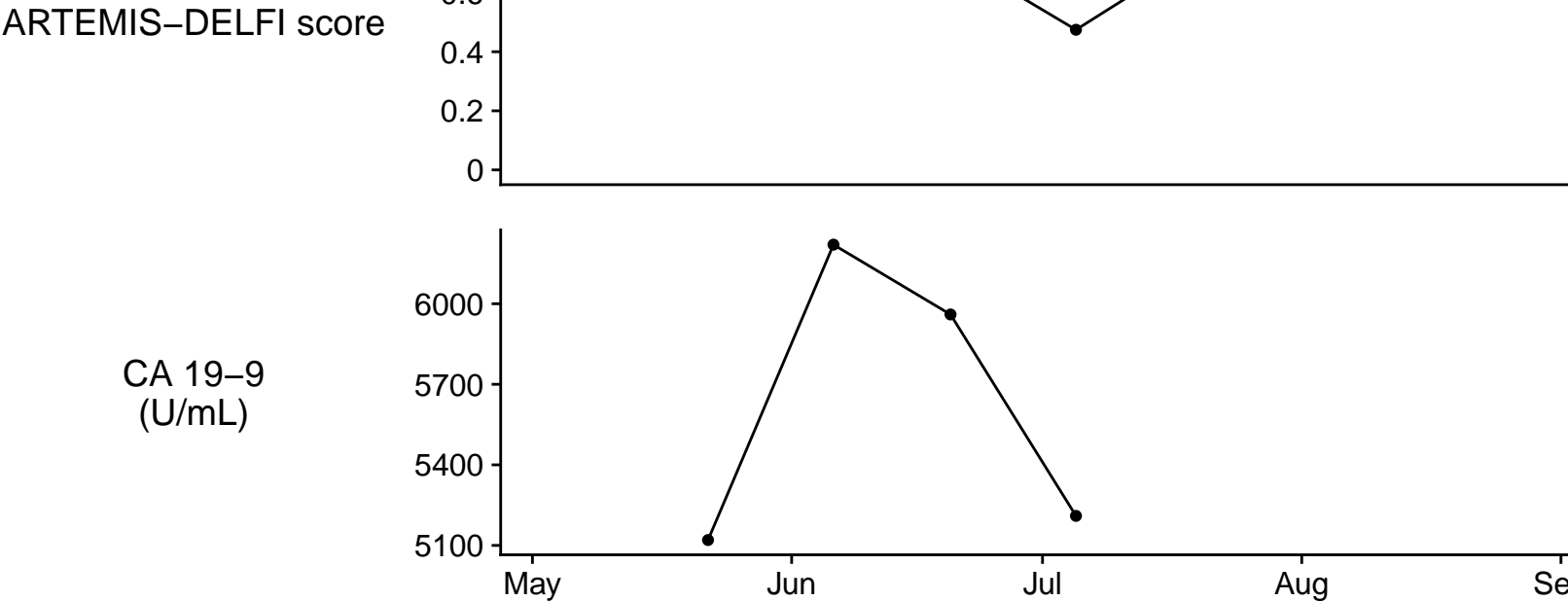
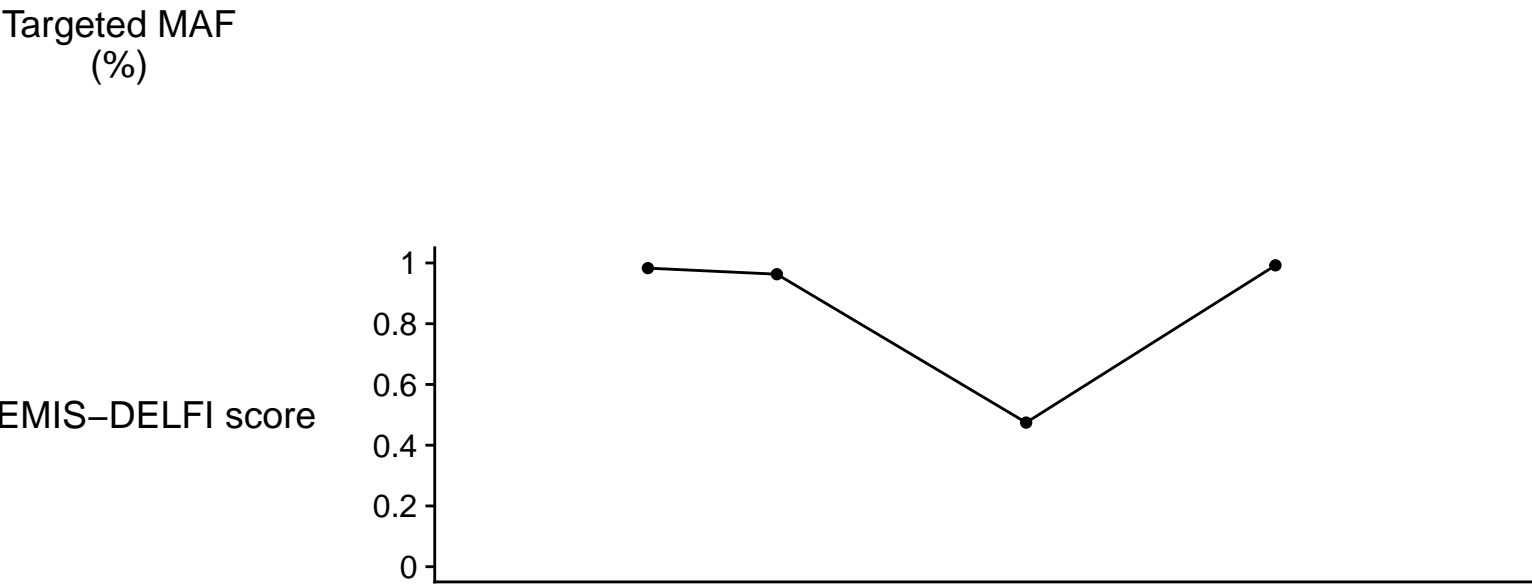
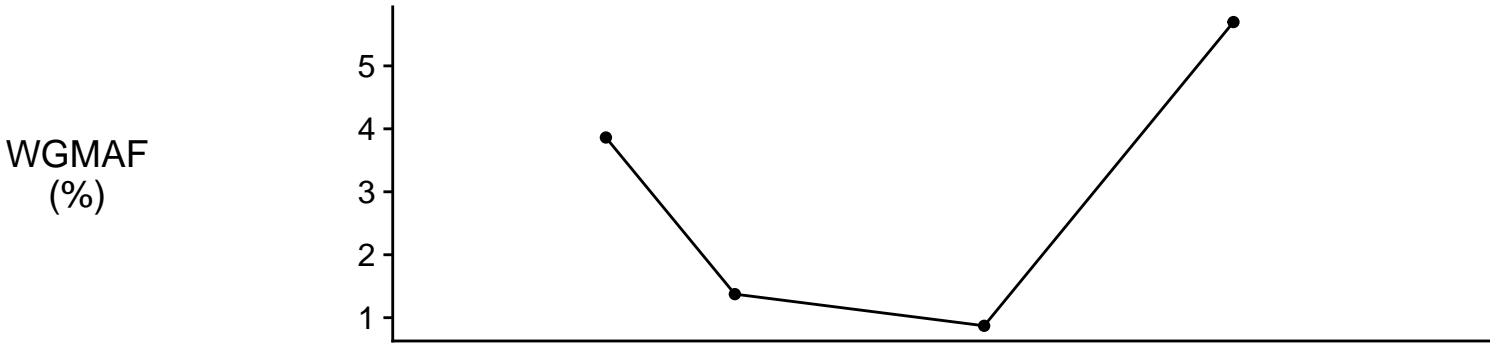
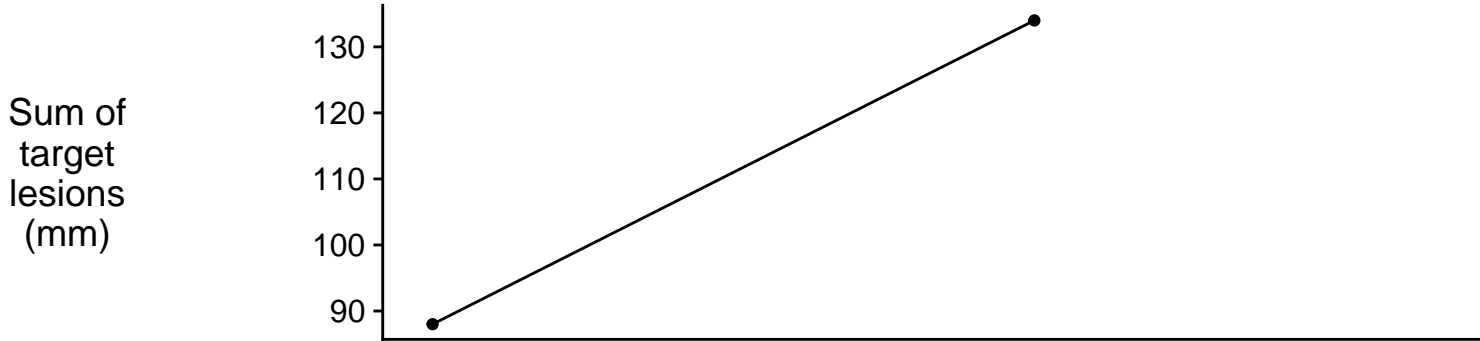
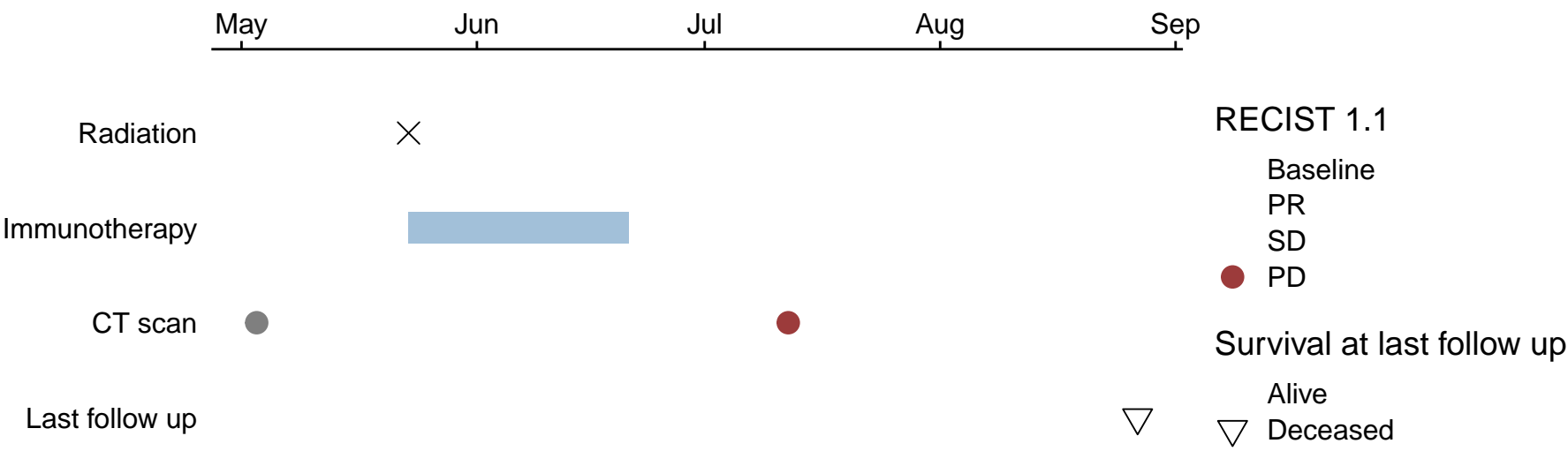
2021



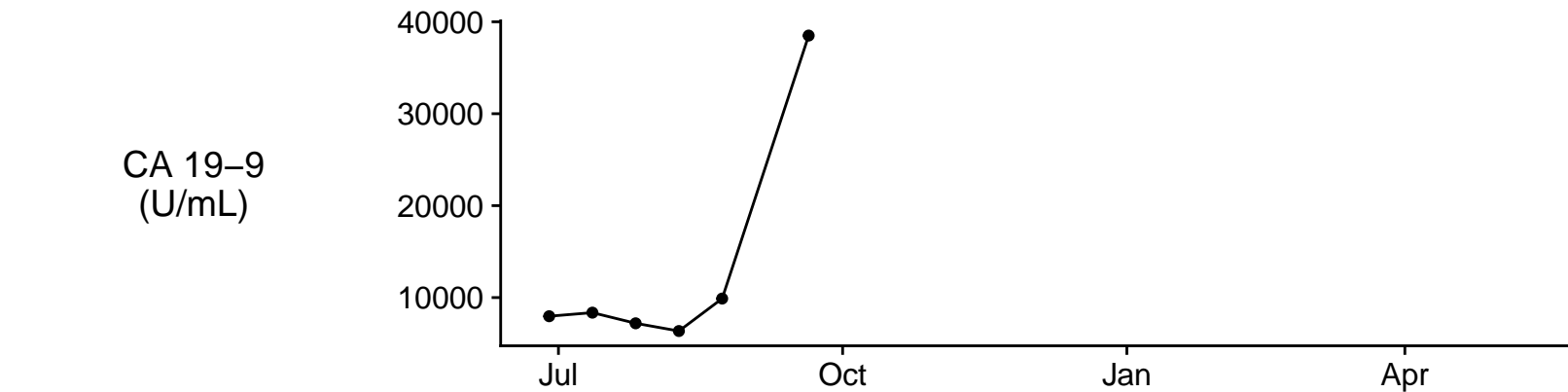
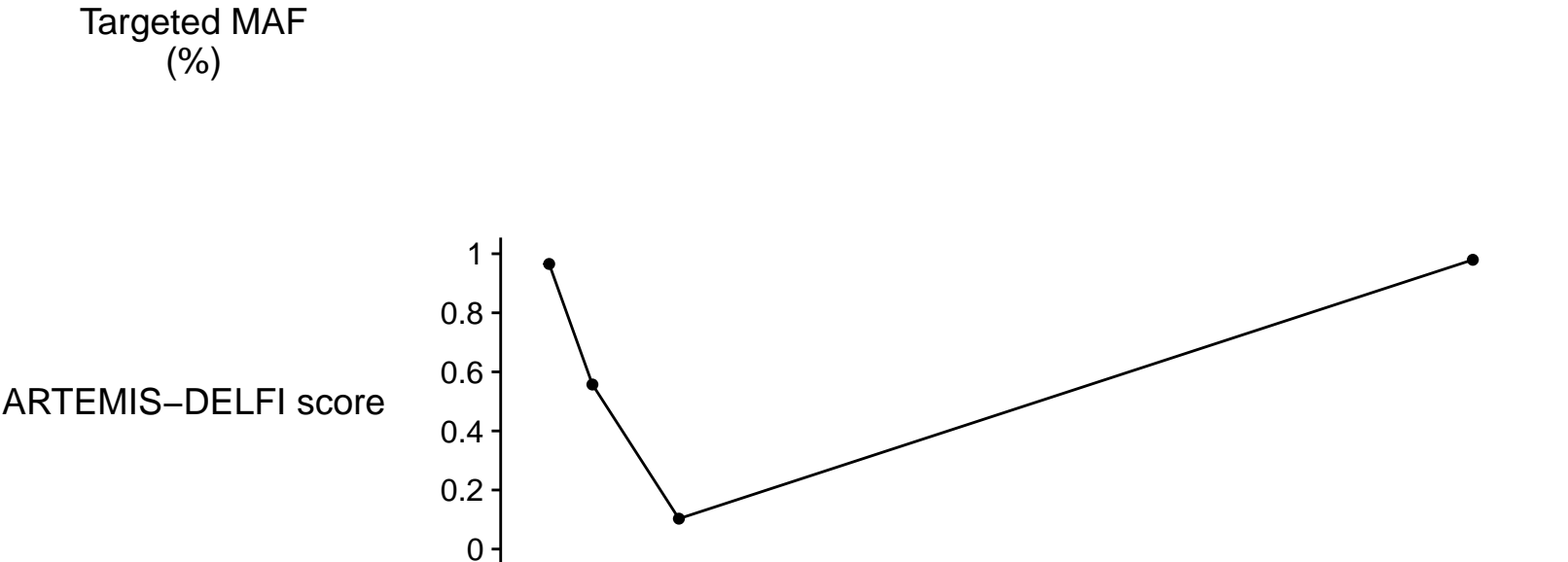
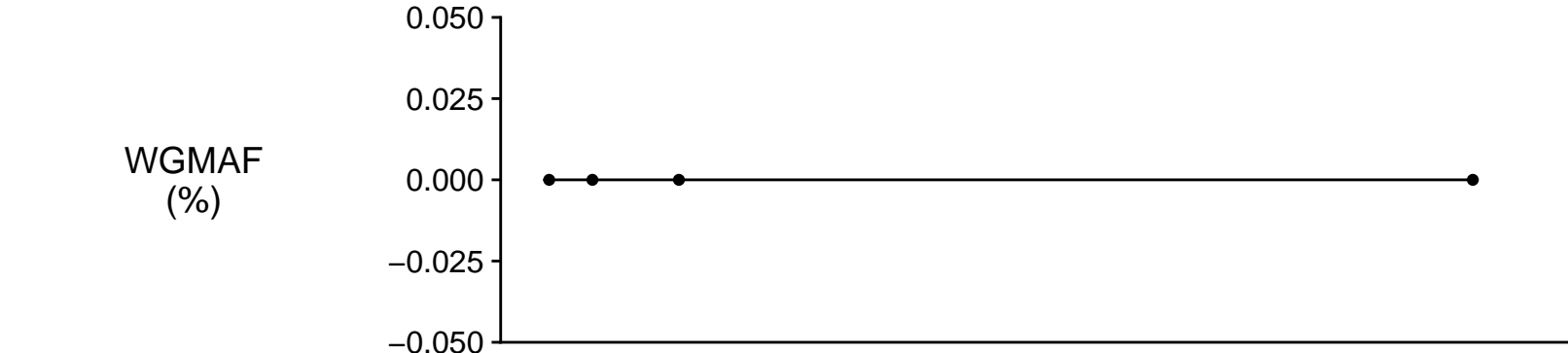
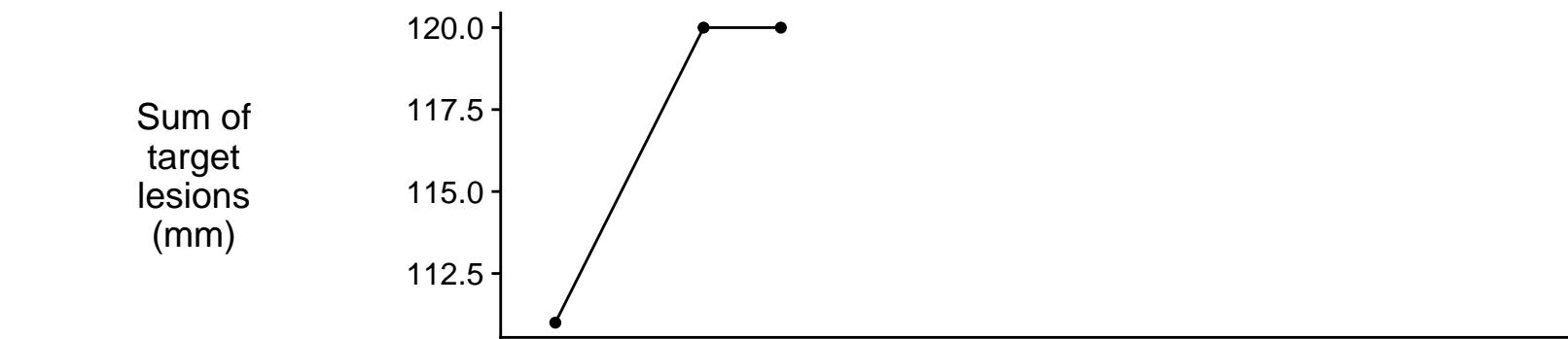
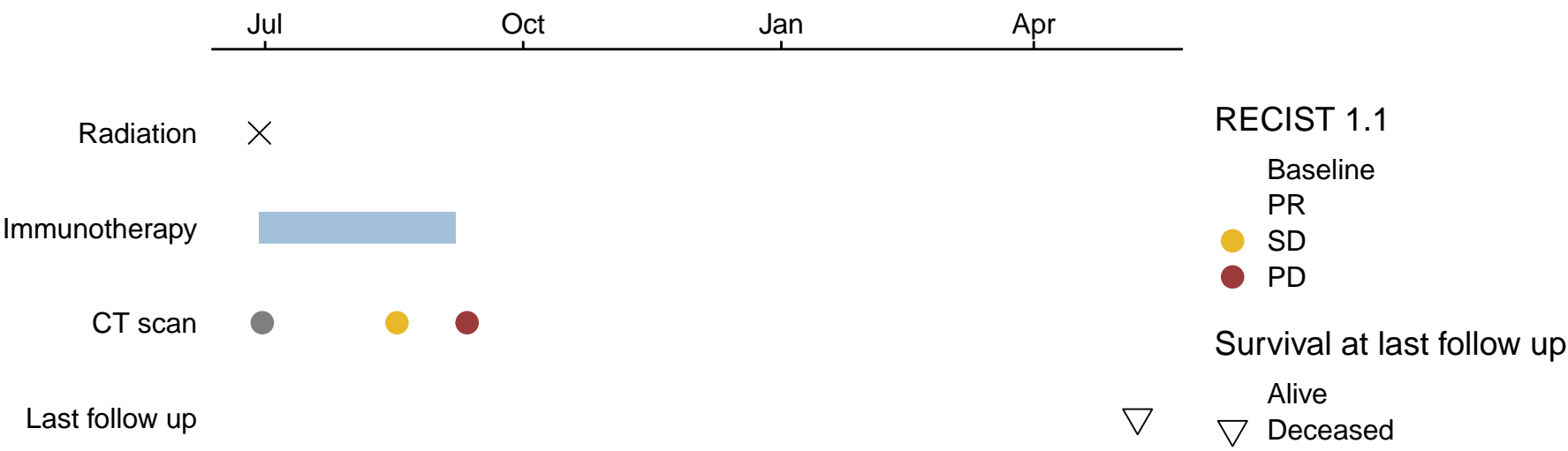
CGPLPA224
WGS data: Tumor (Yes), Normal (Yes), Plasma (25 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 16434, 13403



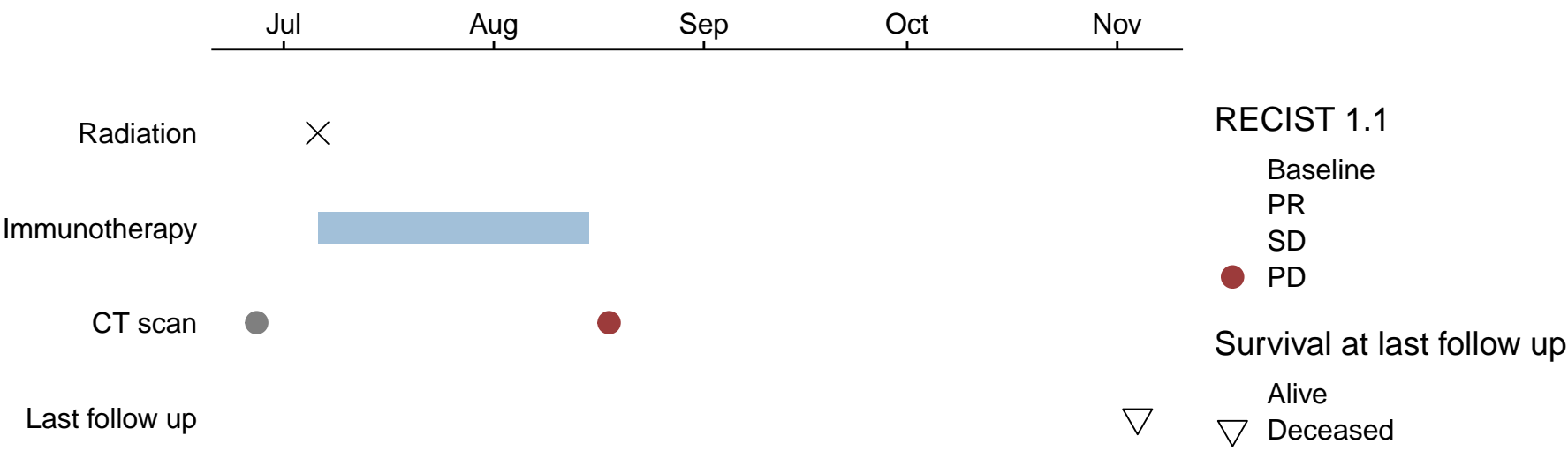
CGPLPA772
WGS data: Tumor (Yes), Normal (Yes), Plasma (4 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 4411, 1763



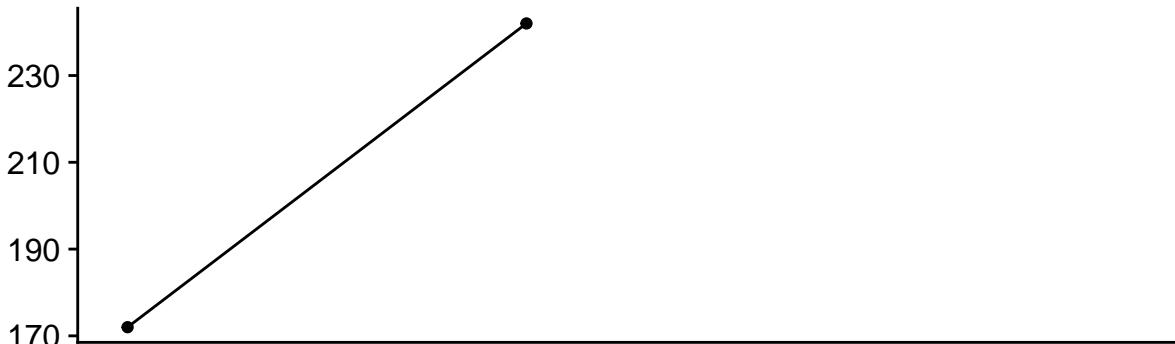
CGPLPA773
WGS data: Tumor (Yes), Normal (Yes), Plasma (4 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 518, 15



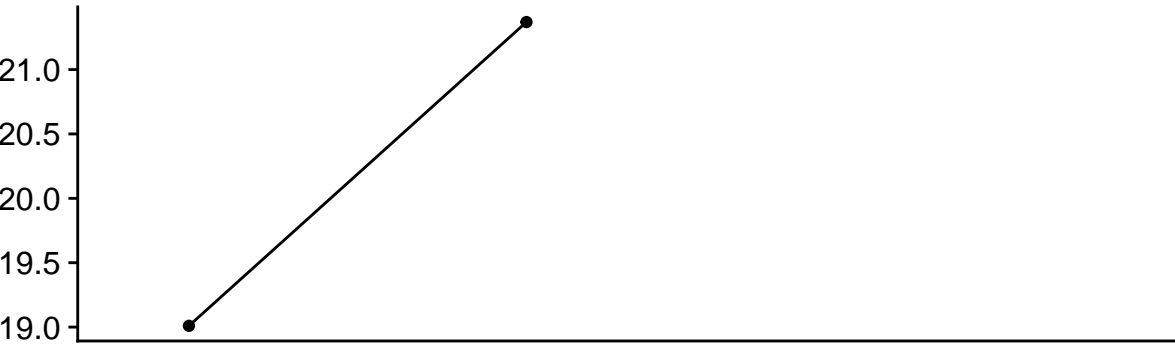
CGPLPA774
WGS data: Tumor (Yes), Normal (Yes), Plasma (2 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 6351, 3446



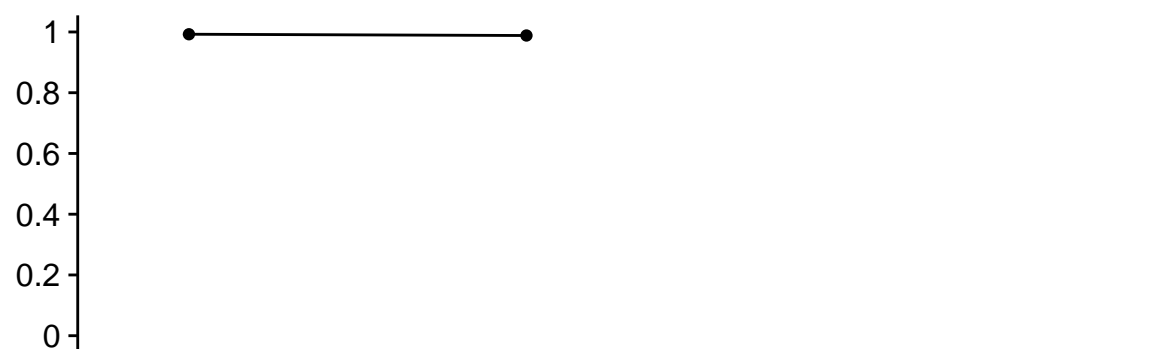
Sum of target lesions (mm)



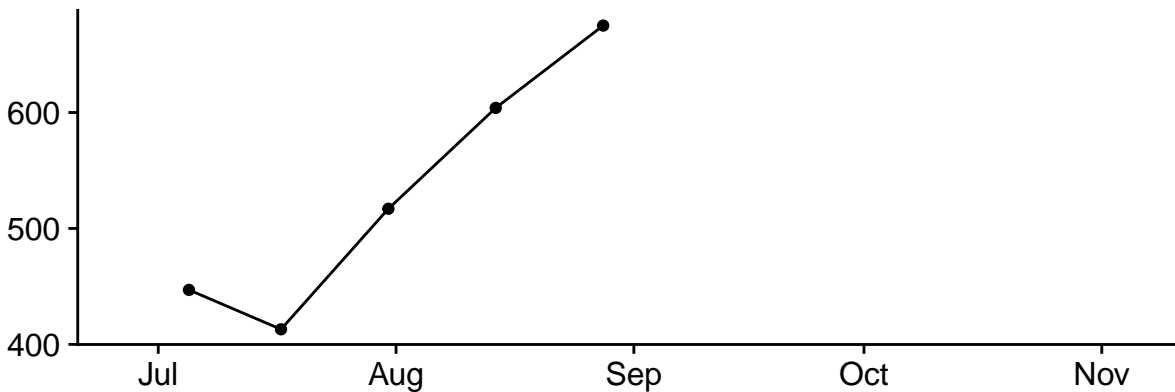
WGMAF (%)

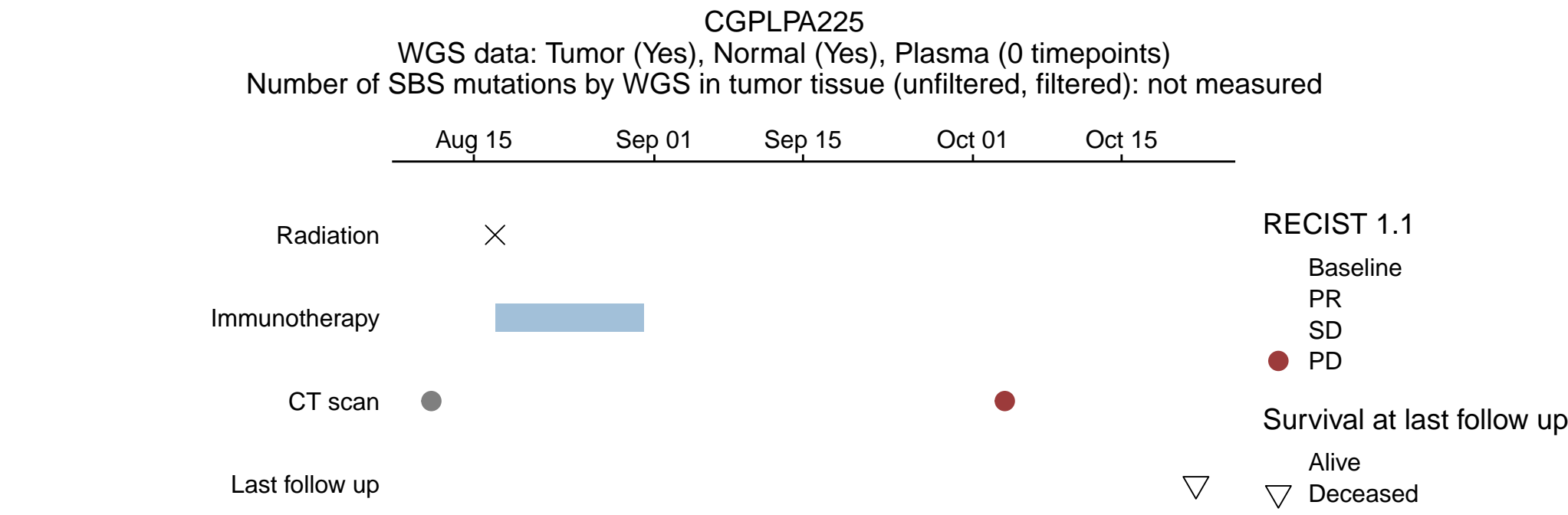


Targeted MAF (%)

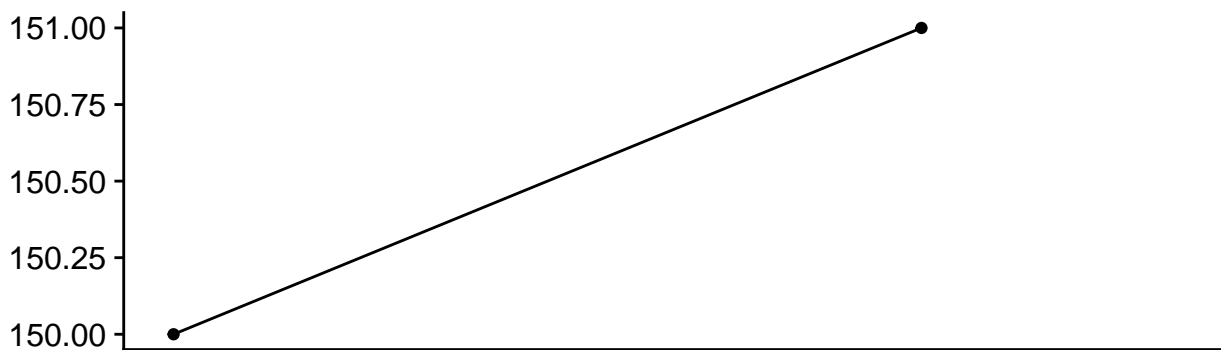


ARTEMIS-DELFI score





Sum of target lesions (mm)

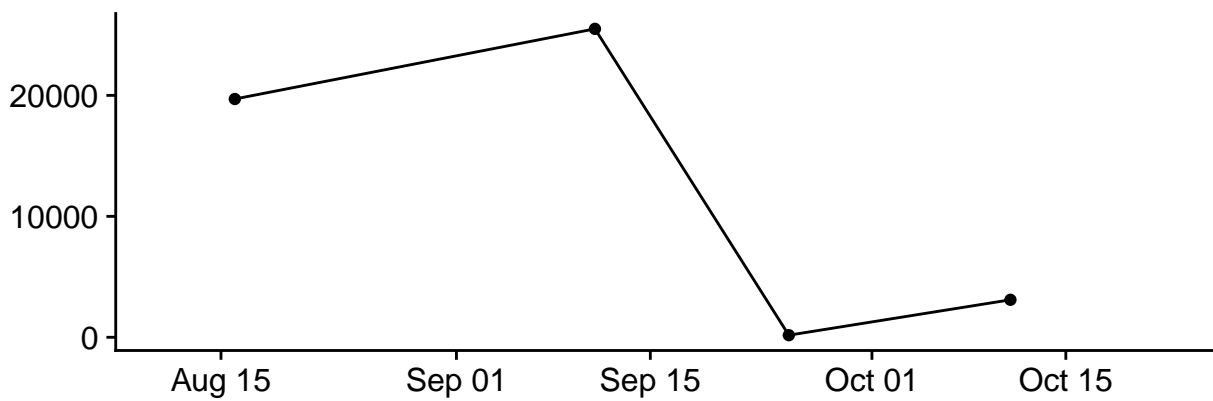


WGMAF (%)

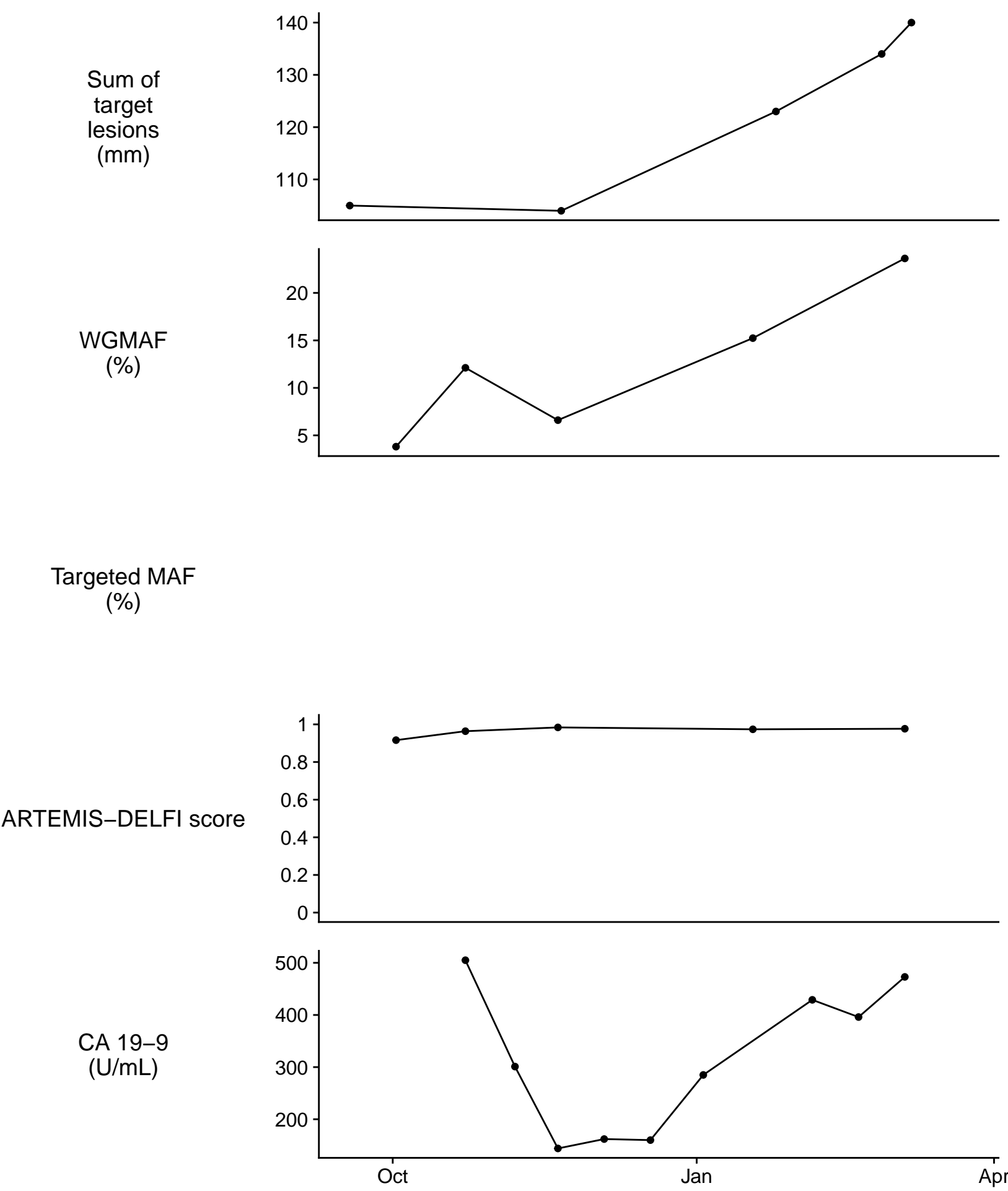
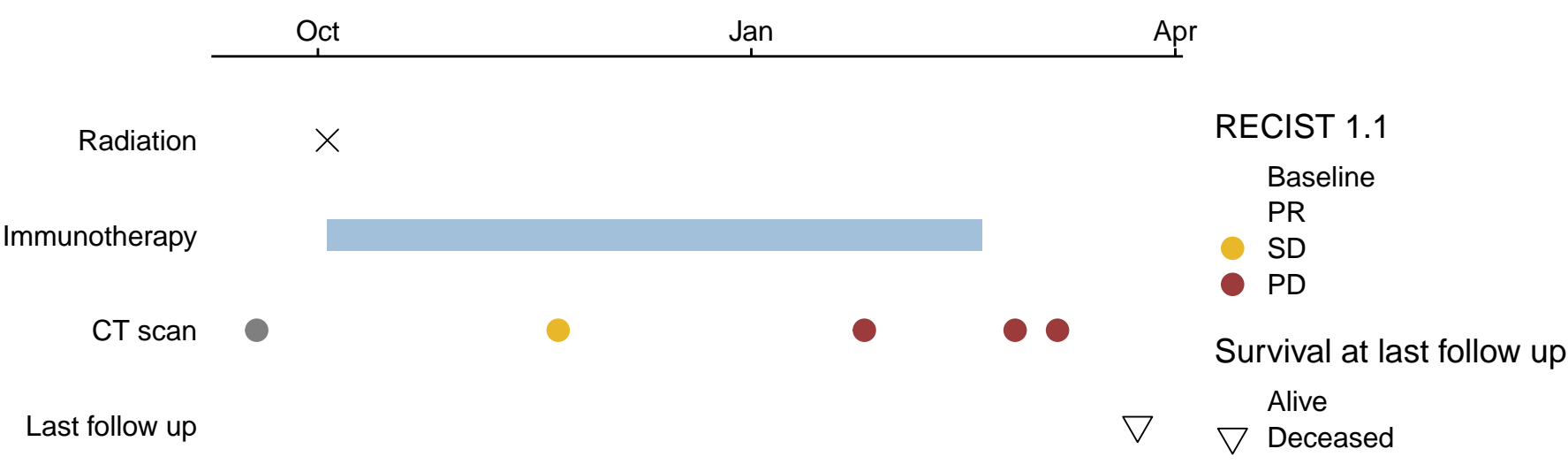
Targeted MAF (%)

ARTEMIS-DELFI score

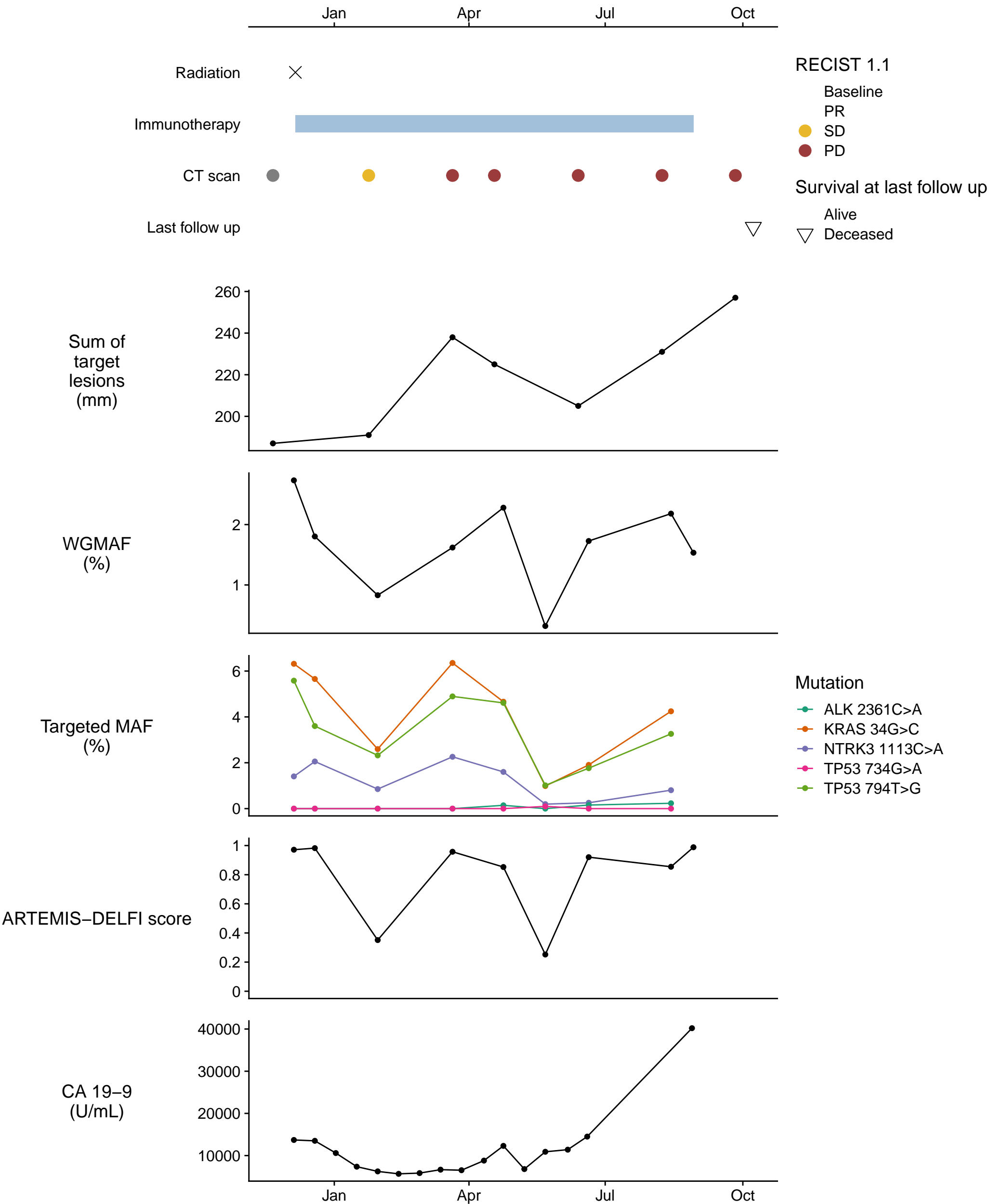
CA 19-9 (U/mL)



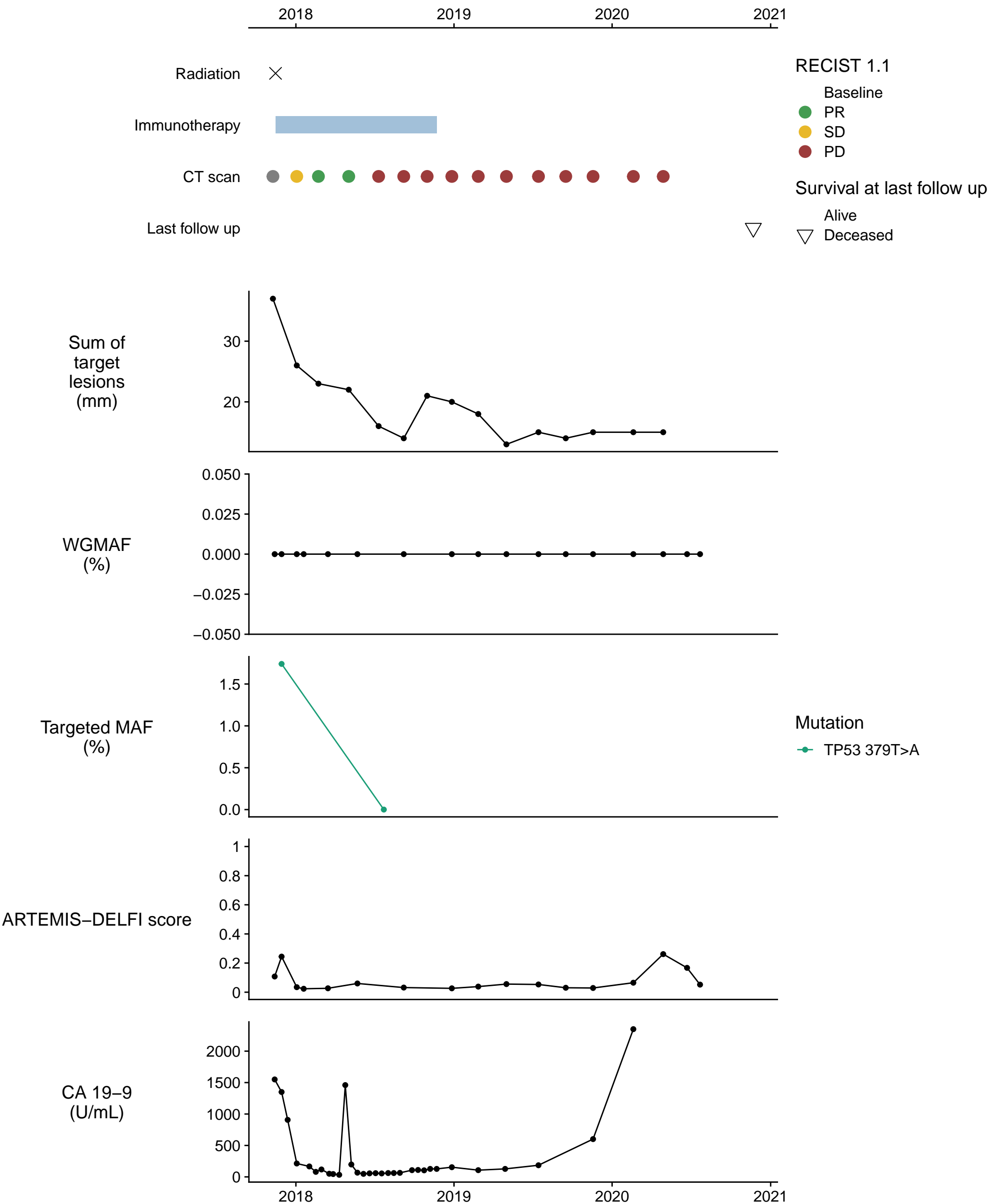
CGPLPA775
WGS data: Tumor (Yes), Normal (Yes), Plasma (5 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 11376, 5757



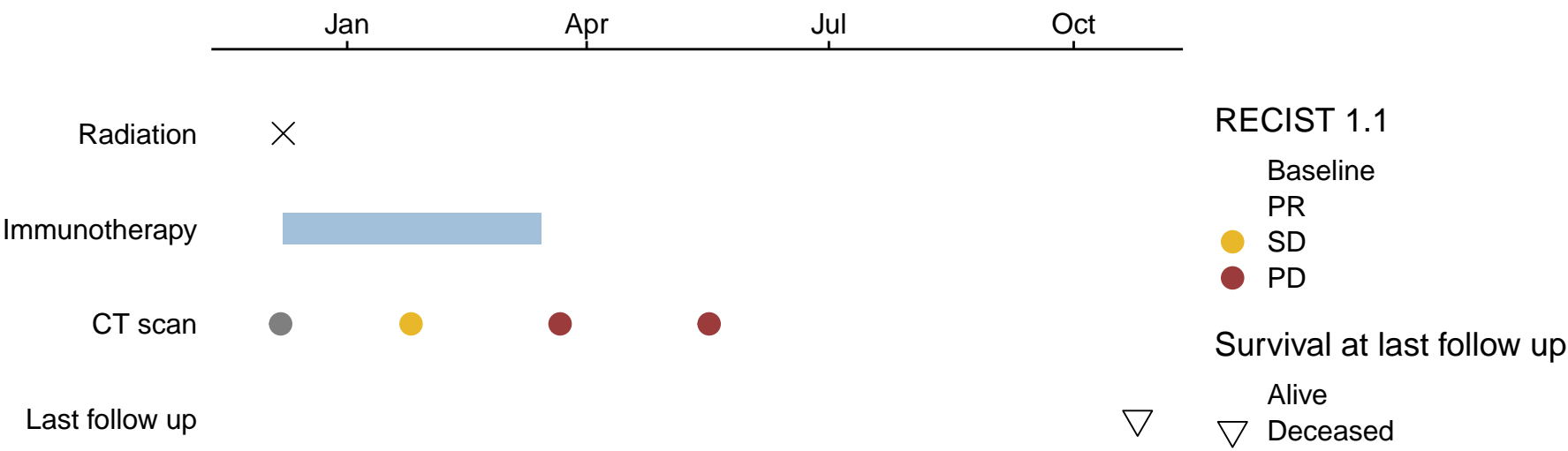
CGPLPA217
WGS data: Tumor (Yes), Normal (Yes), Plasma (9 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 15589, 13432



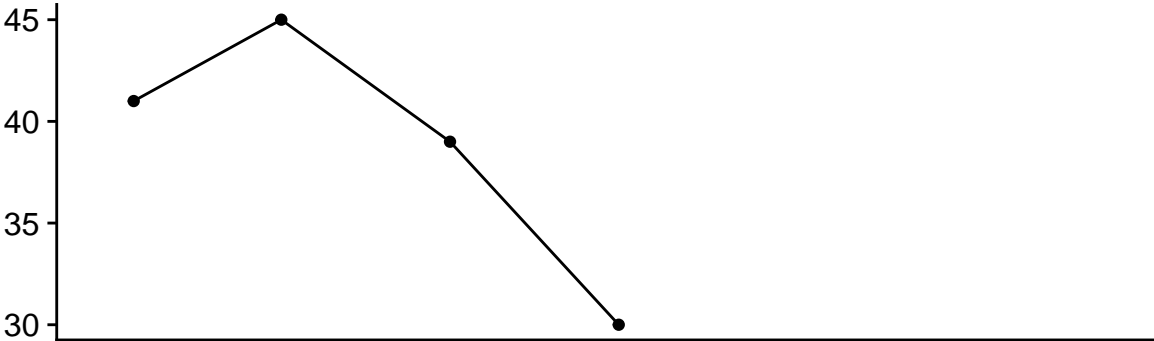
CGPLPA222
WGS data: Tumor (Yes), Normal (Yes), Plasma (17 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 99, 9



CGPLPA776
WGS data: Tumor (No), Normal (Yes), Plasma (5 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured



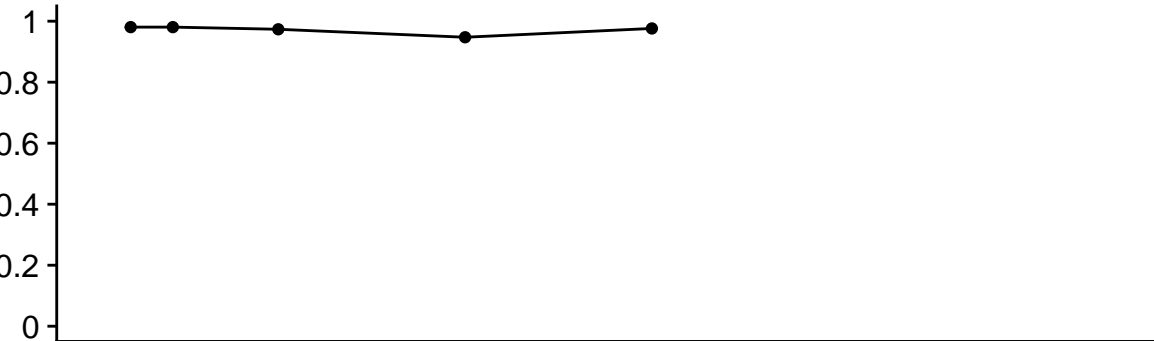
Sum of target lesions (mm)



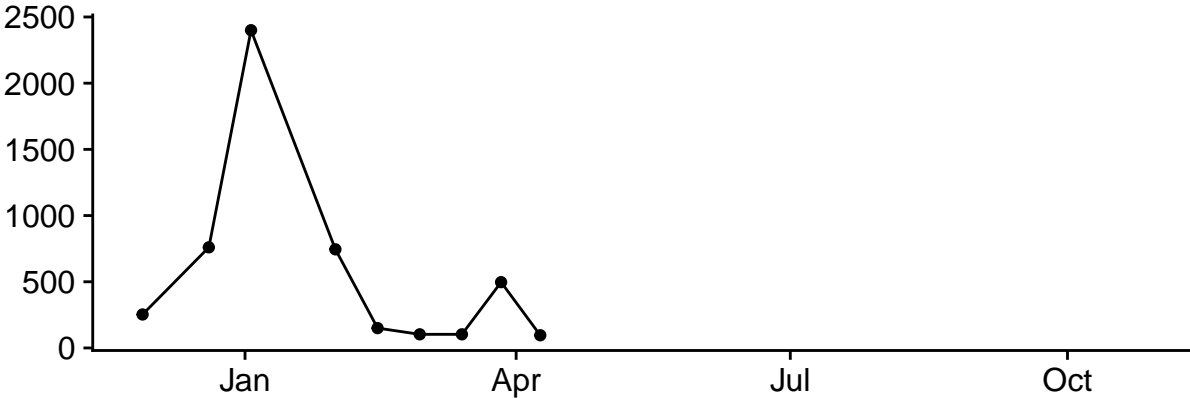
WGMAF (%)

Targeted MAF (%)

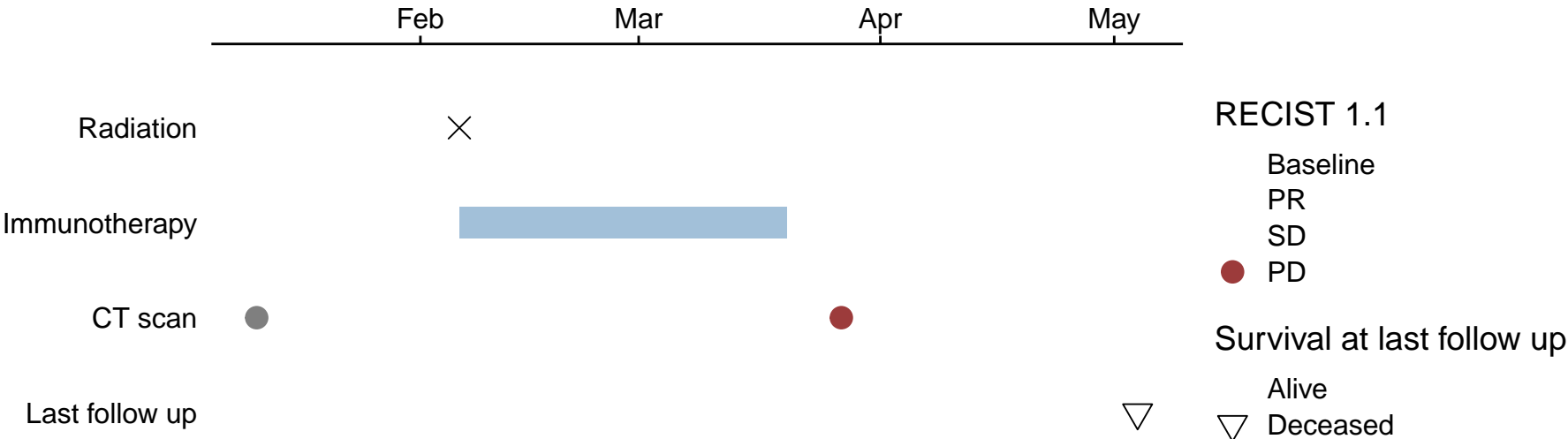
ARTEMIS-DELFI score



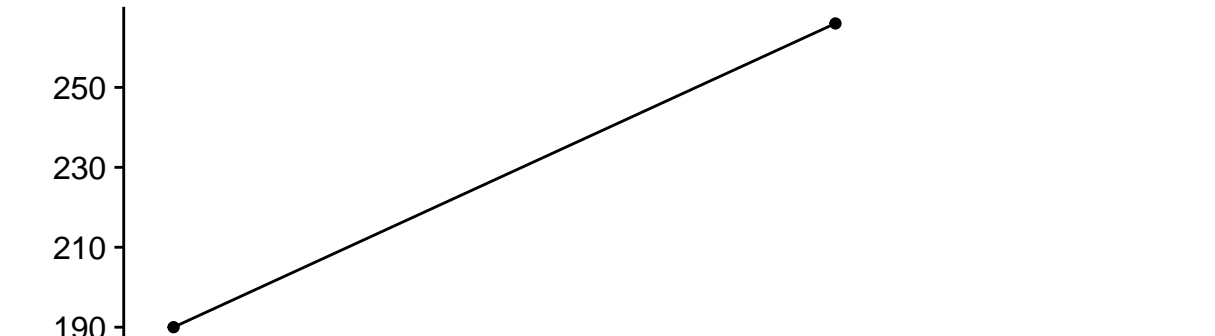
CA 19-9 (U/mL)



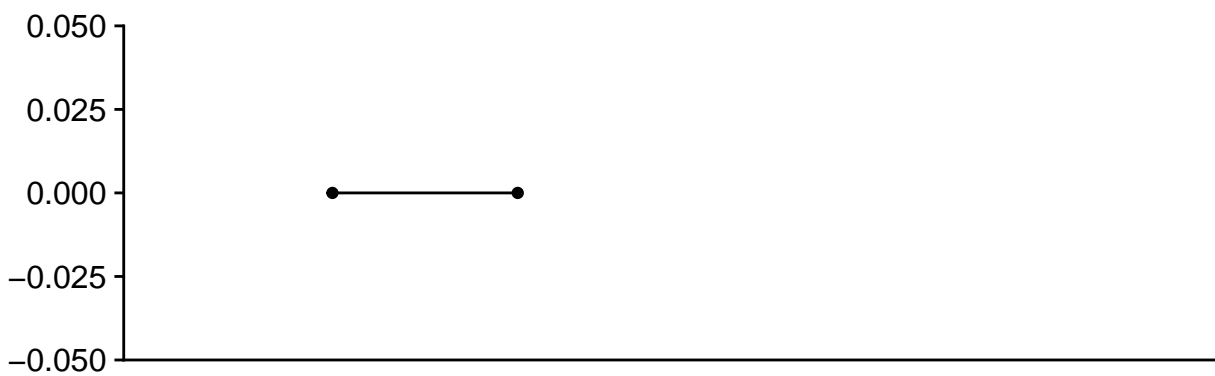
CGPLPA777
WGS data: Tumor (Yes), Normal (Yes), Plasma (2 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 726, 42



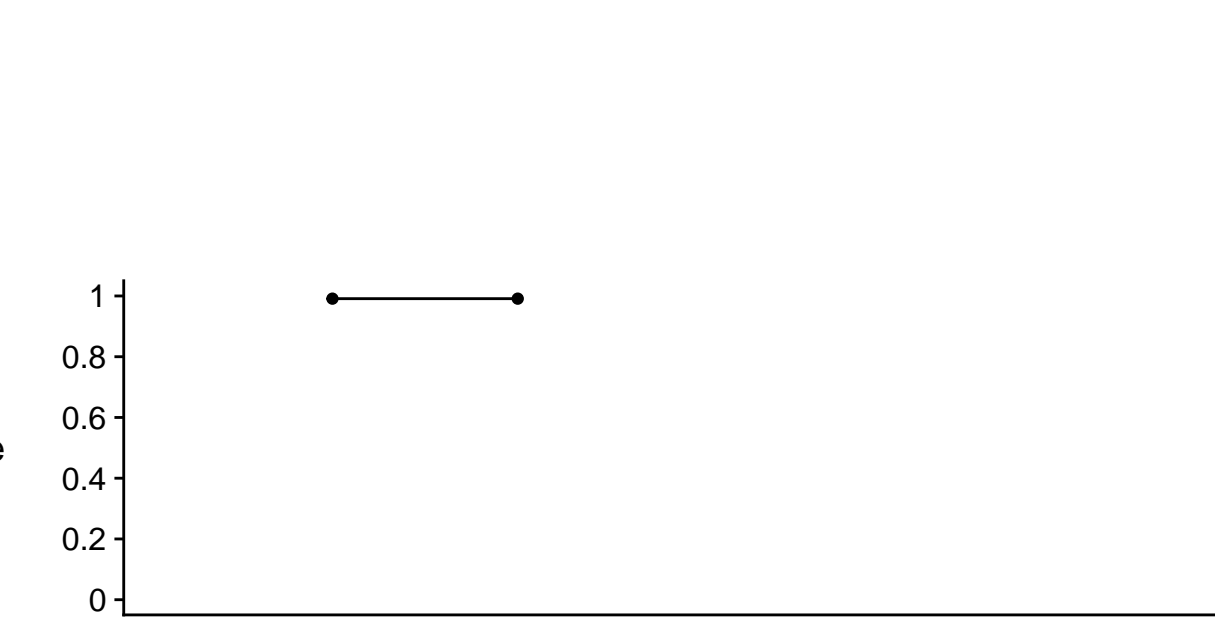
Sum of target lesions (mm)



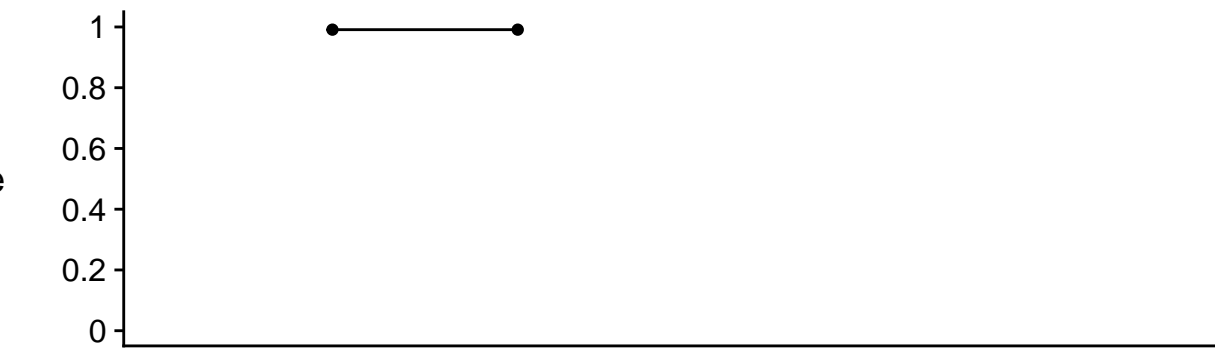
WGMAF (%)



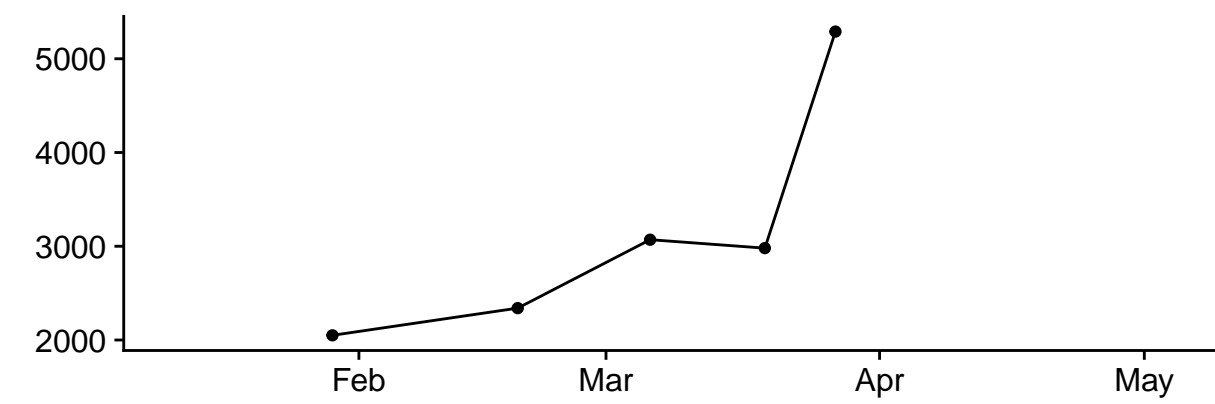
Targeted MAF (%)



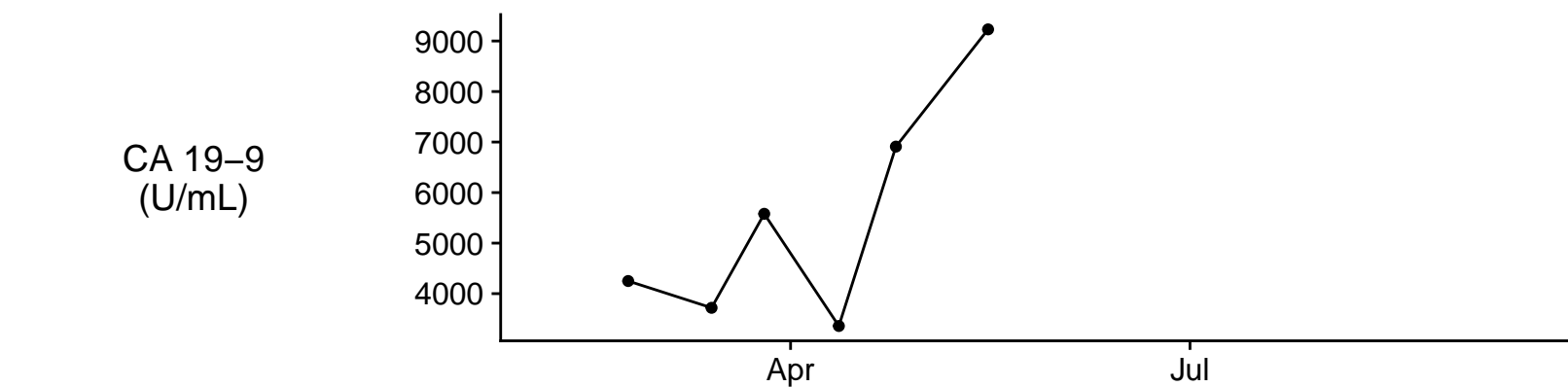
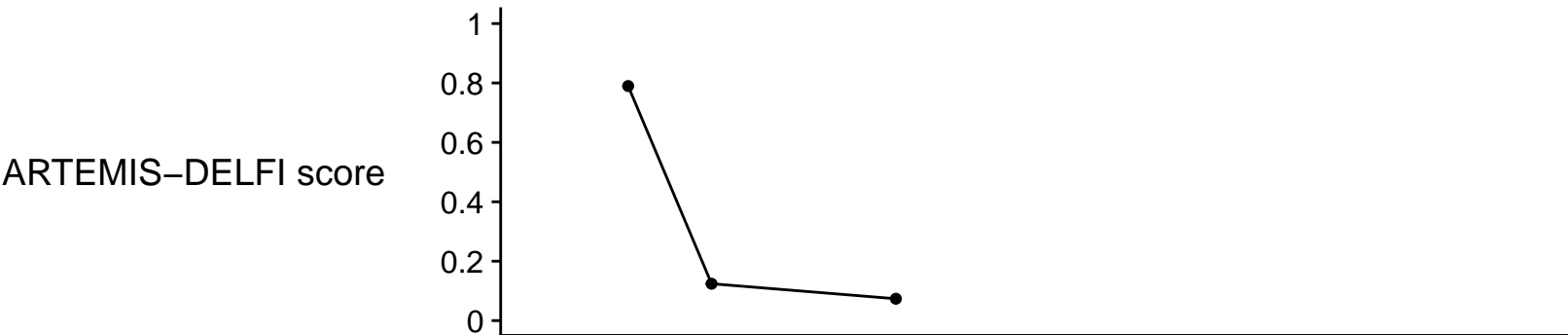
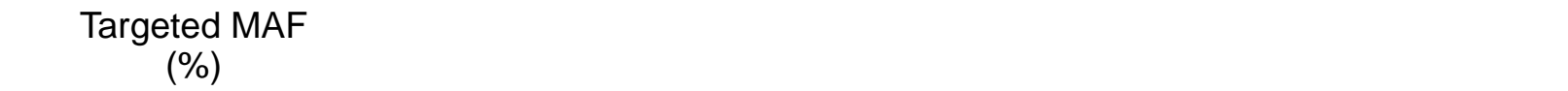
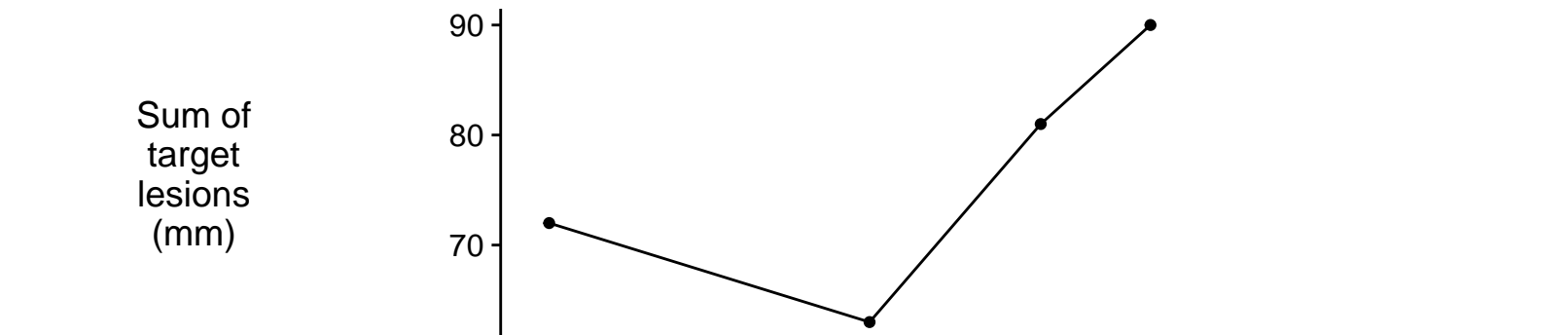
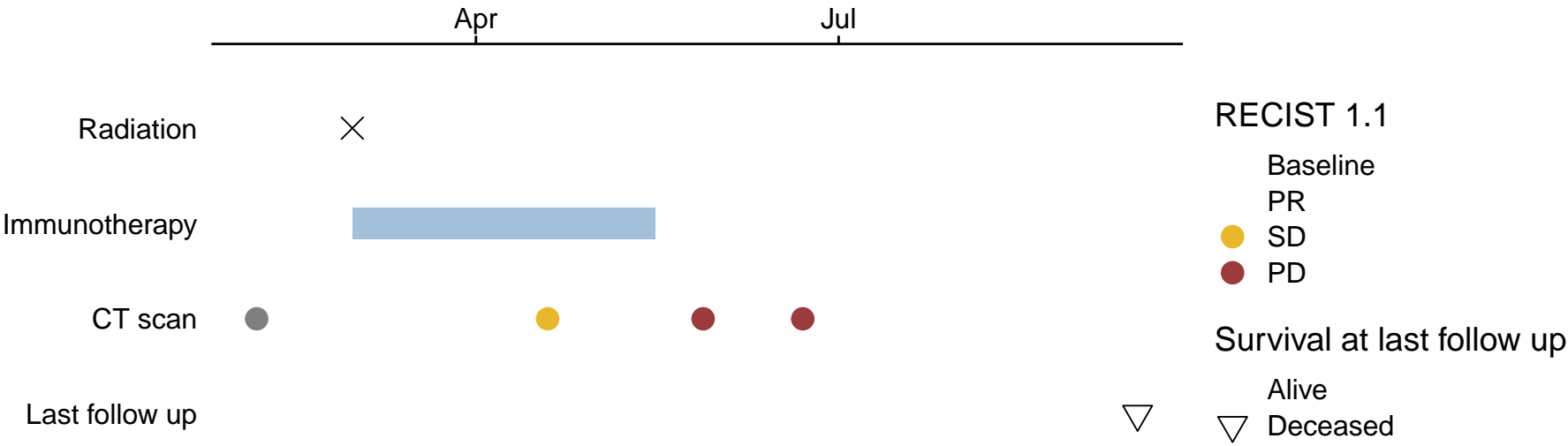
ARTEMIS-DELFI score



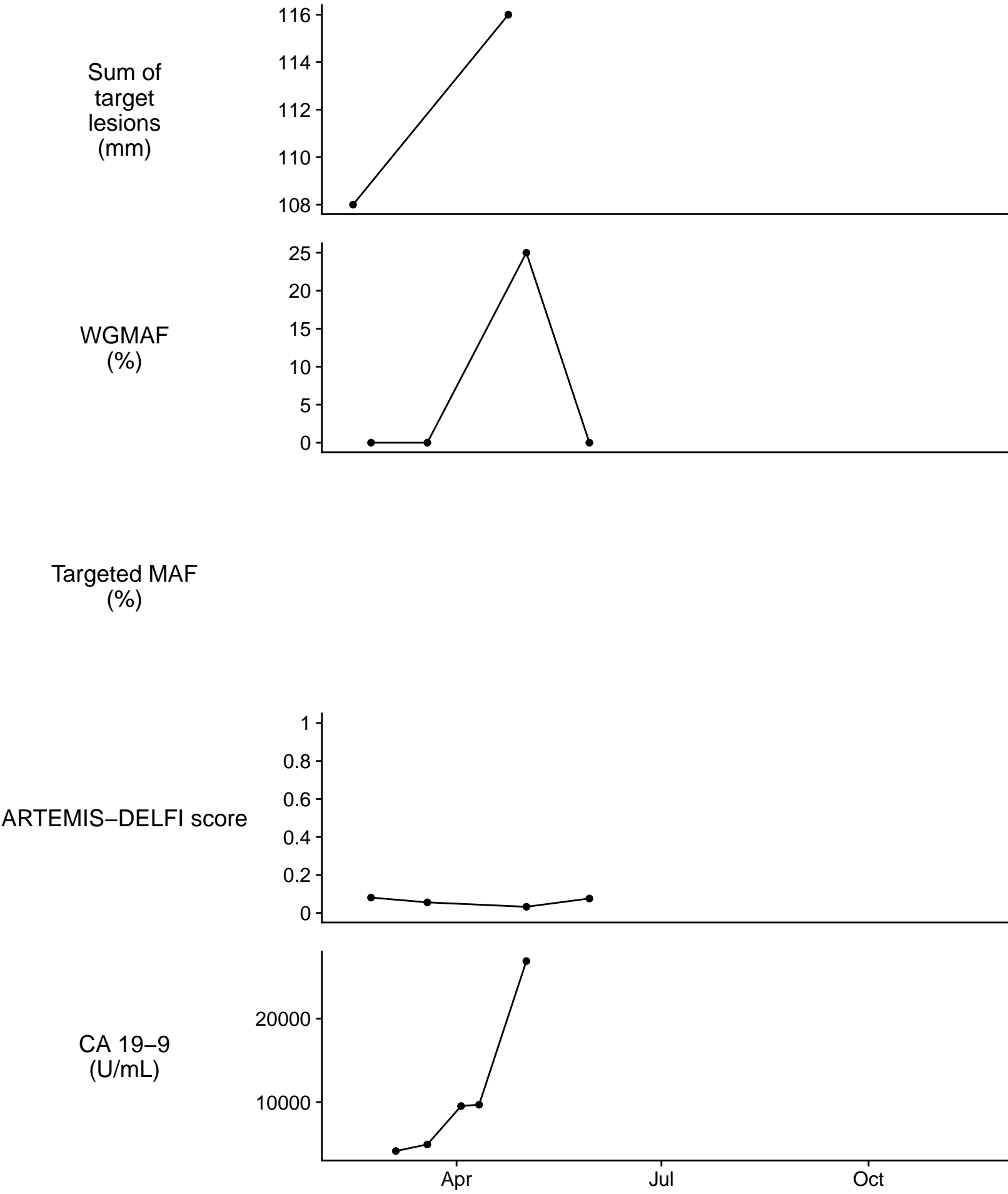
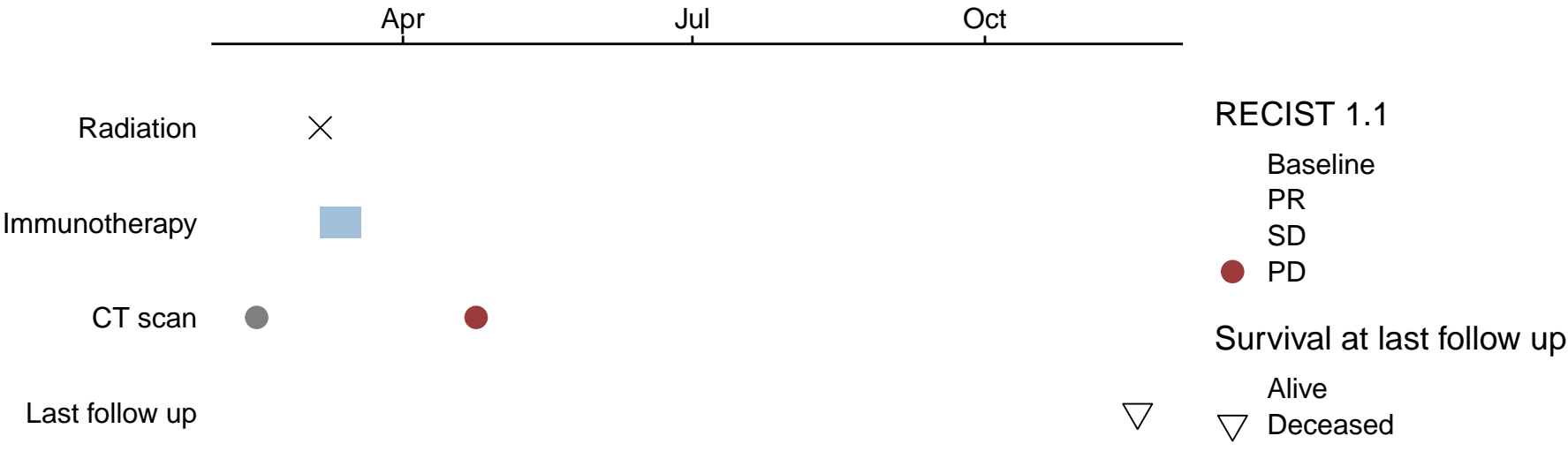
CA 19-9 (U/mL)



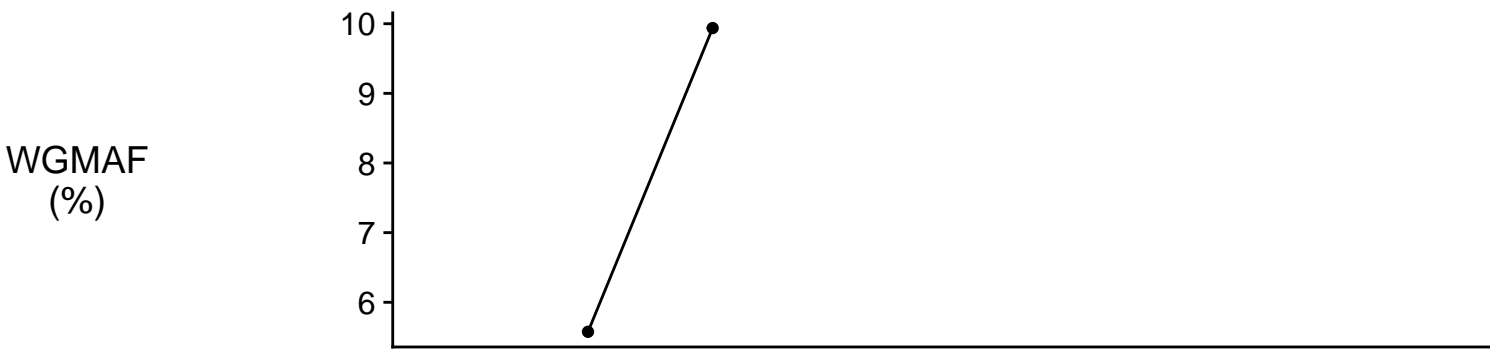
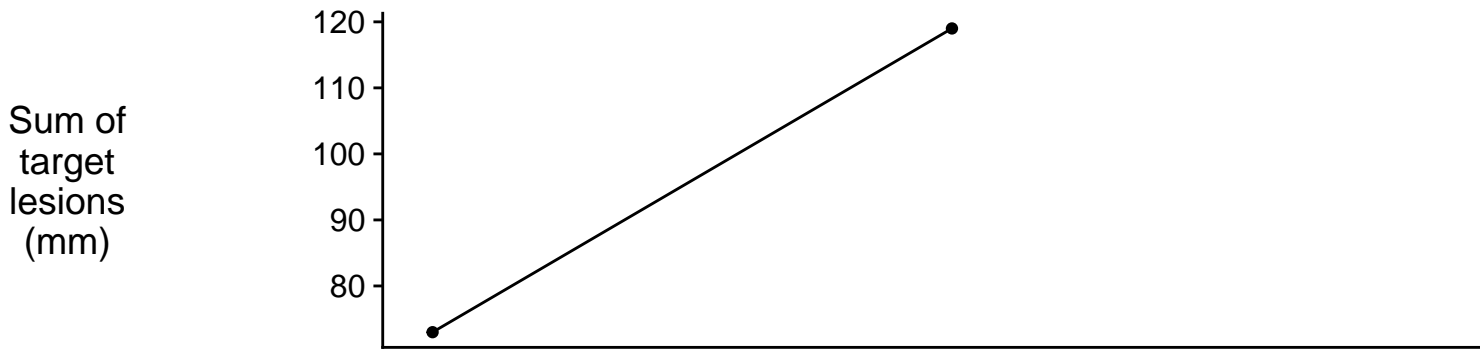
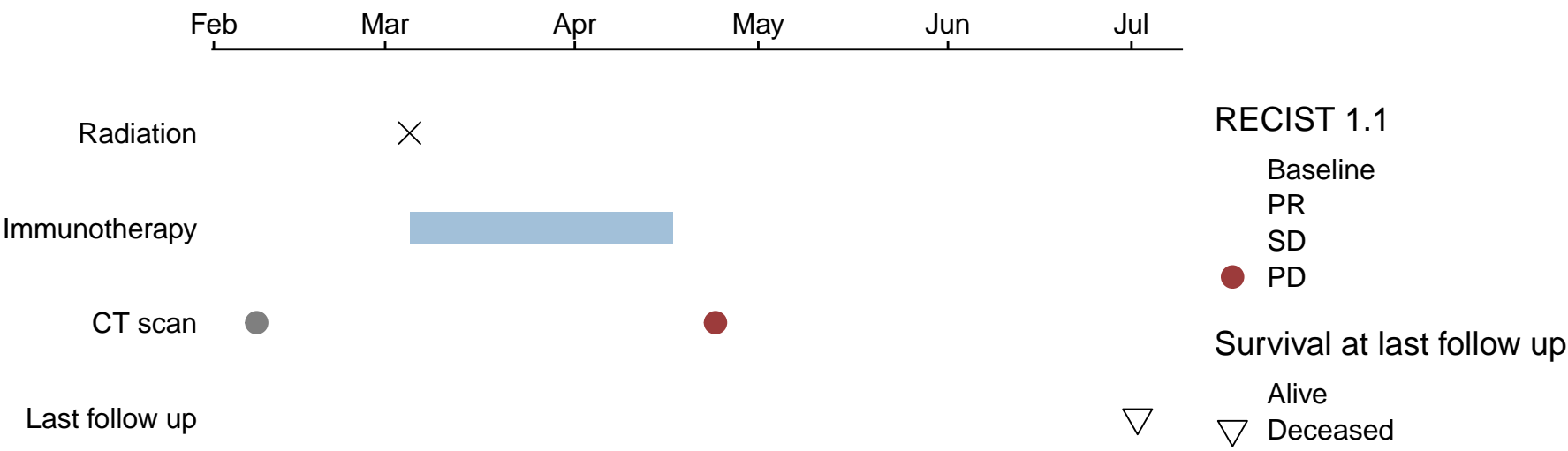
CGPLPA778
WGS data: Tumor (No), Normal (No), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured



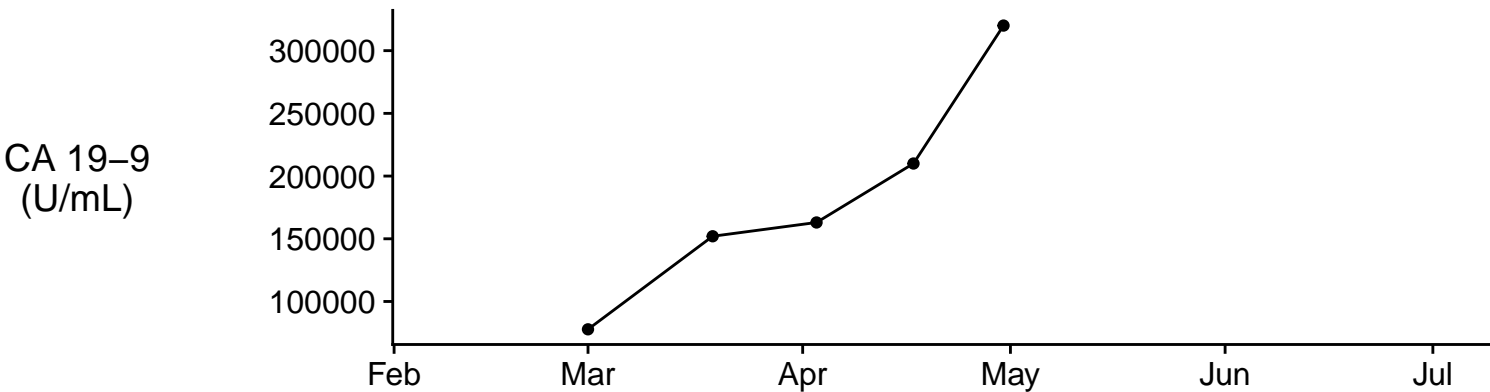
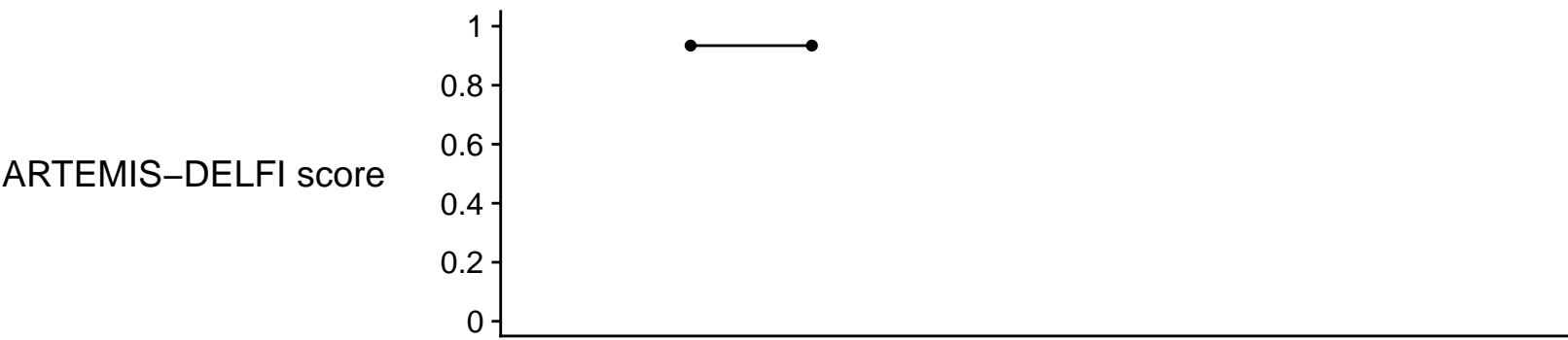
CGPLPA779
WGS data: Tumor (Yes), Normal (Yes), Plasma (4 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 581, 11



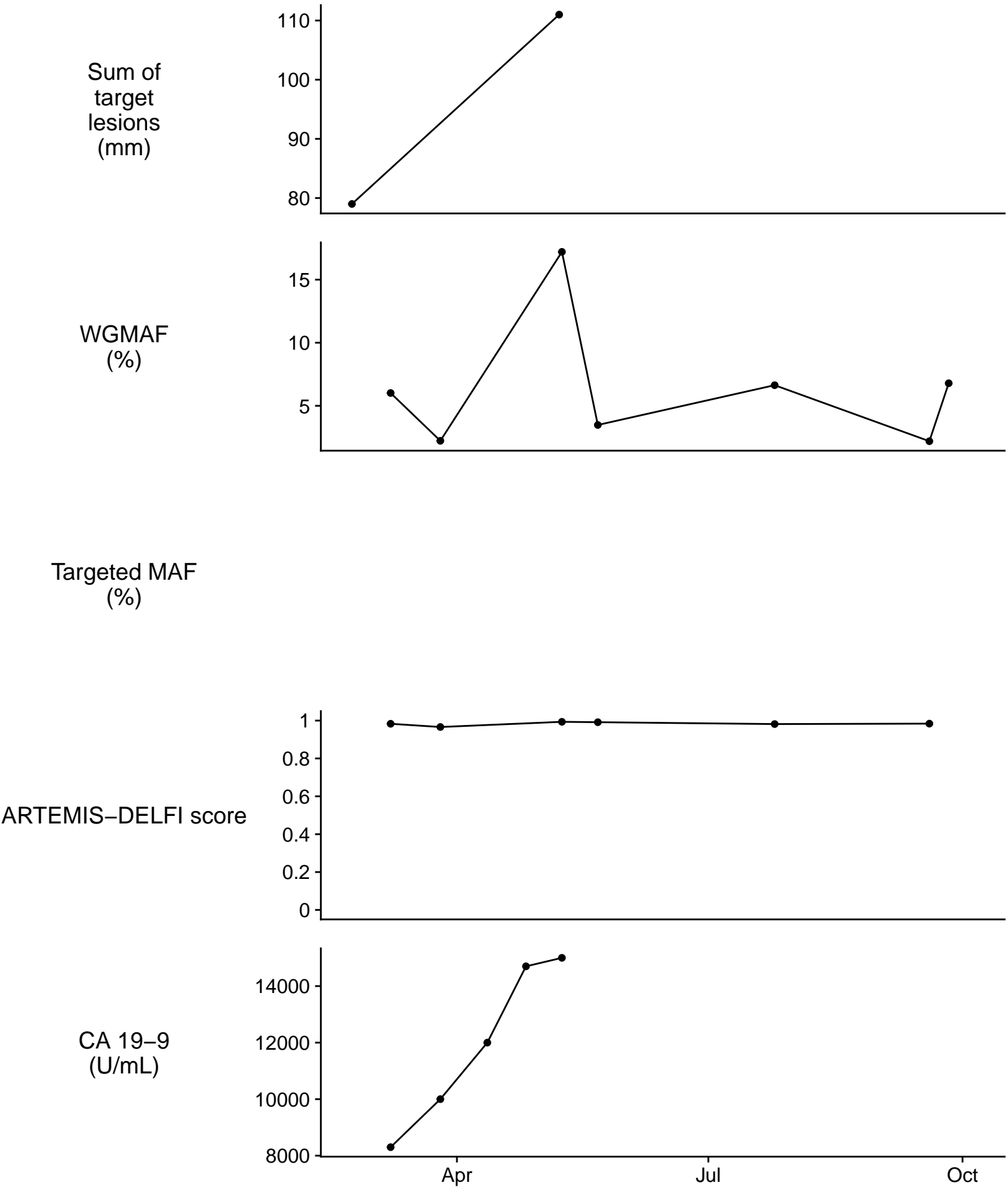
CGPLPA780
WGS data: Tumor (Yes), Normal (Yes), Plasma (2 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 6996, 1450



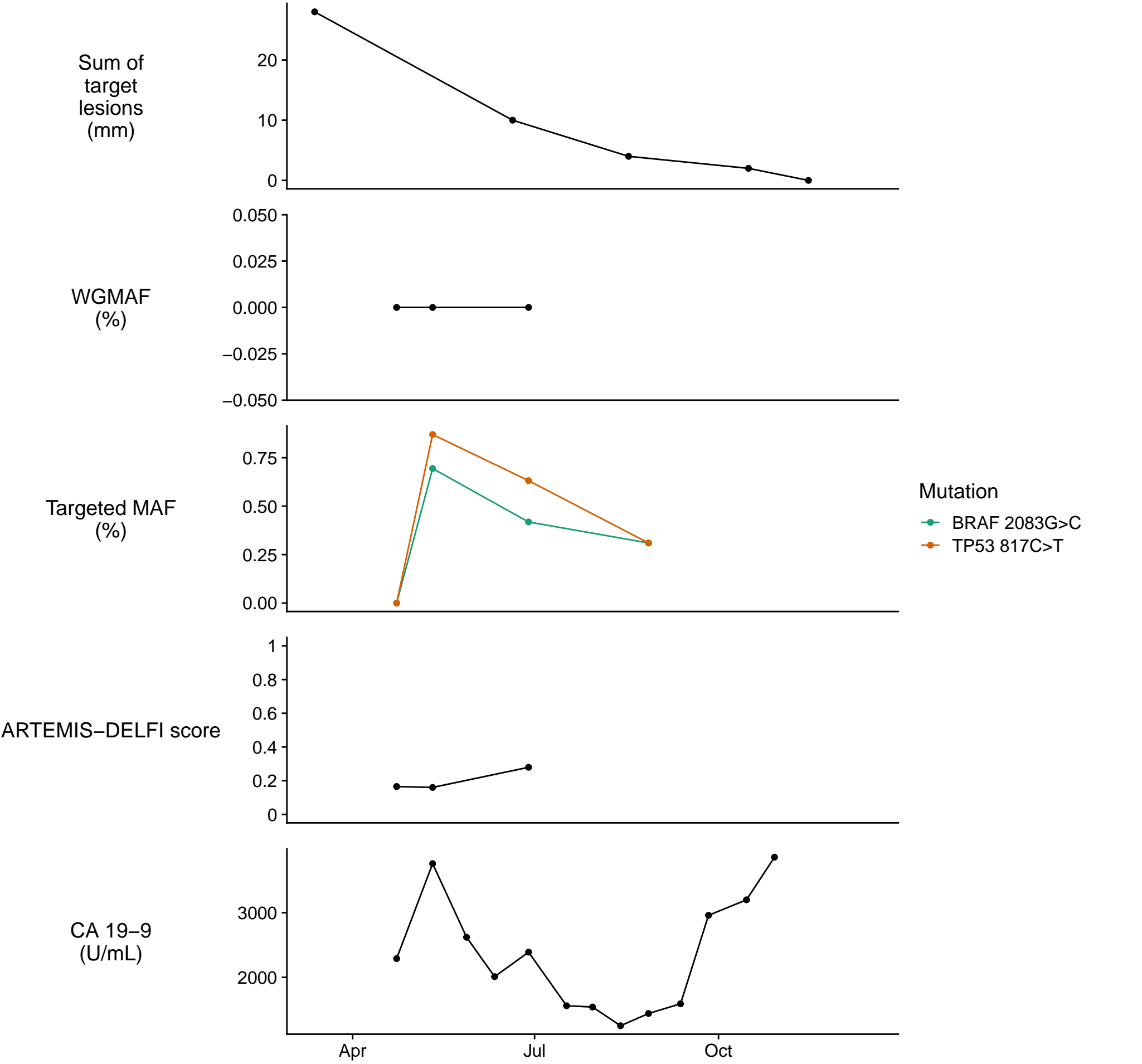
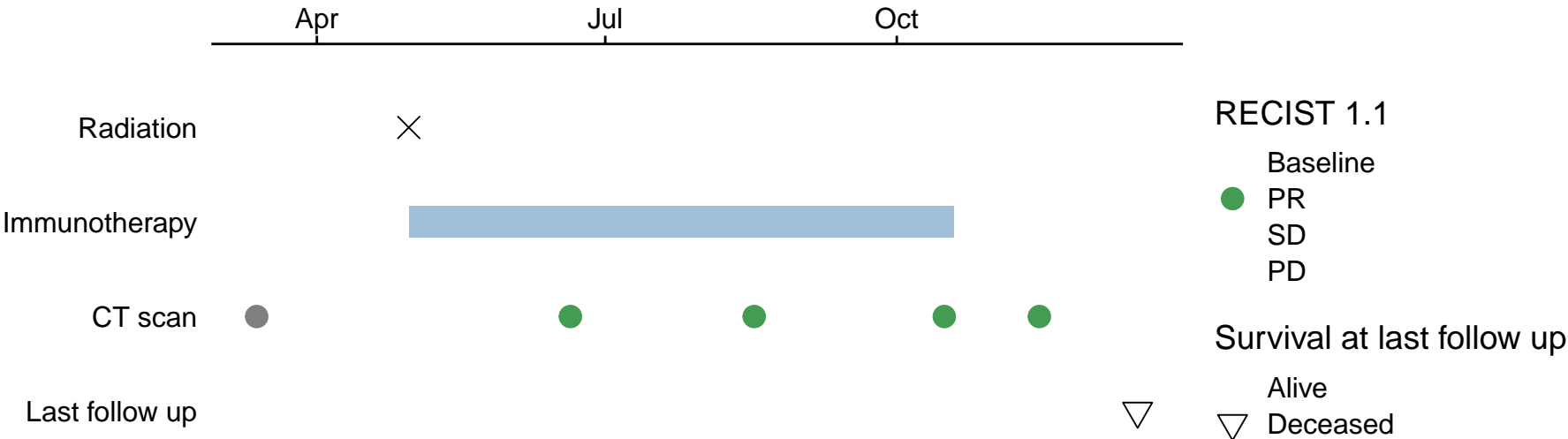
Targeted MAF (%)



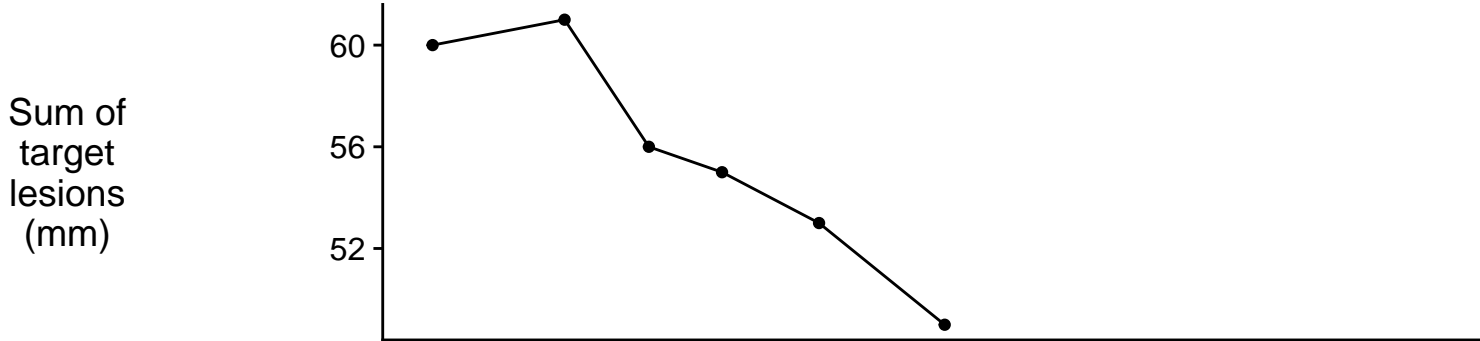
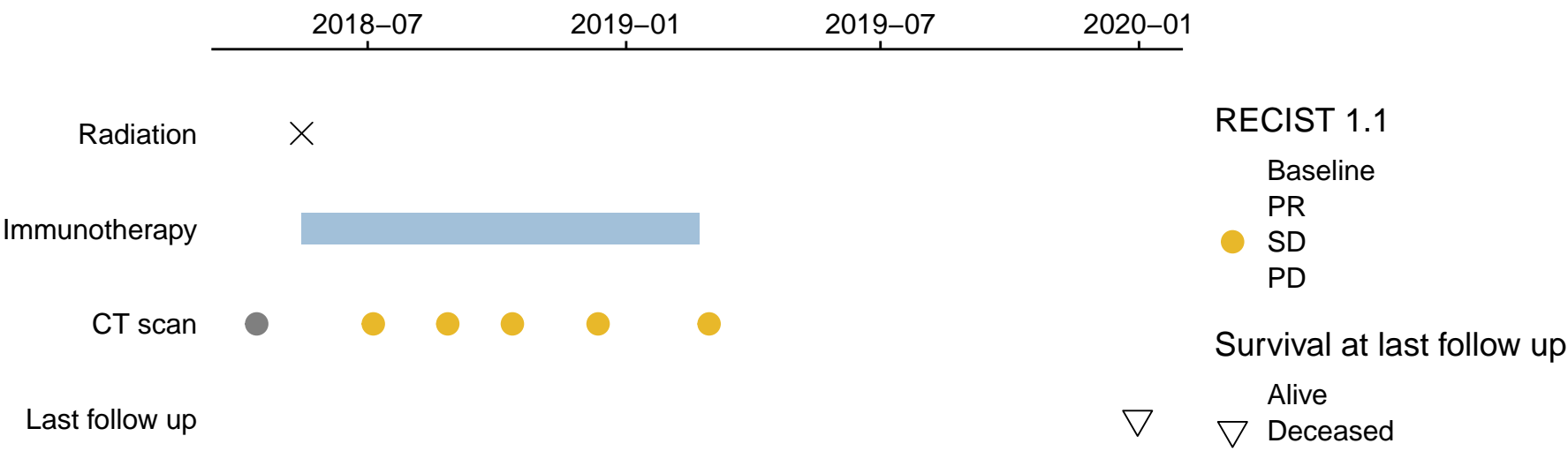
CGPLPA781
WGS data: Tumor (Yes), Normal (Yes), Plasma (7 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 4501, 1416



CGPLPA226
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 55, 11

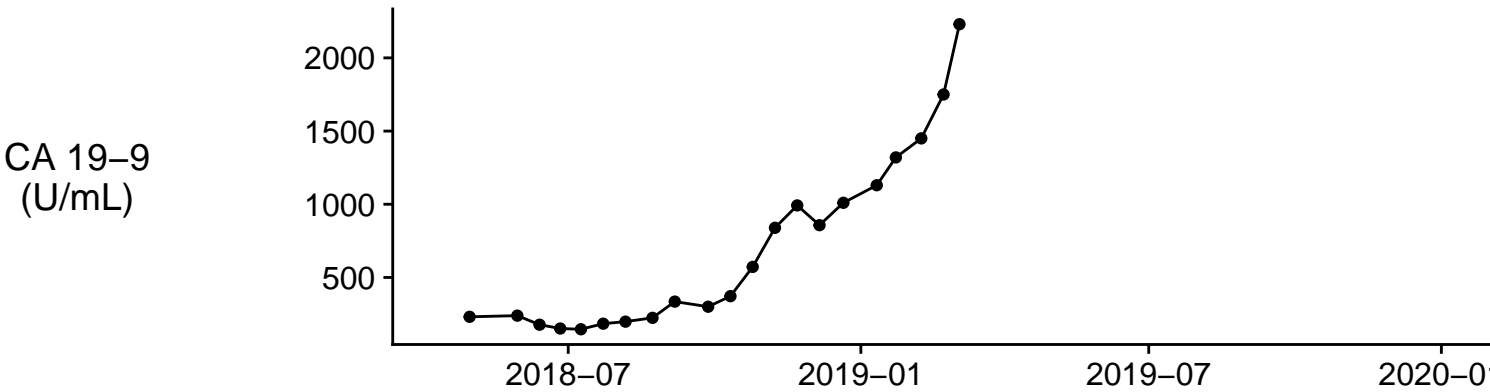
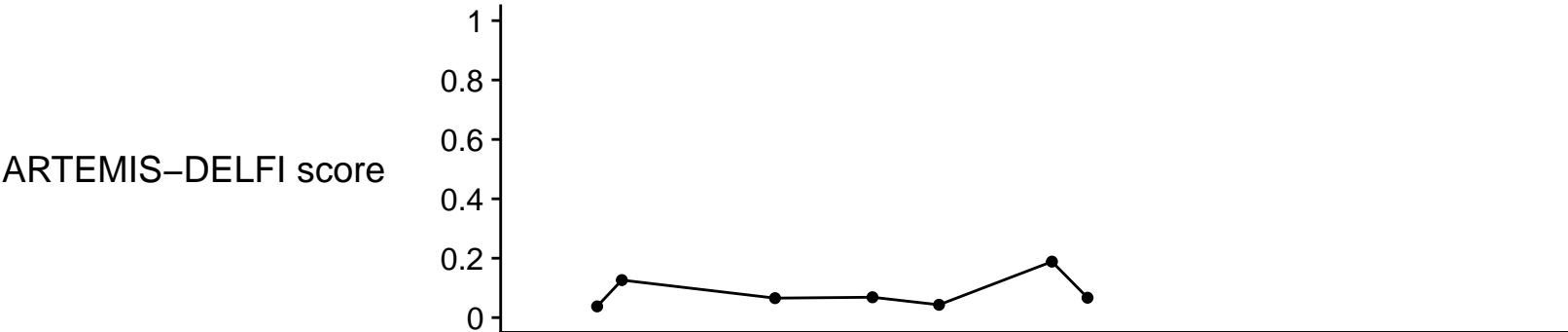


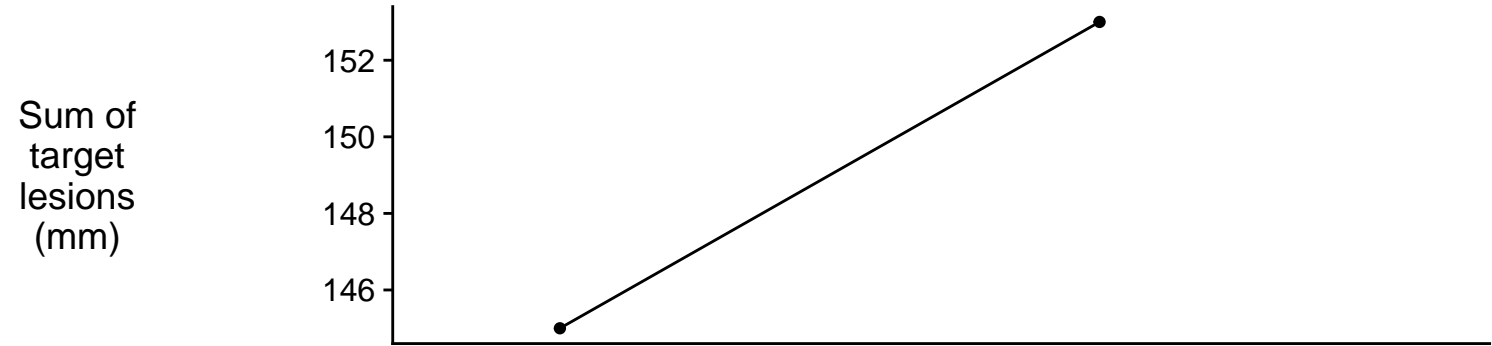
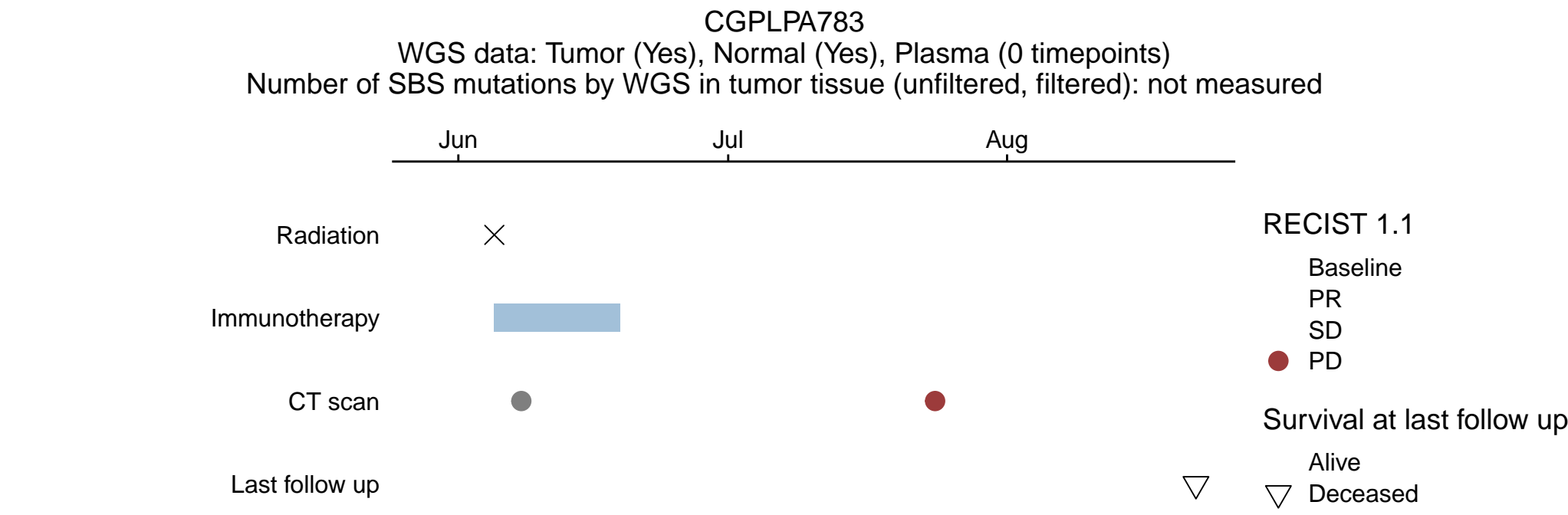
CGPLPA782
WGS data: Tumor (No), Normal (No), Plasma (7 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured



WGMAF (%)

Targeted MAF (%)

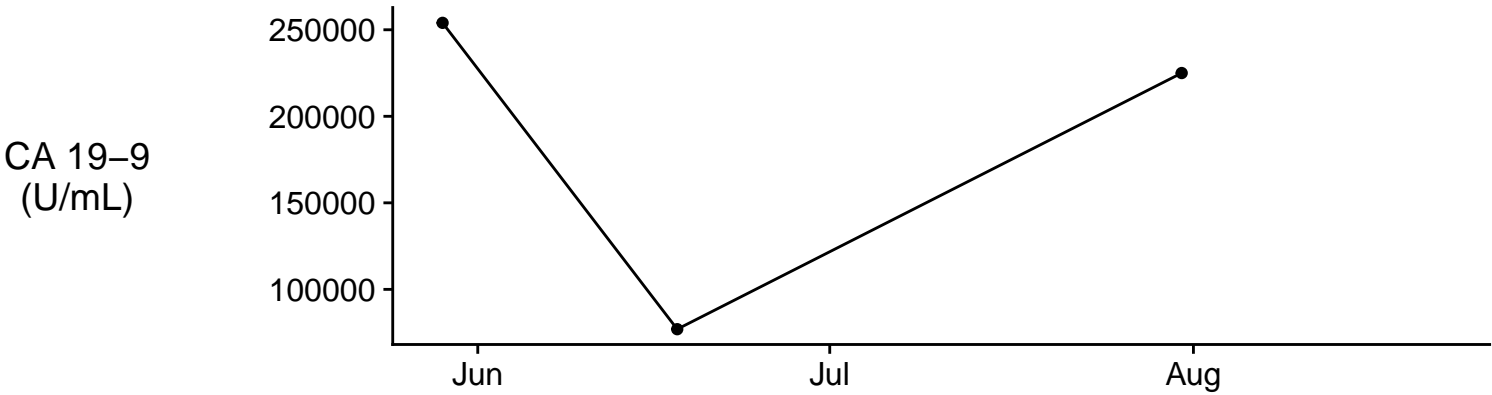




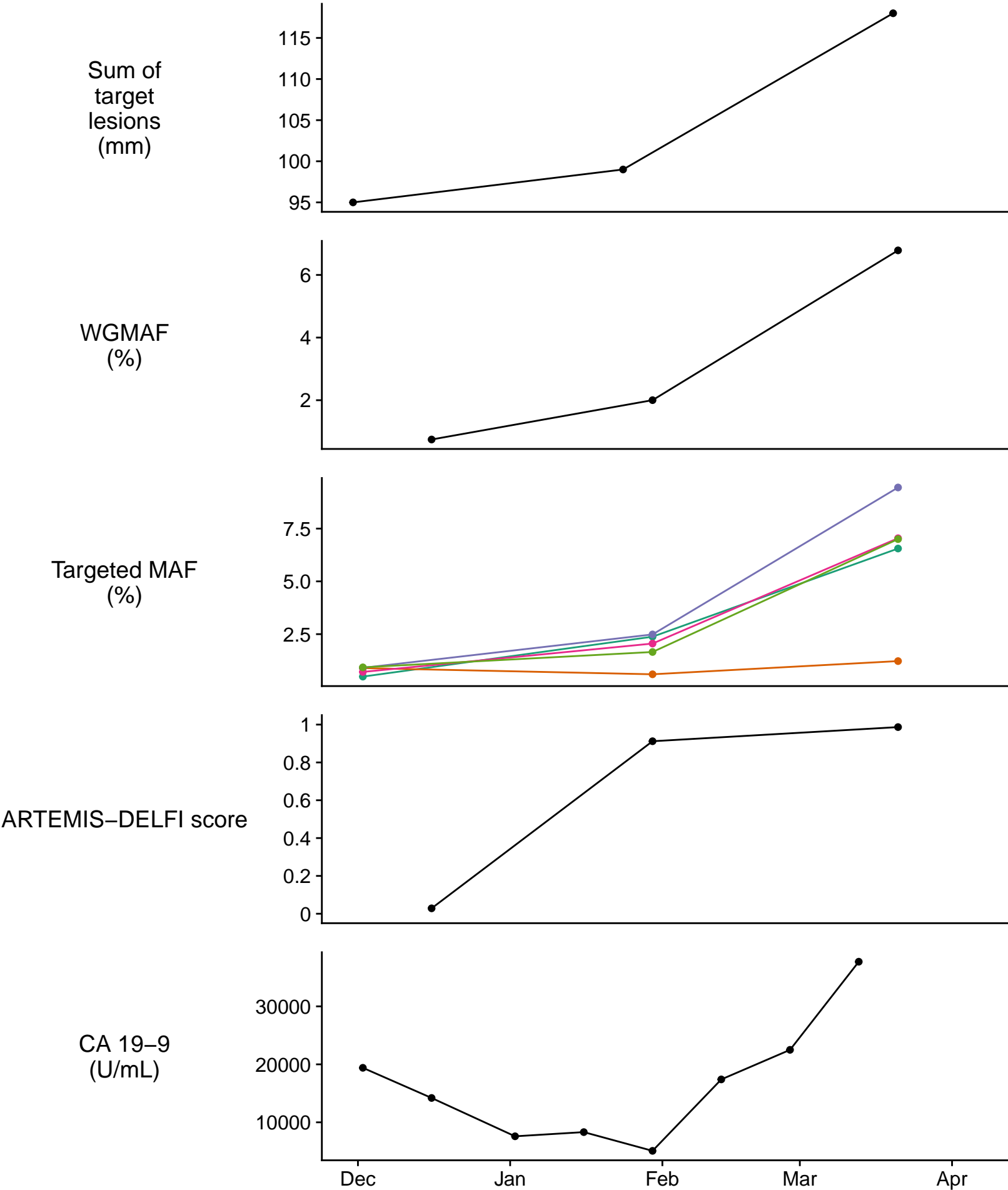
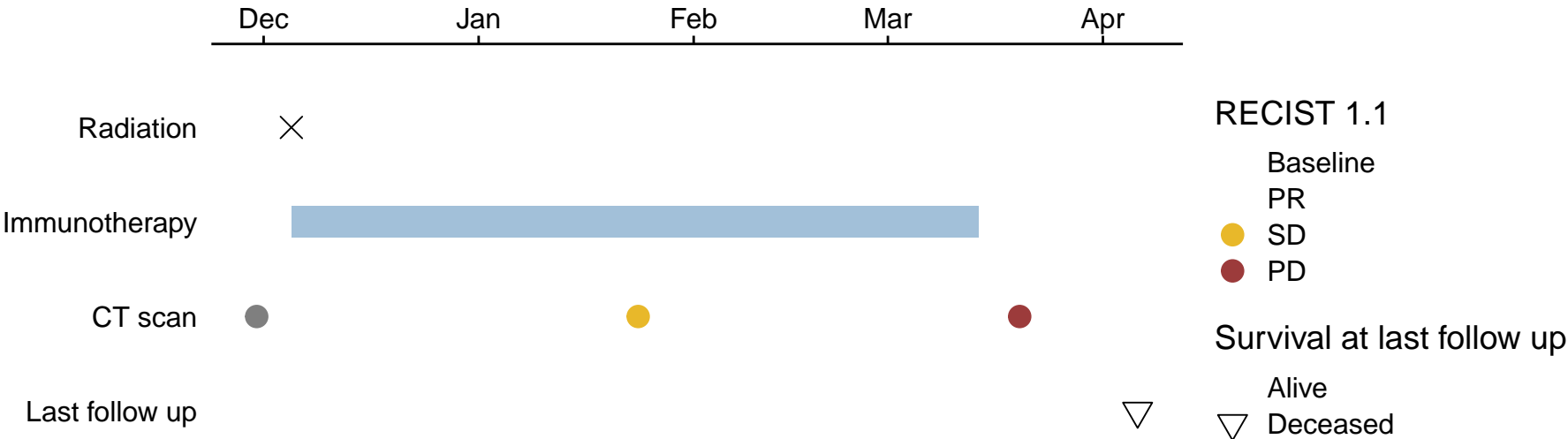
WGMAF (%)

Targeted MAF (%)

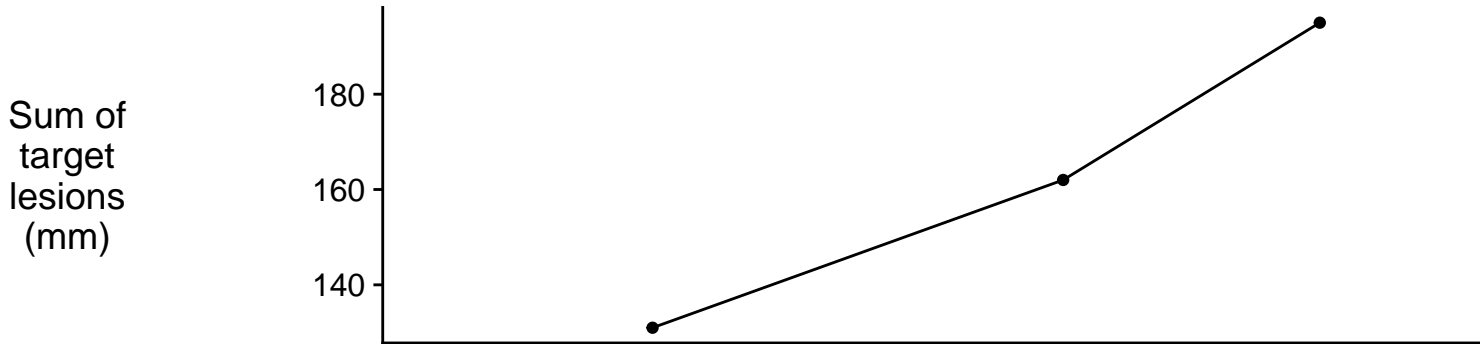
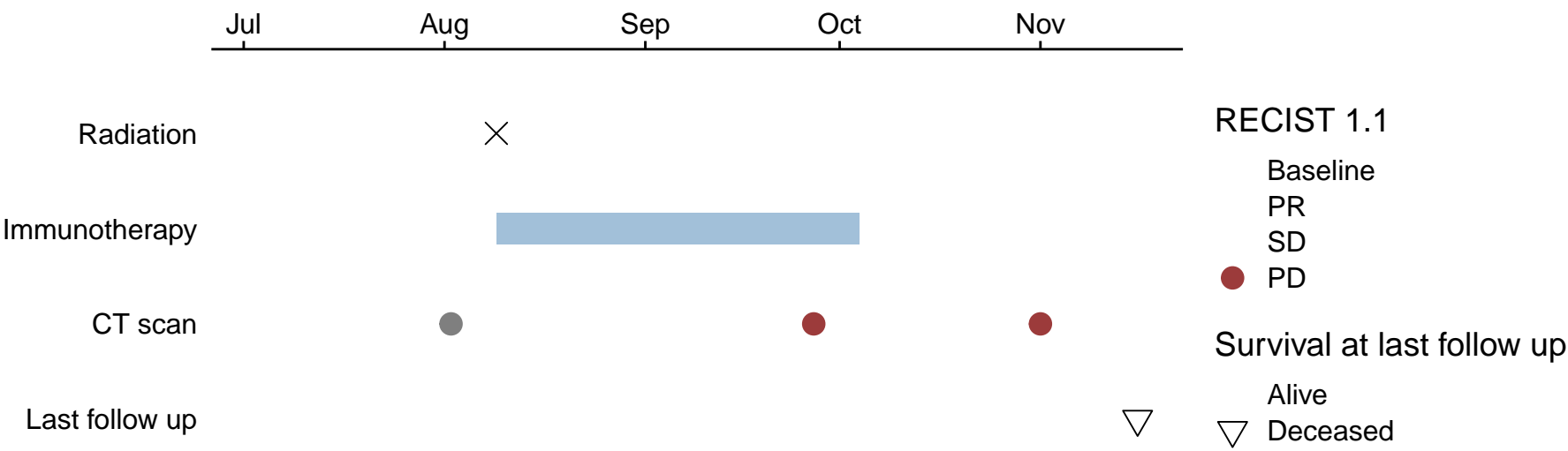
ARTEMIS-DELFI score



CGPLPA220
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 4171, 3288

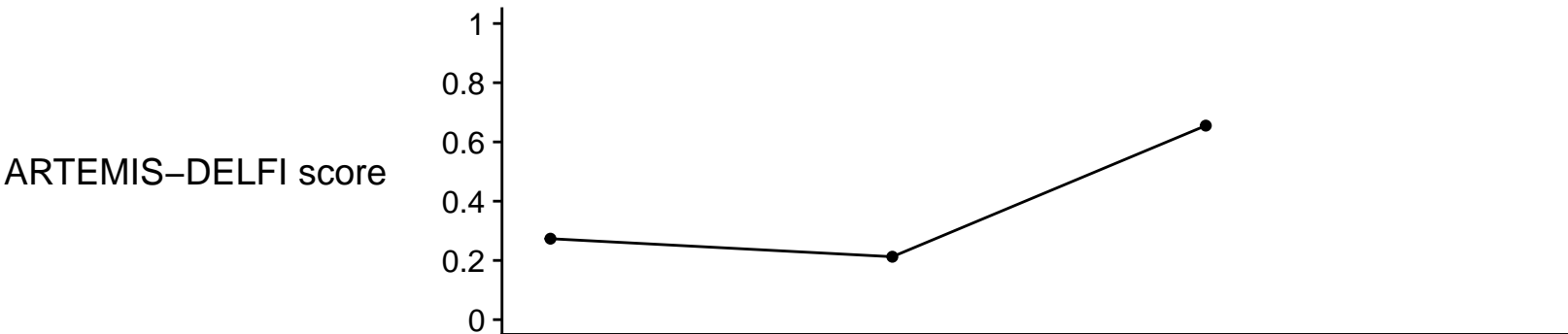


CGPLPA784
WGS data: Tumor (No), Normal (No), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured

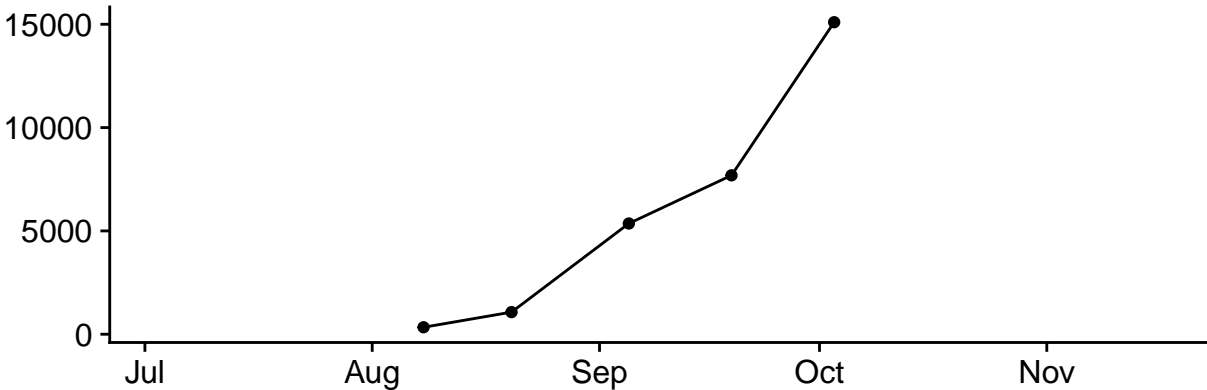


WGMAF (%)

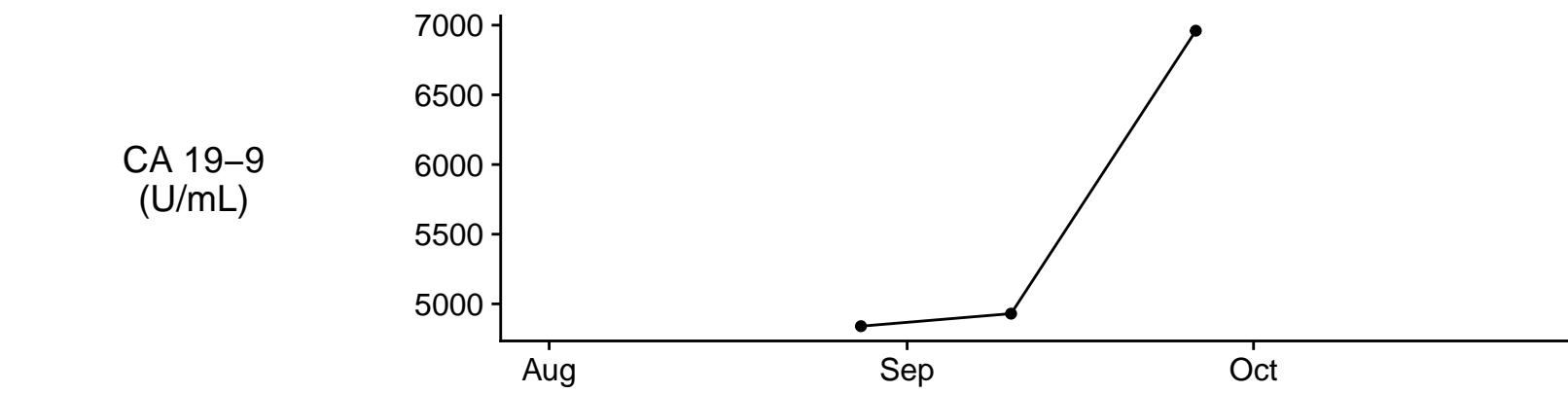
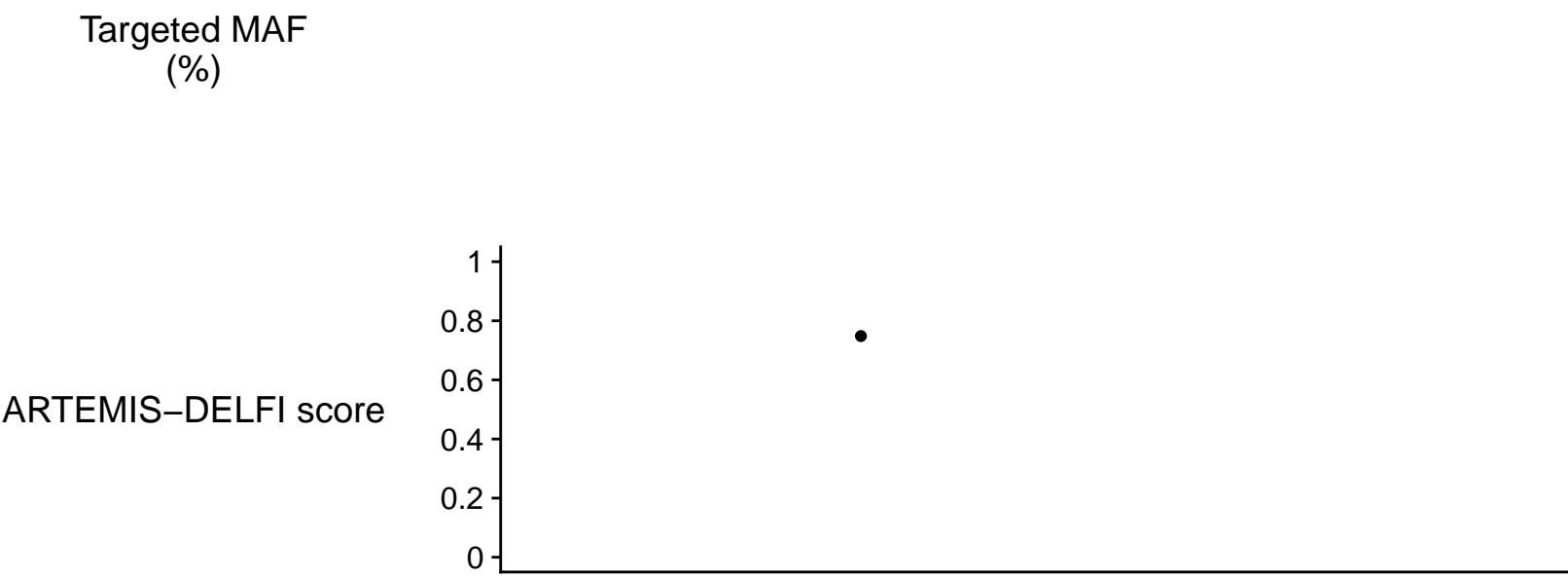
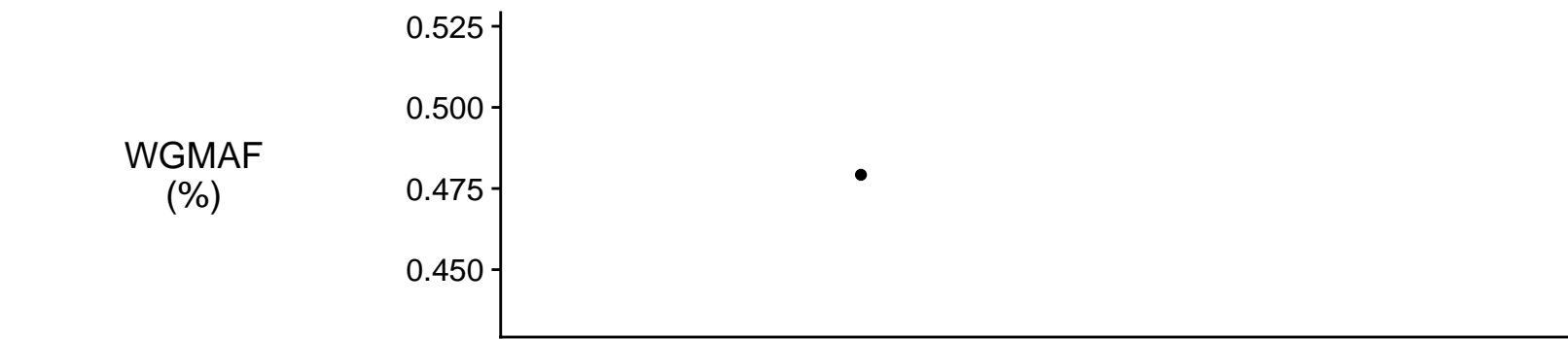
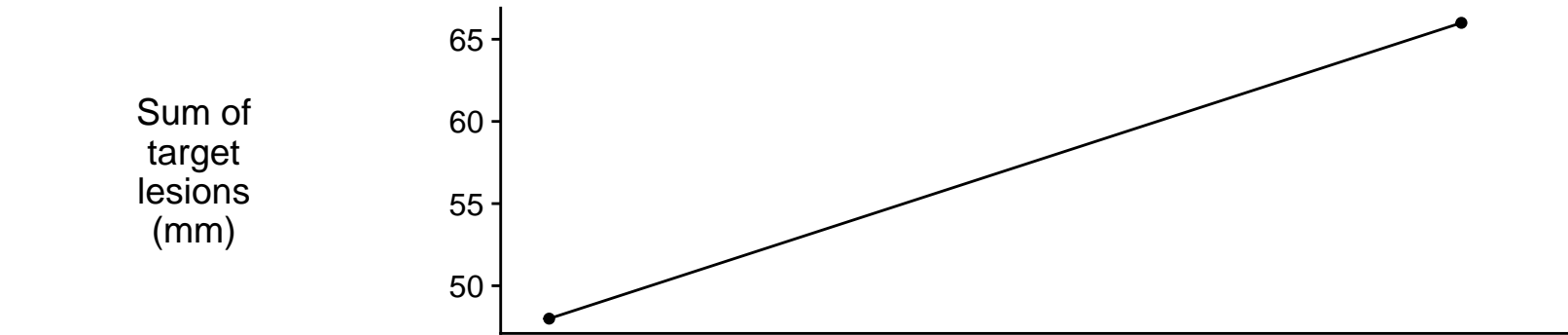
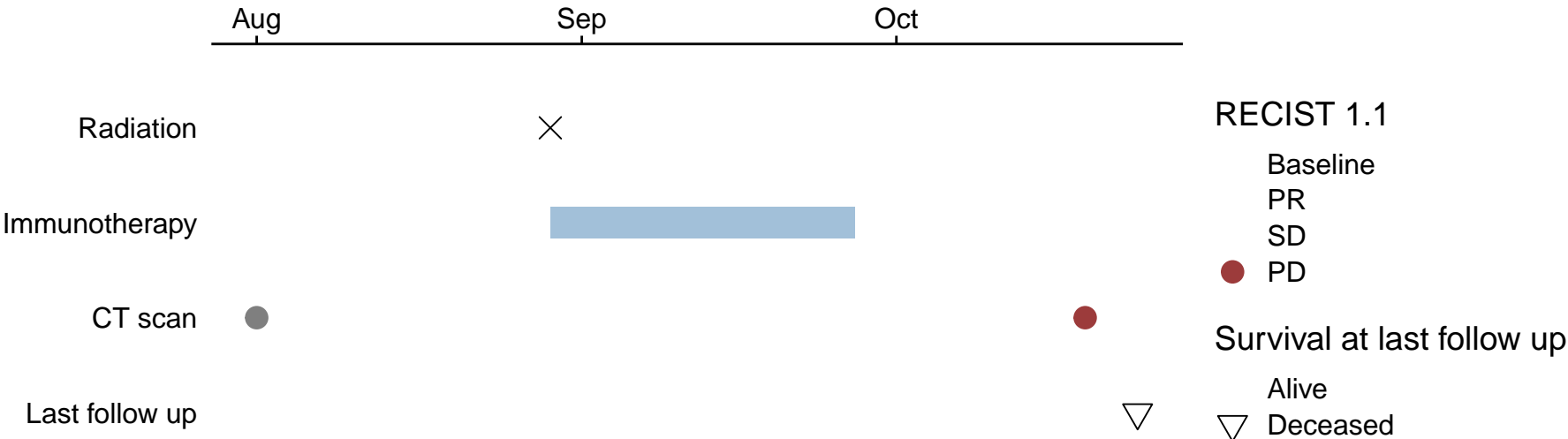
Targeted MAF (%)



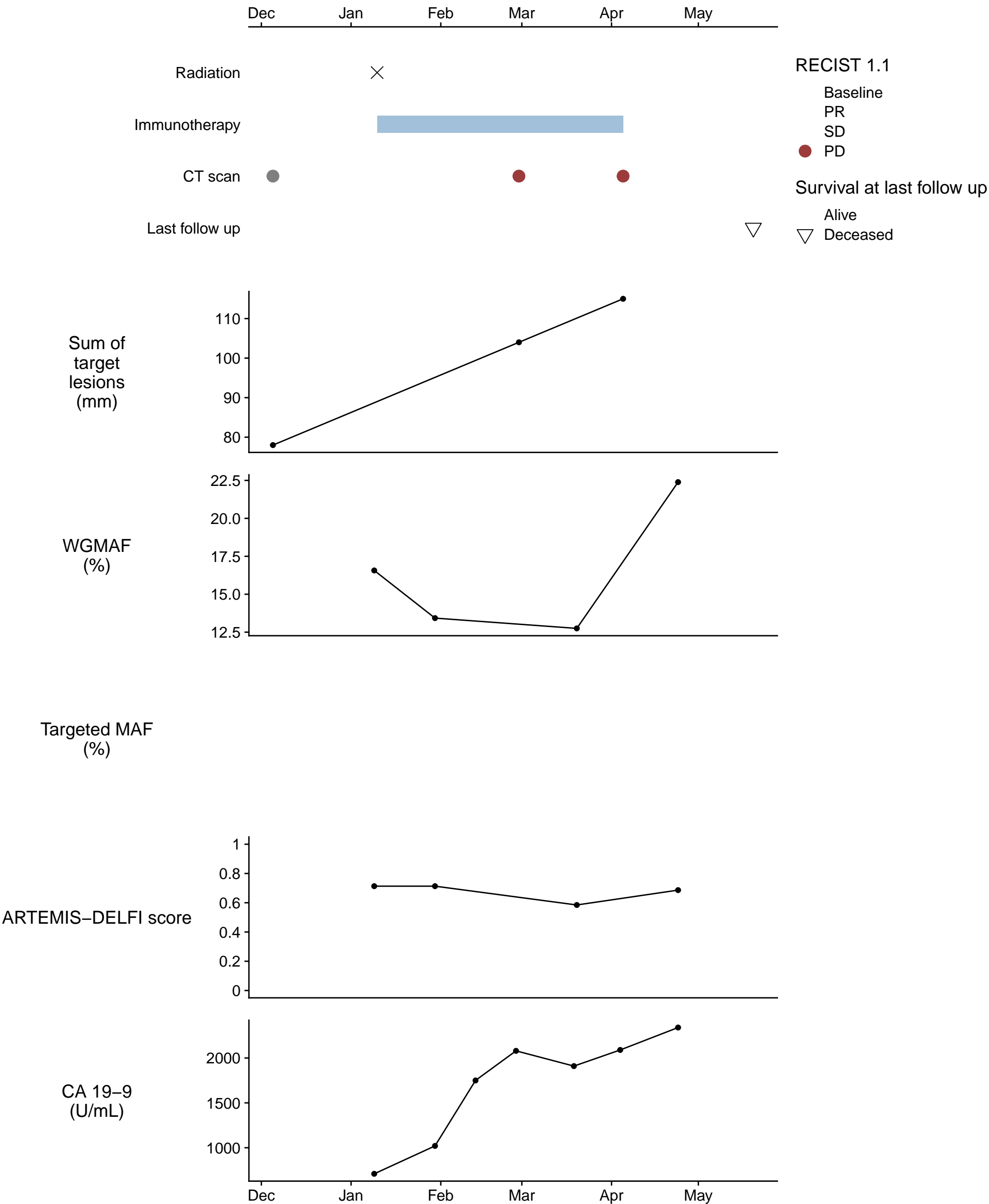
CA 19-9 (U/mL)



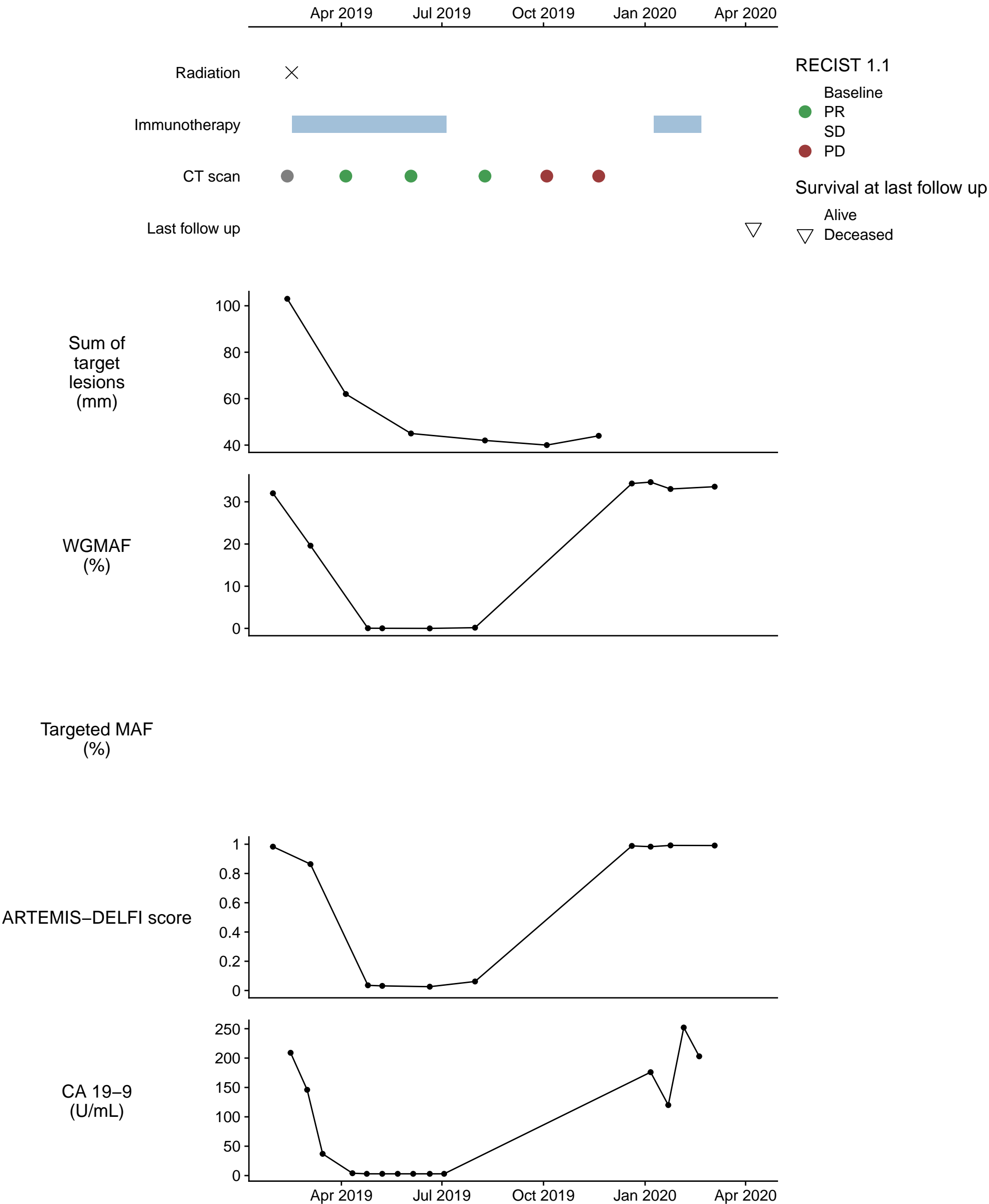
CGPLPA785
WGS data: Tumor (Yes), Normal (Yes), Plasma (1 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 3486, 2105



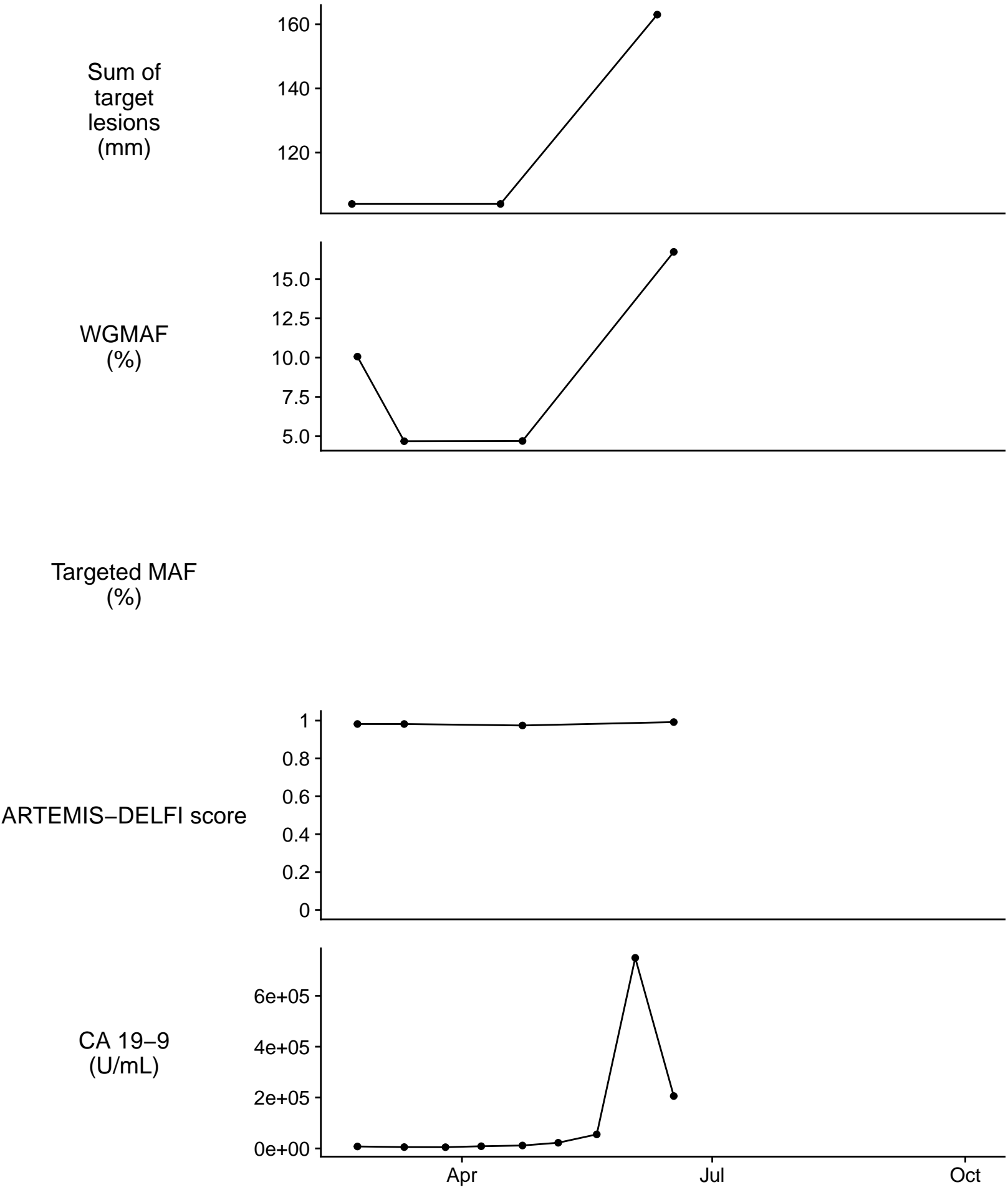
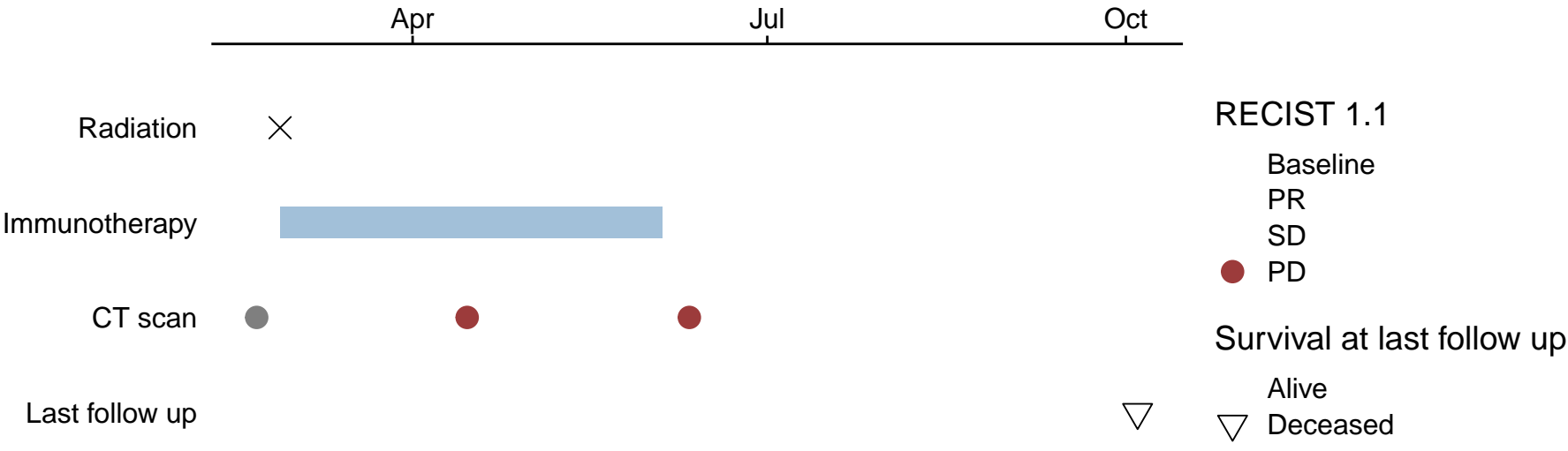
CGPLPA786
WGS data: Tumor (Yes), Normal (Yes), Plasma (4 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 4764, 1554



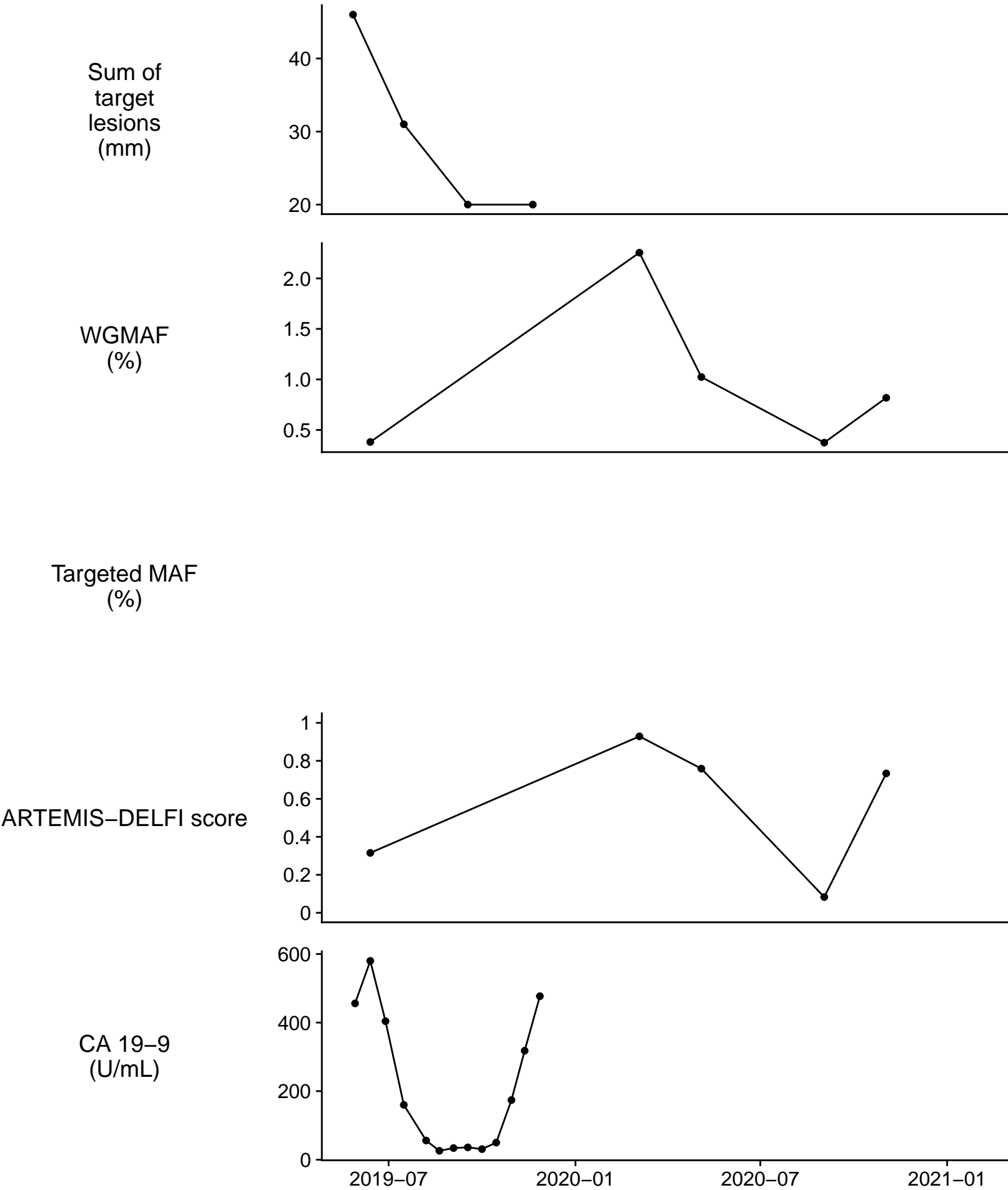
CGPLPA787
WGS data: Tumor (Yes), Normal (Yes), Plasma (10 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 18331, 13335



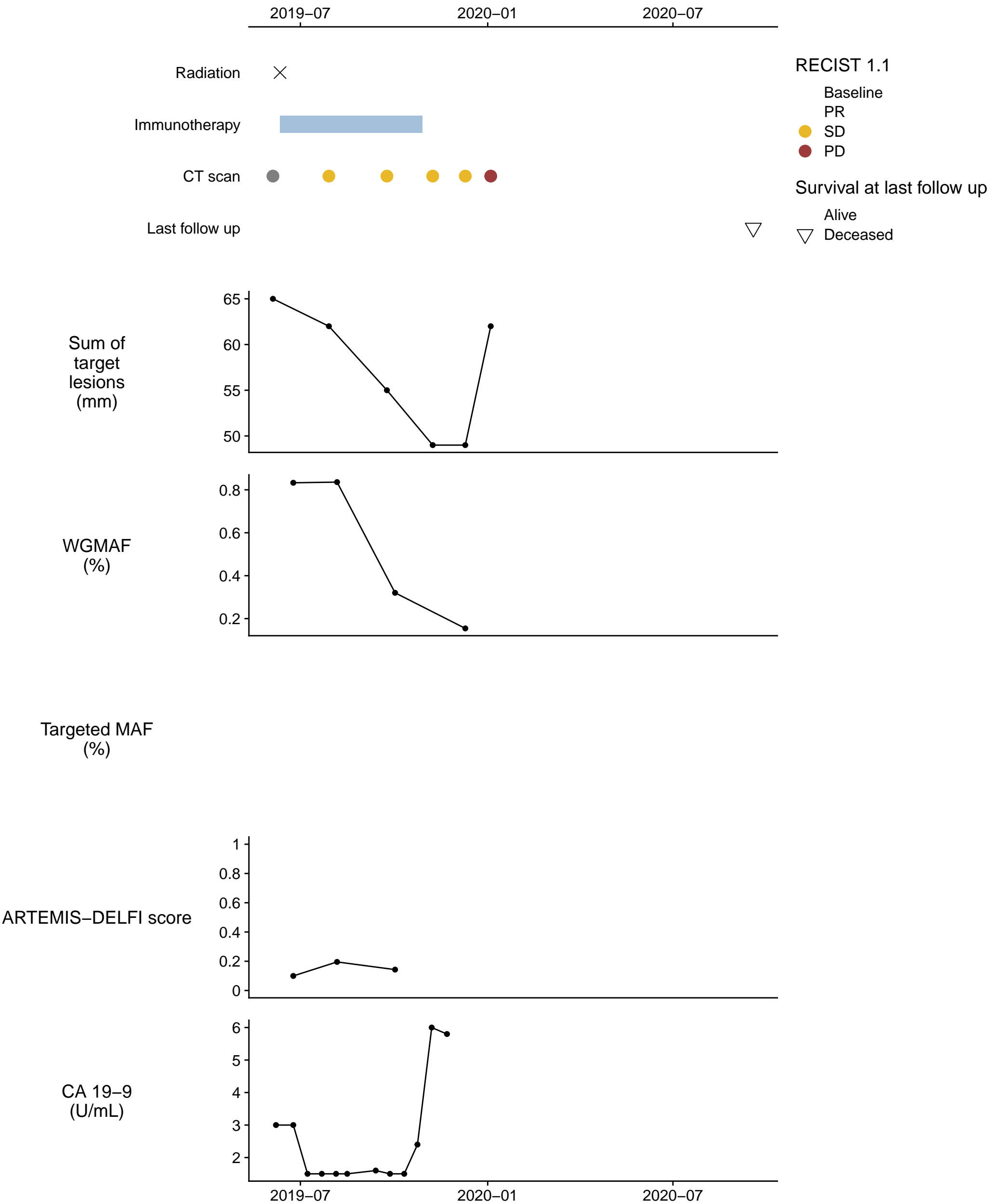
CGPLPA788
WGS data: Tumor (Yes), Normal (Yes), Plasma (4 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 5187, 1124



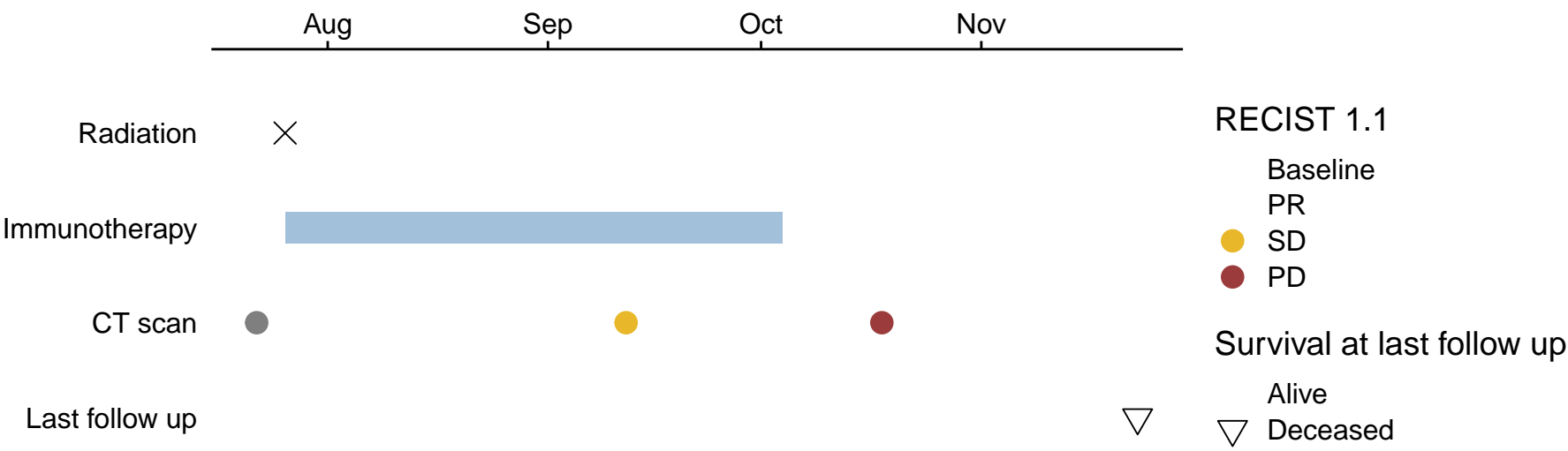
CGPLPA789
WGS data: Tumor (Yes), Normal (Yes), Plasma (5 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 5343, 2806



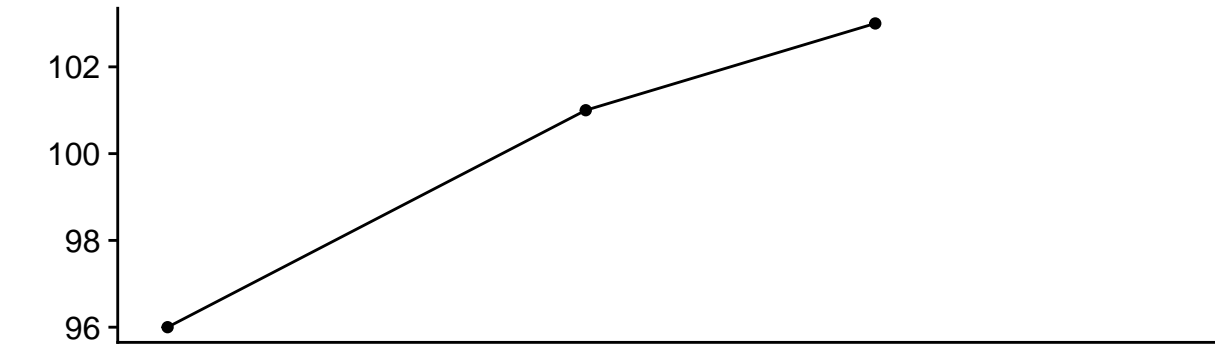
CGPLPA790
WGS data: Tumor (Yes), Normal (Yes), Plasma (4 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 15640, 5730



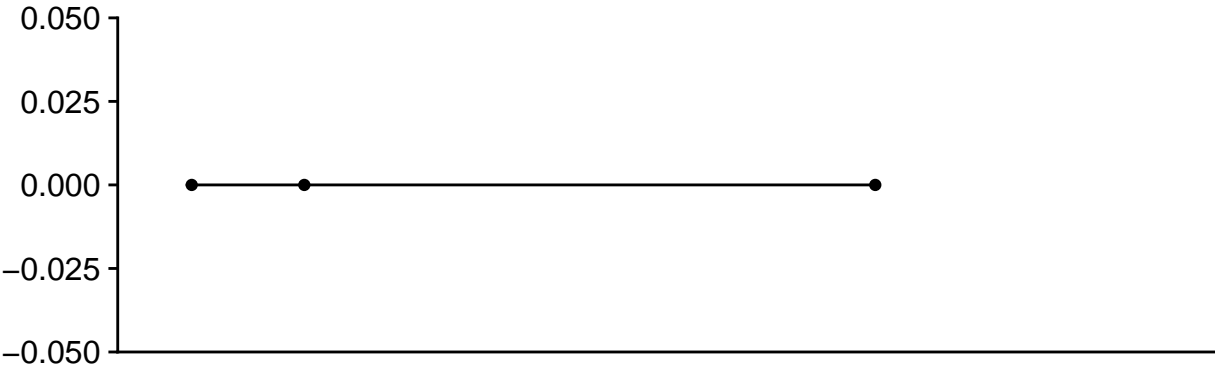
CGPLPA791
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 318, 17



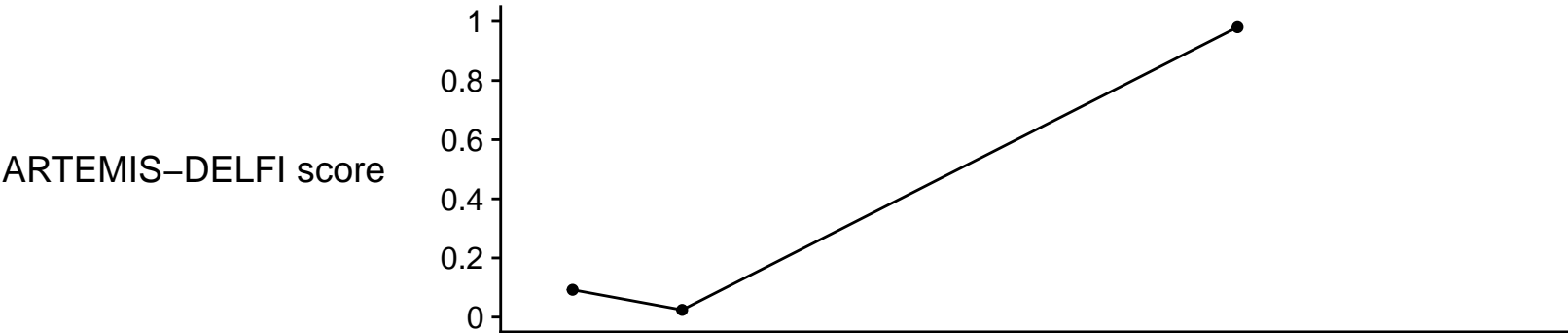
Sum of target lesions (mm)



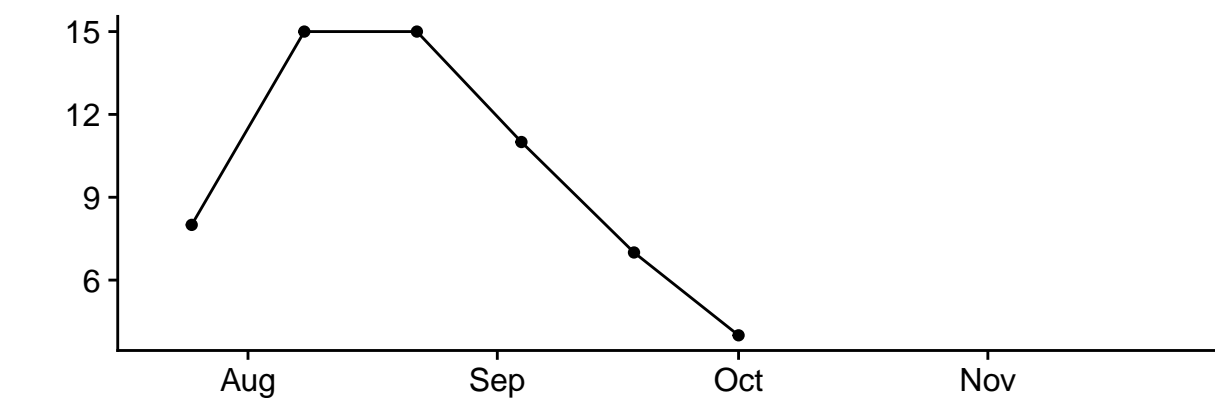
WGMAF (%)



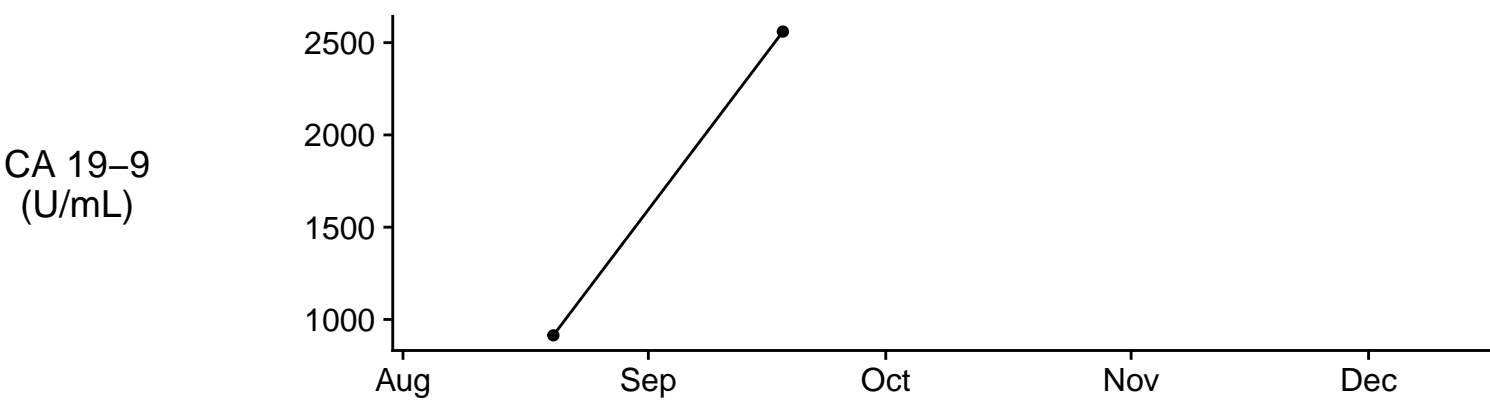
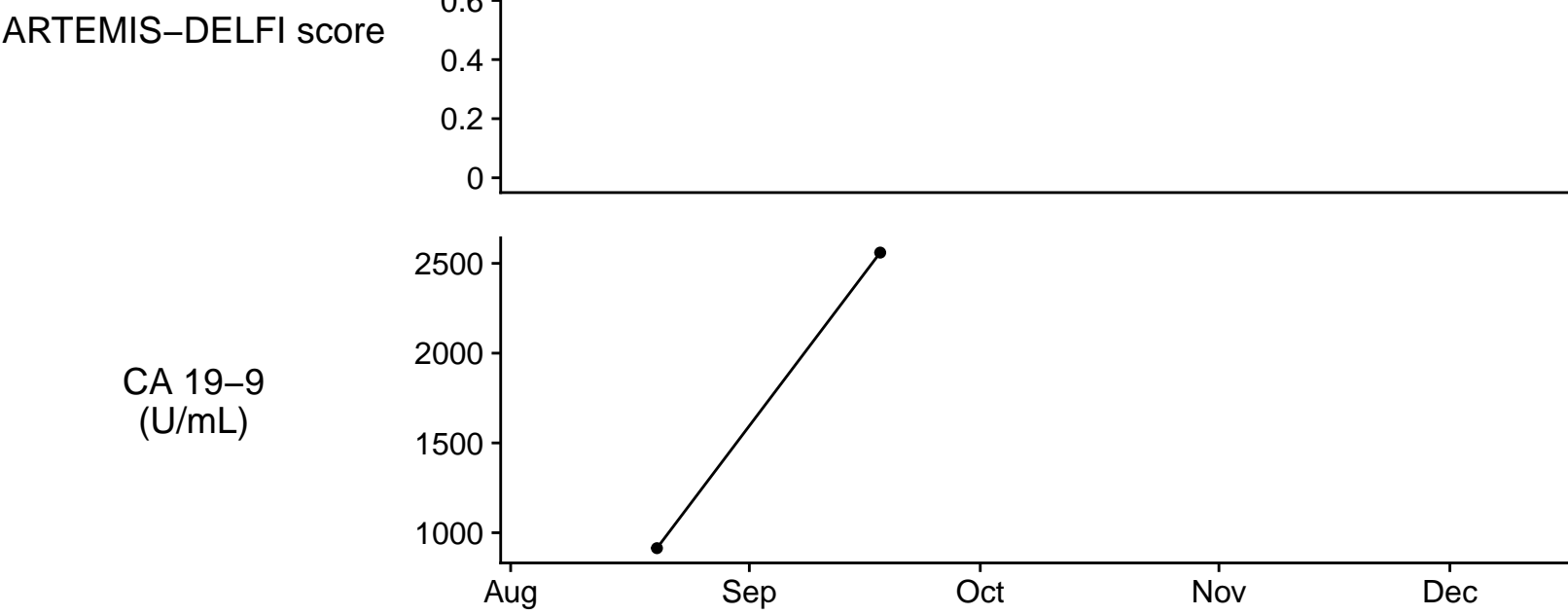
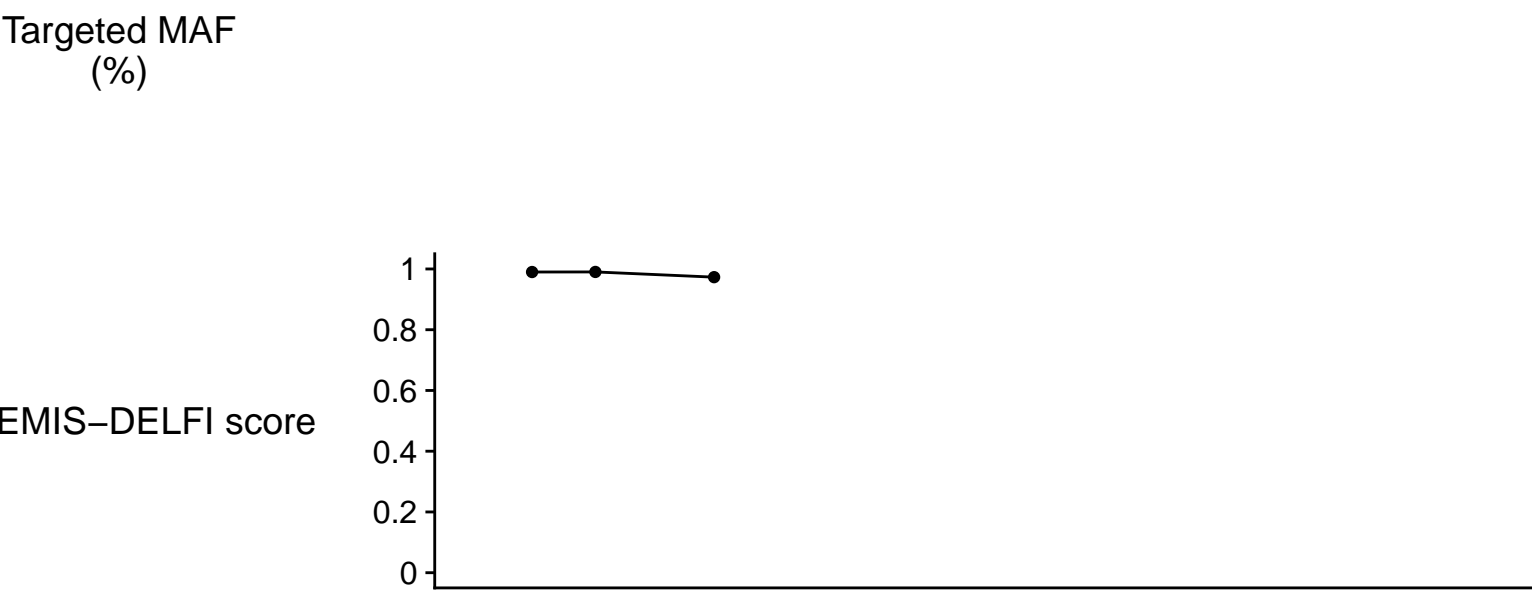
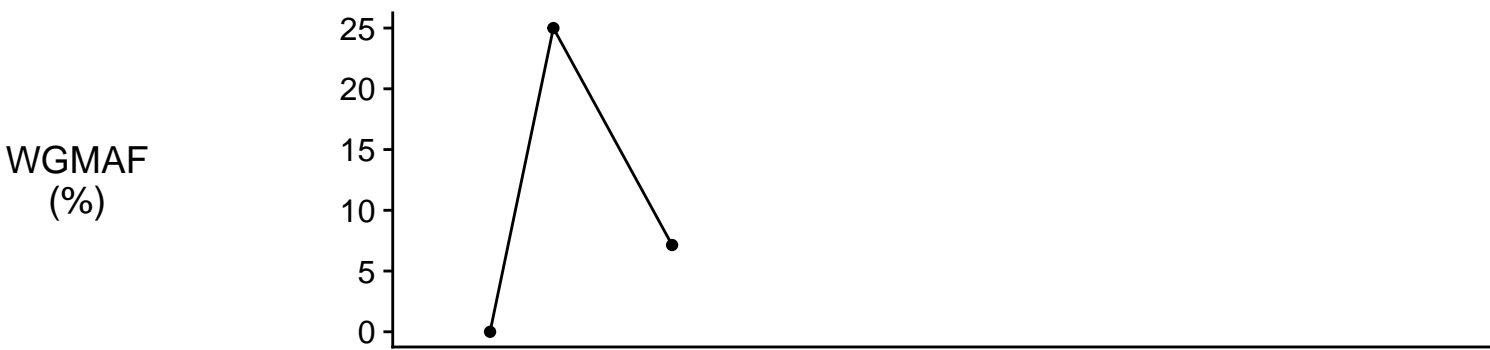
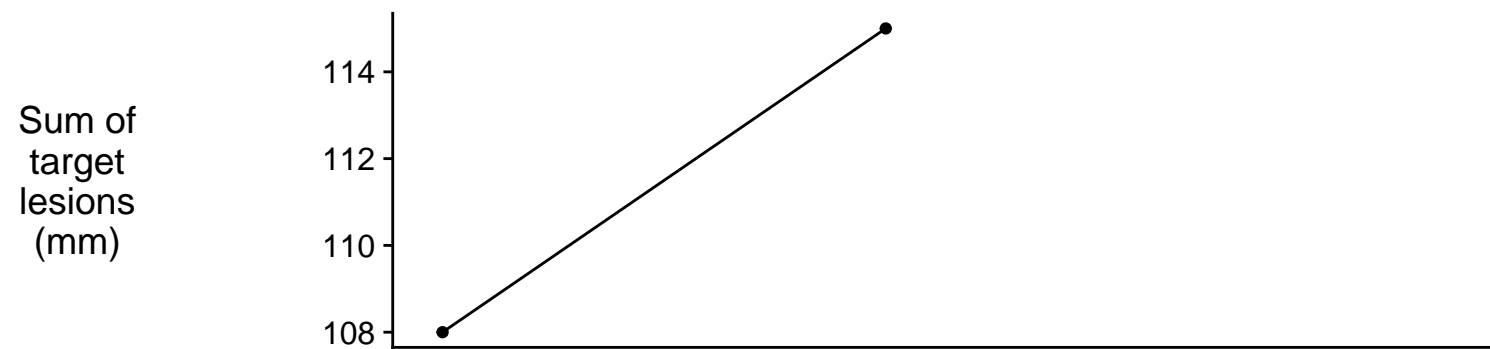
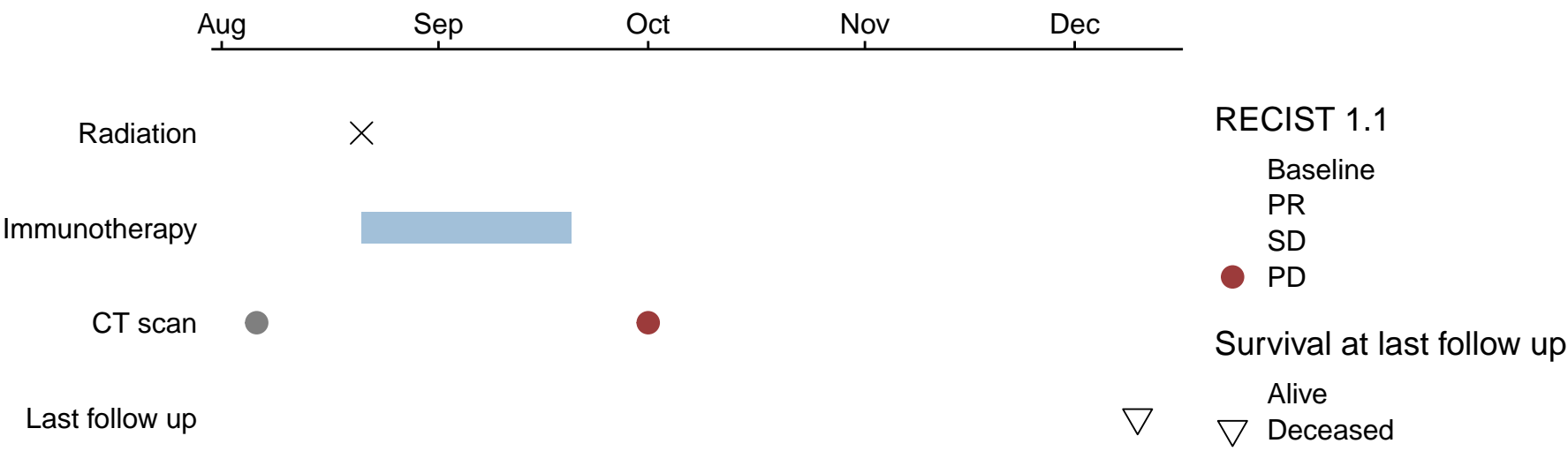
Targeted MAF (%)



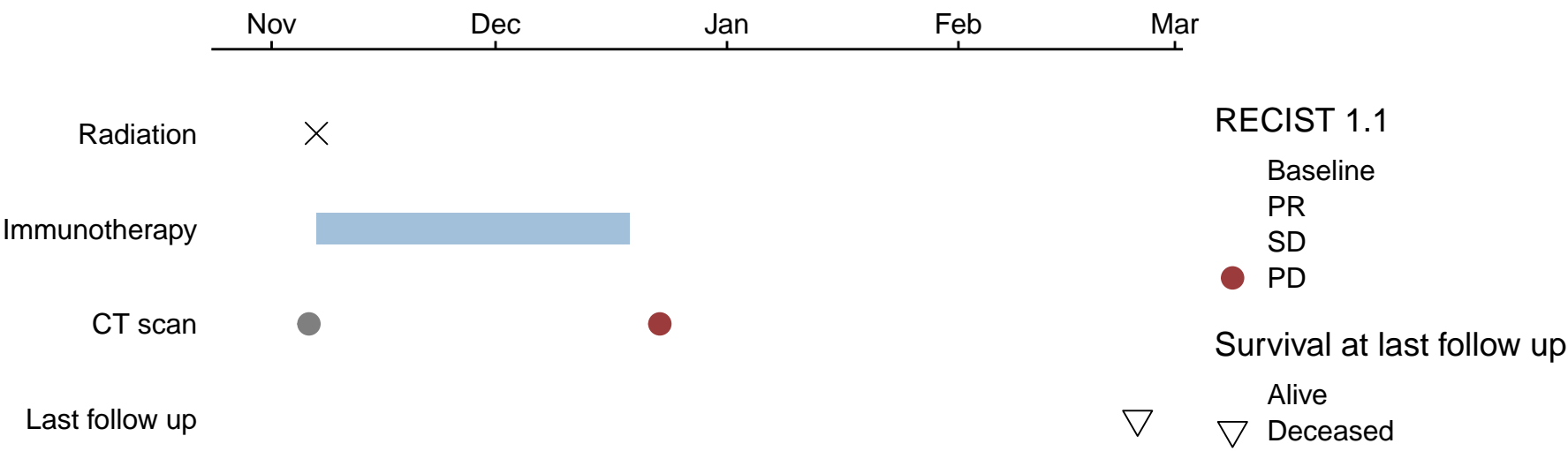
CA 19-9 (U/mL)



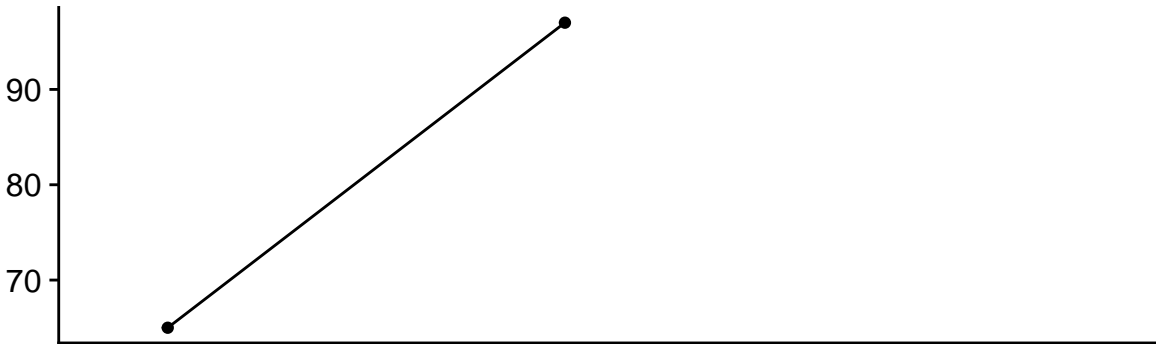
CGPLPA792
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 625, 47



CGPLPA793
WGS data: Tumor (No), Normal (No), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): not measured



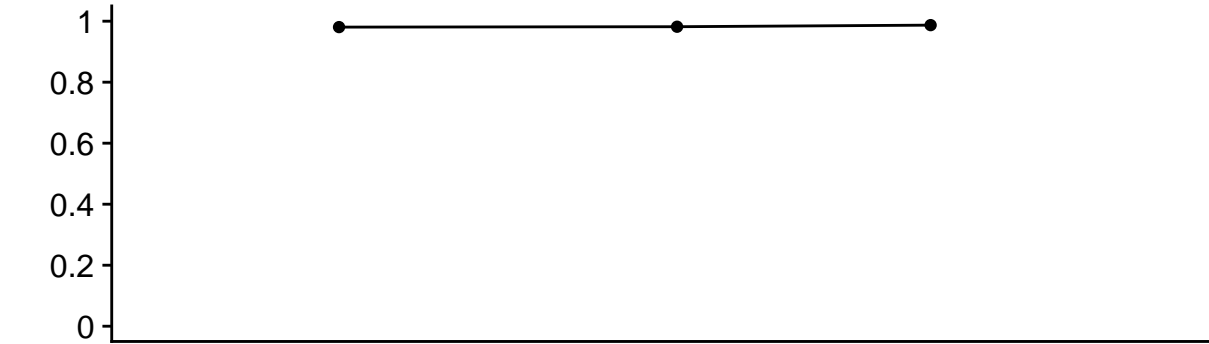
Sum of target lesions (mm)



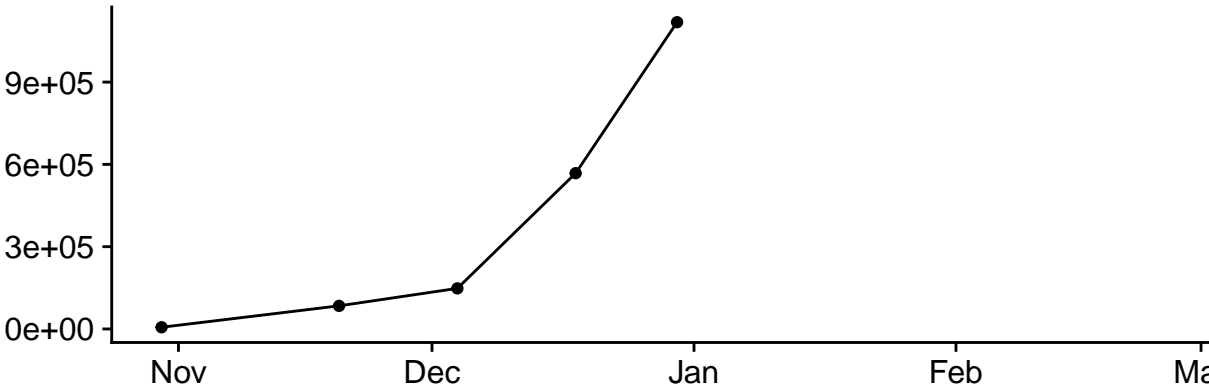
WGMAF (%)

Targeted MAF (%)

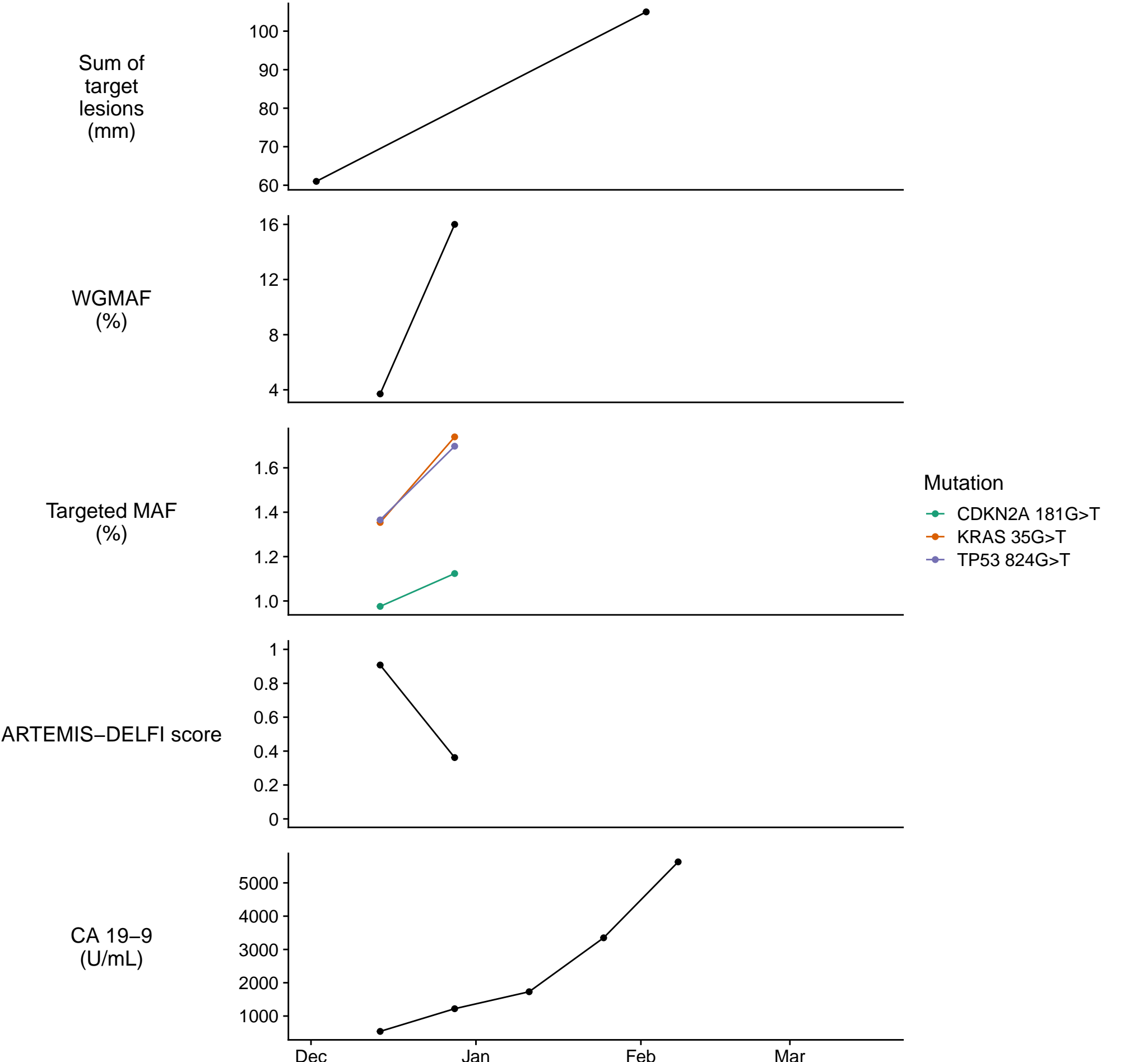
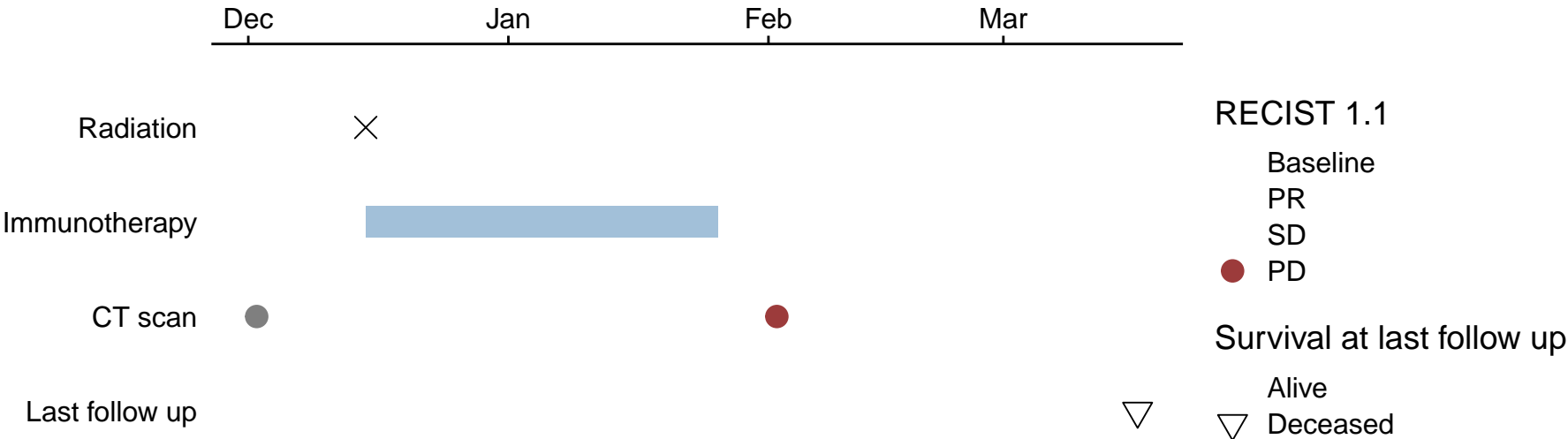
ARTEMIS-DELFI score



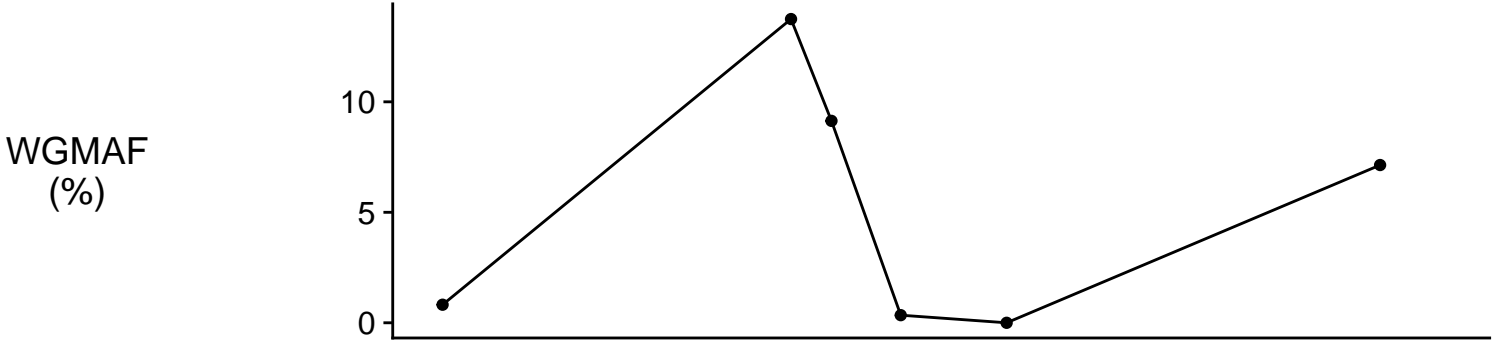
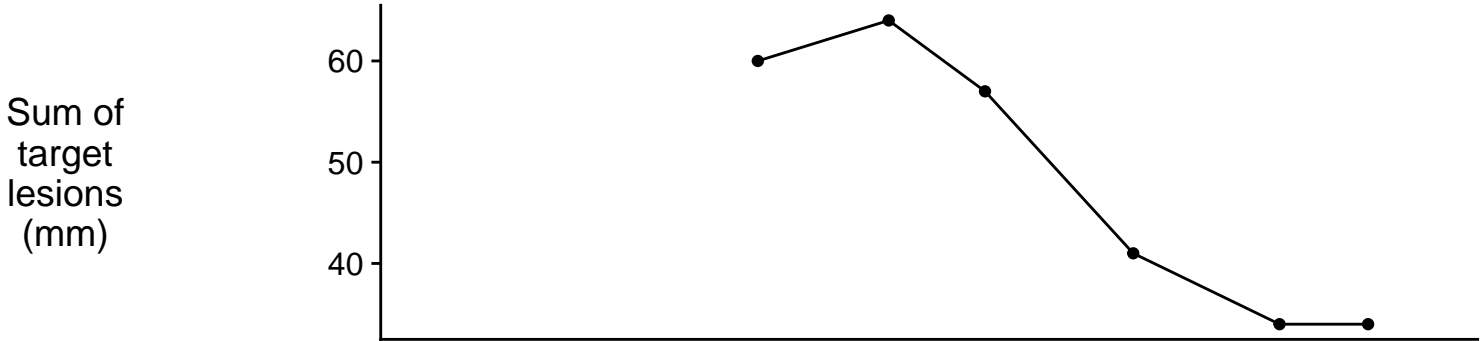
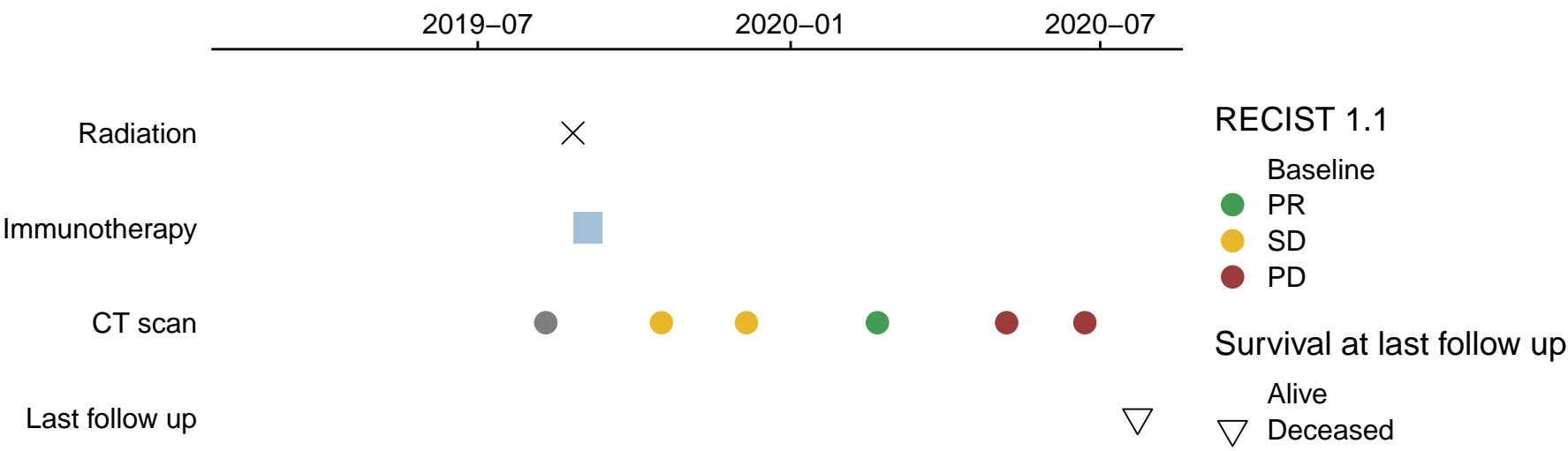
CA 19-9 (U/mL)



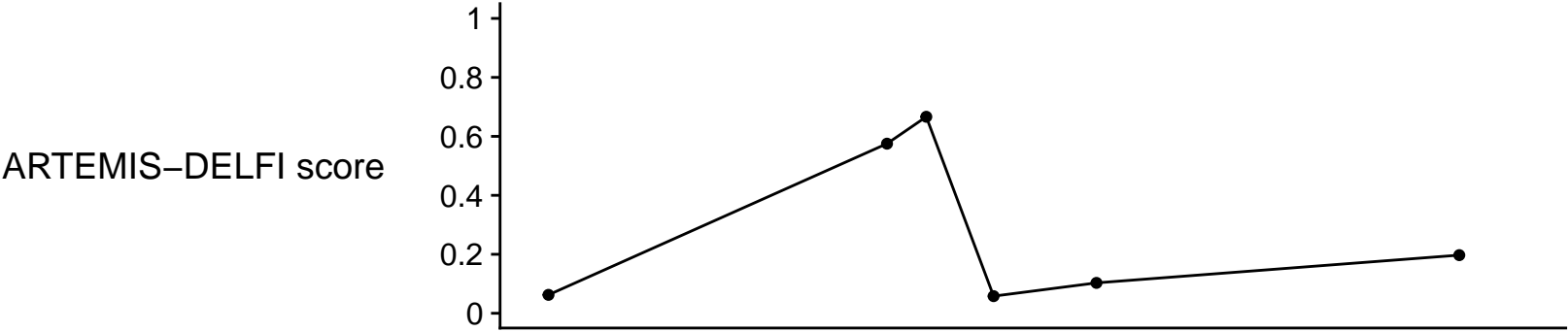
CGPLPA218
WGS data: Tumor (Yes), Normal (Yes), Plasma (2 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 194, 96



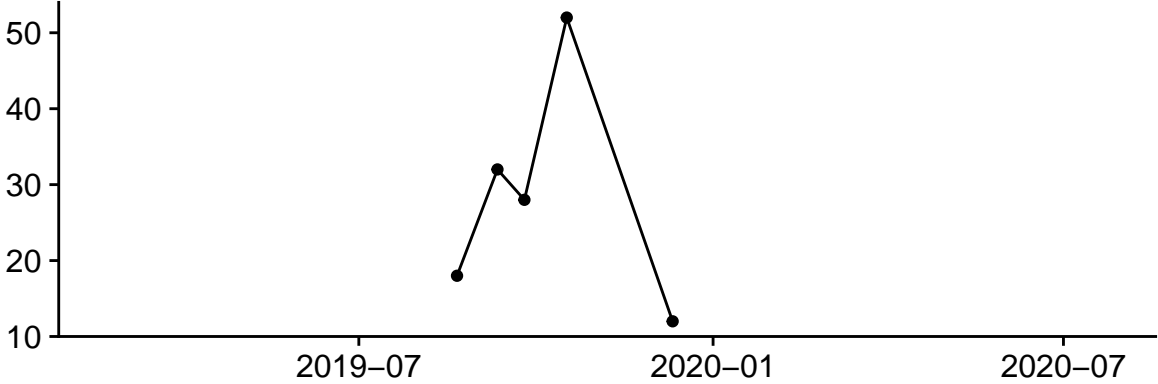
CGPLPA794
WGS data: Tumor (Yes), Normal (Yes), Plasma (6 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 8185, 5648

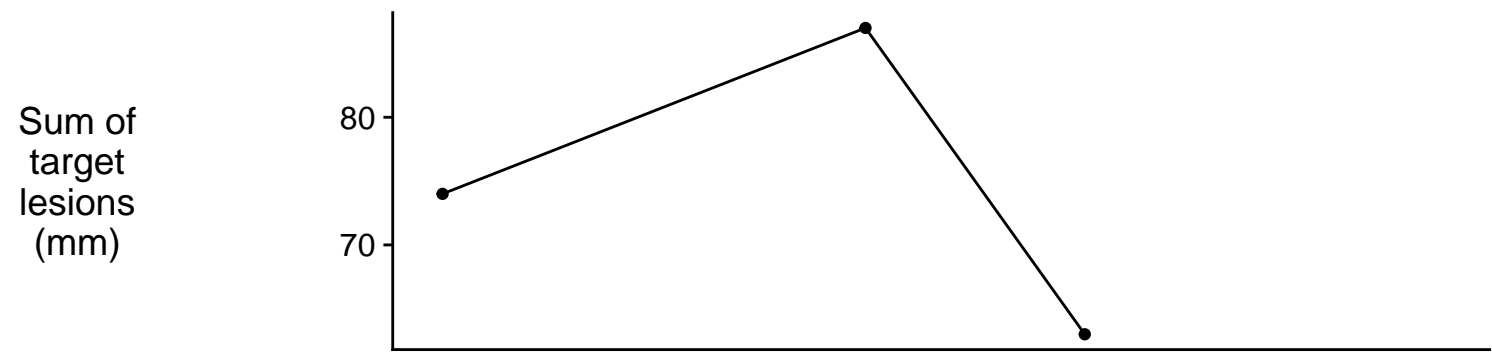
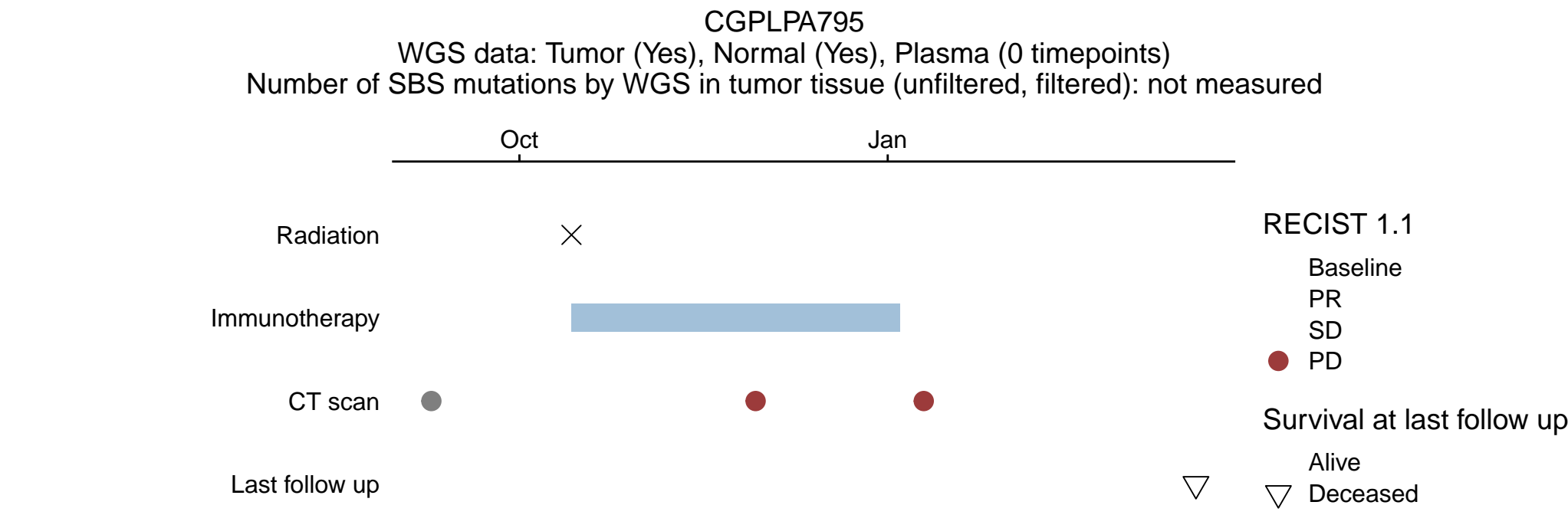


Targeted MAF (%)



CA 19-9 (U/mL)

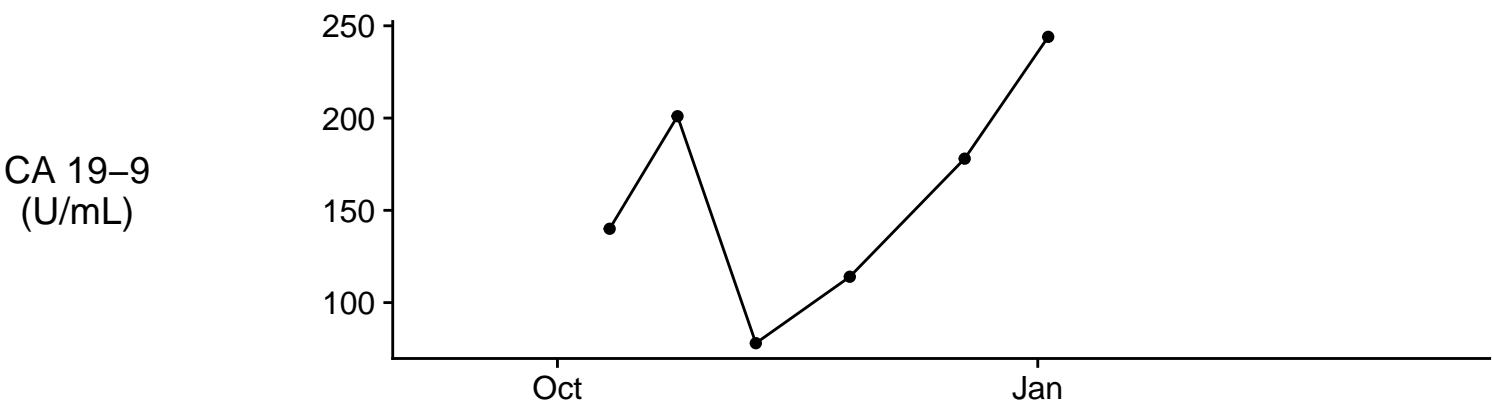




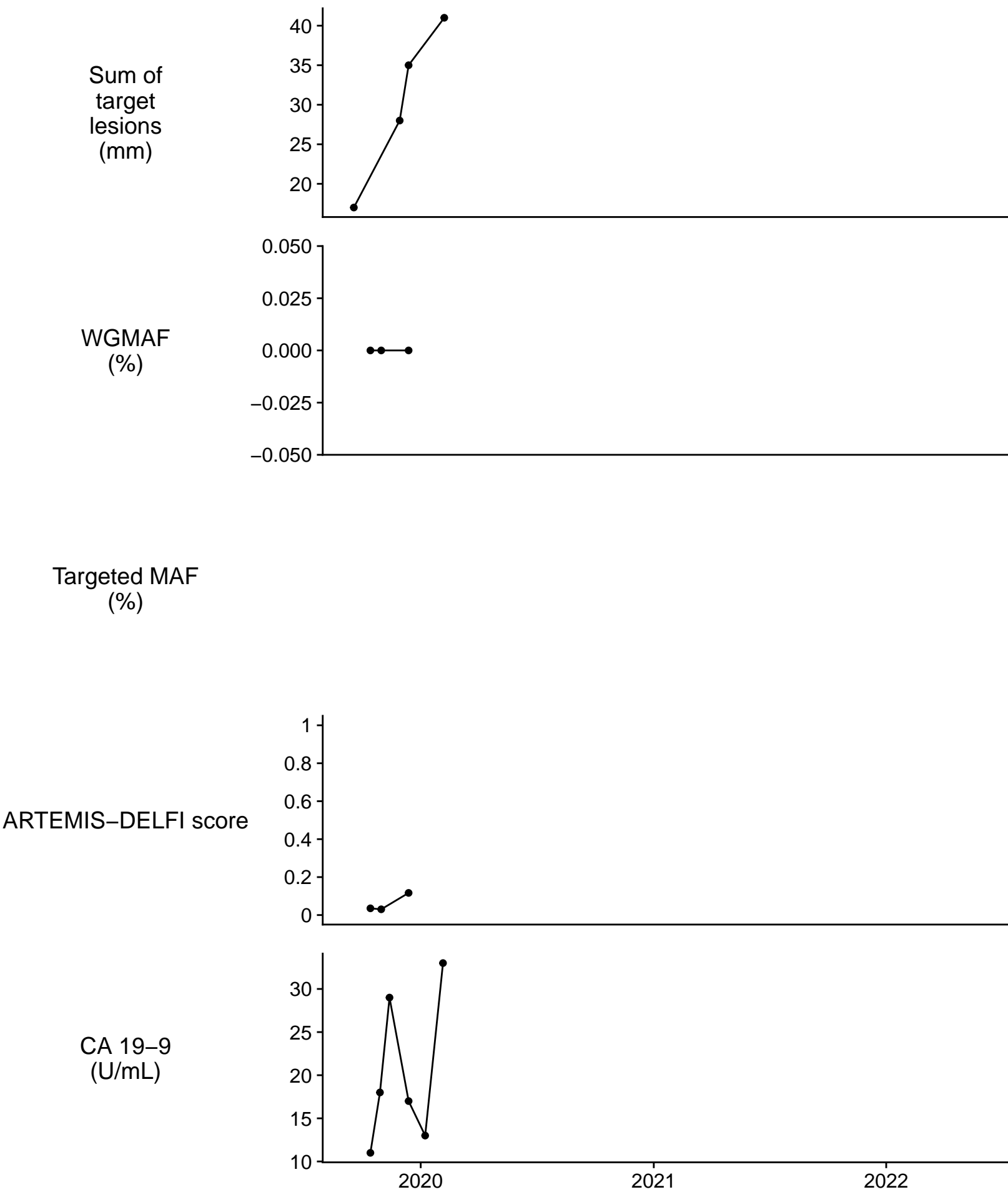
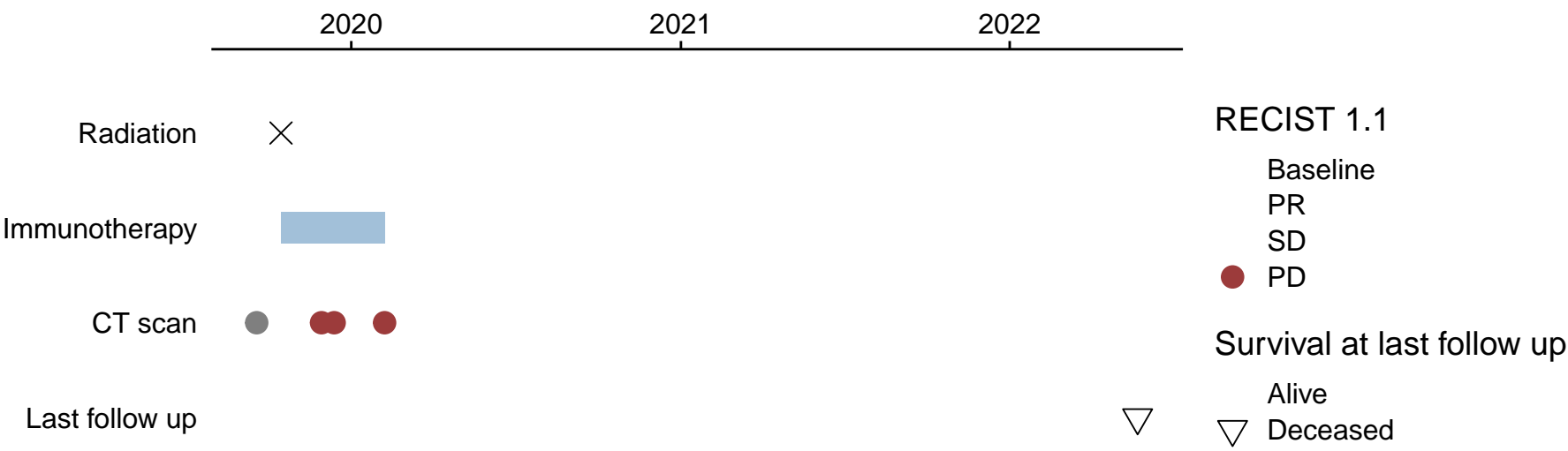
WGMAF (%)

Targeted MAF (%)

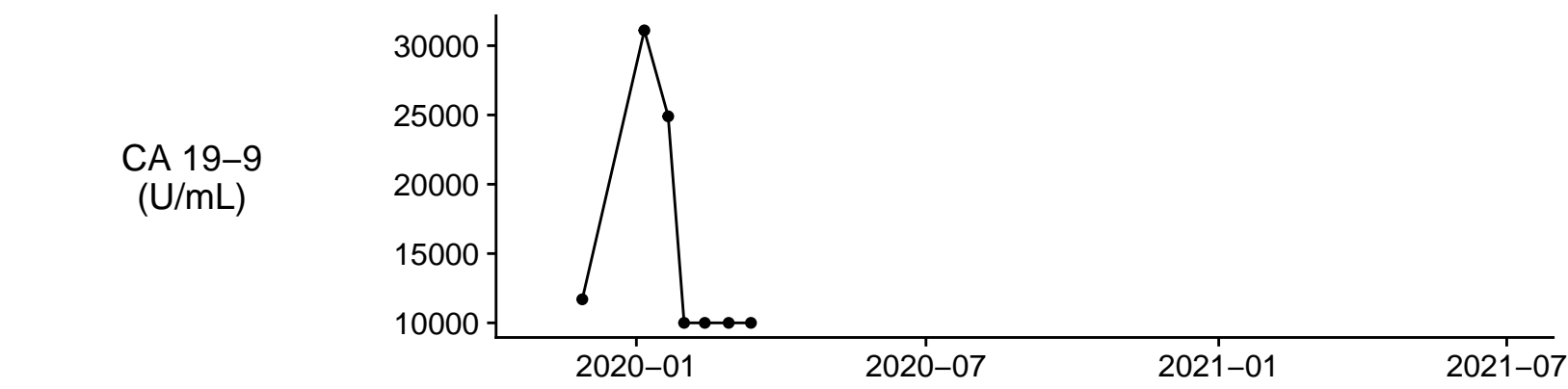
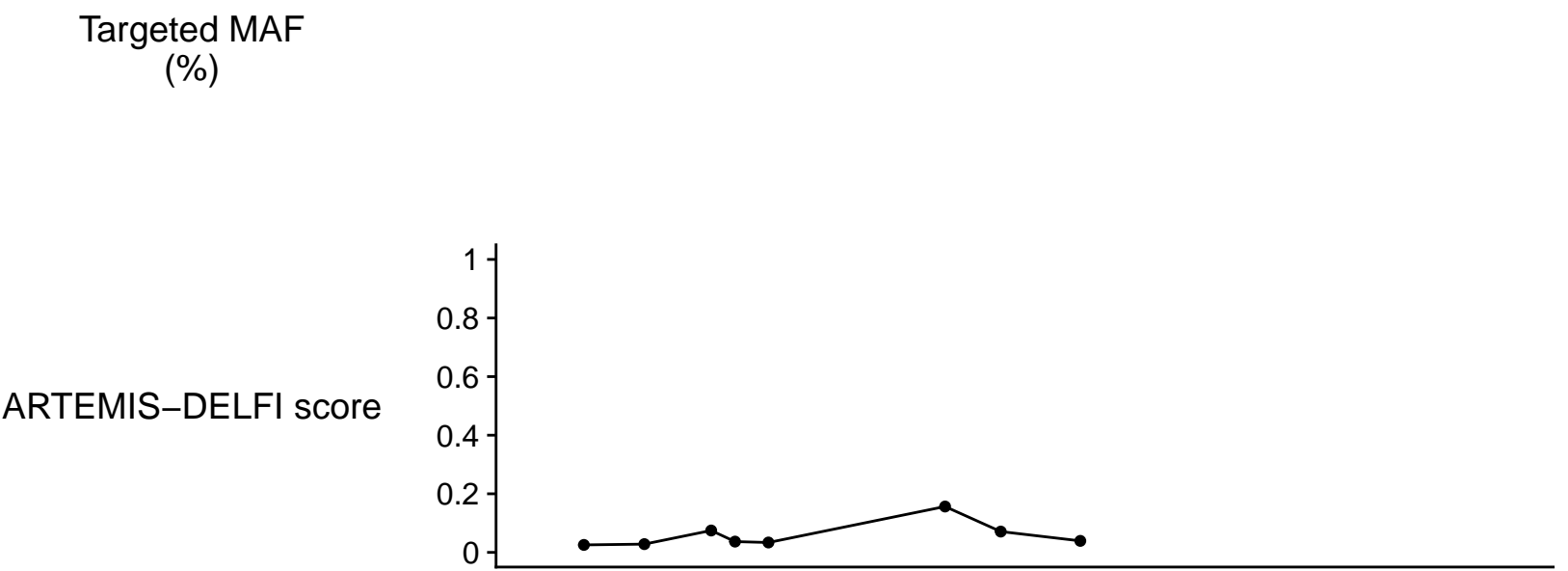
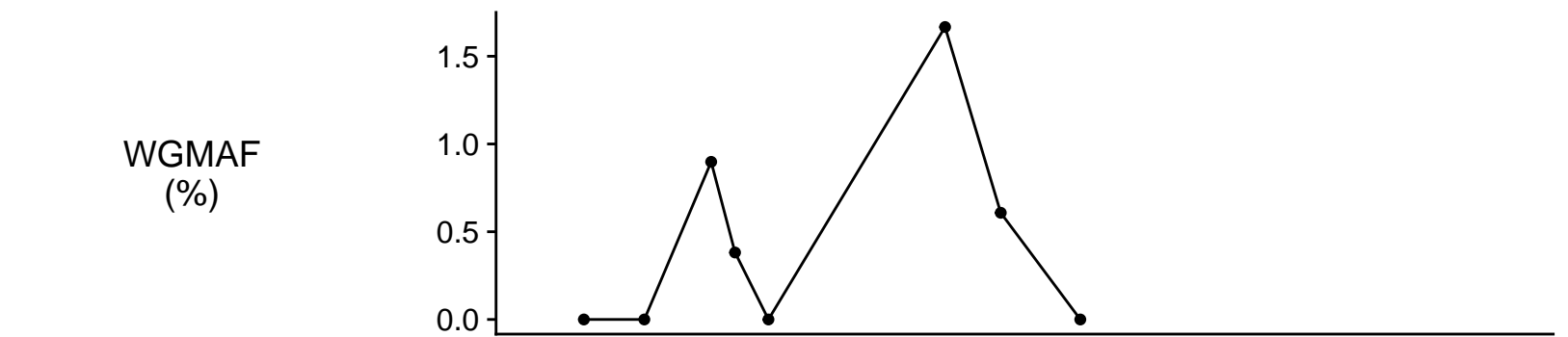
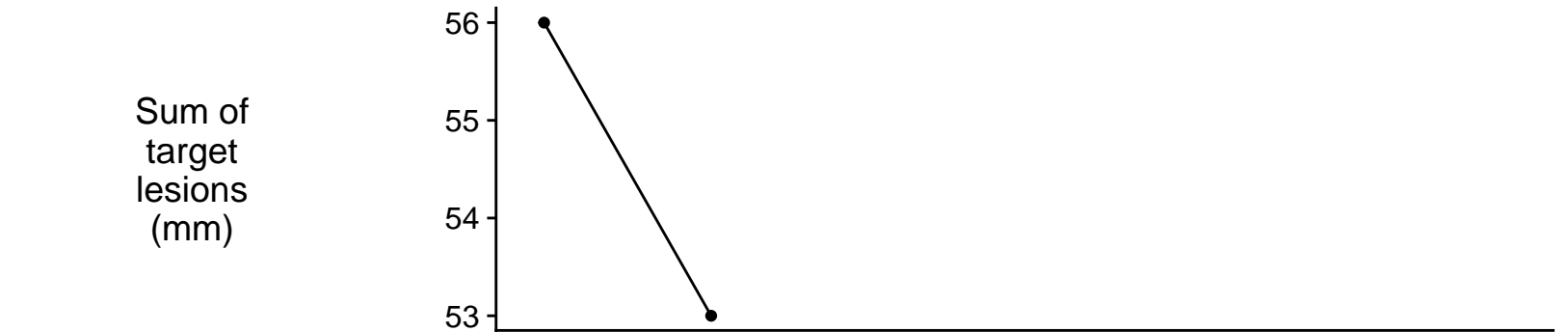
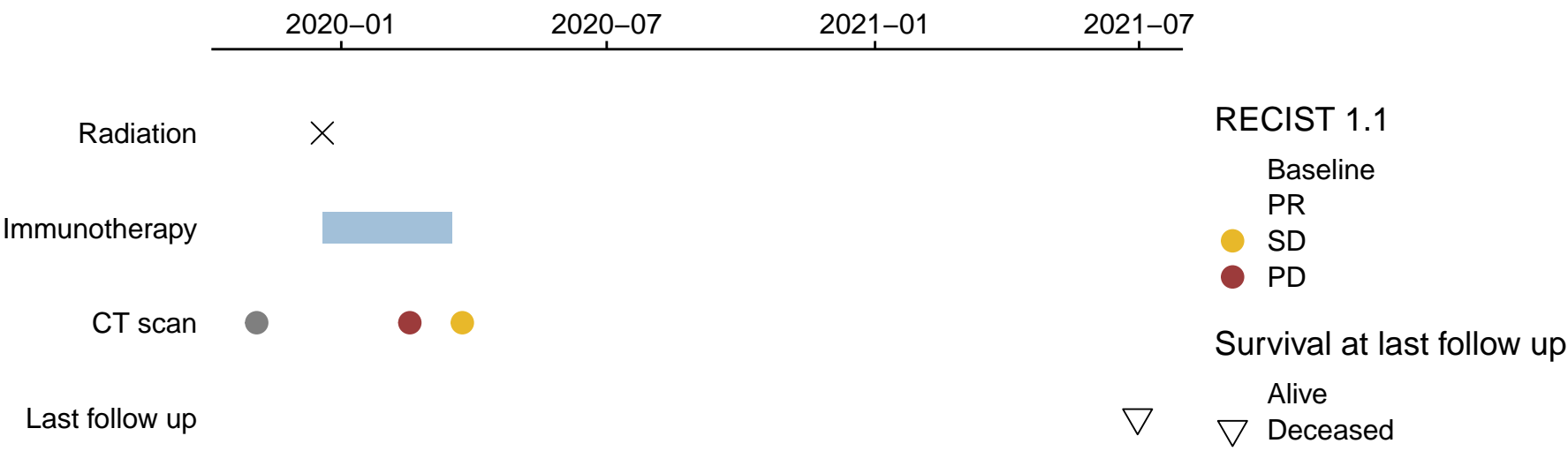
ARTEMIS-DELFI score



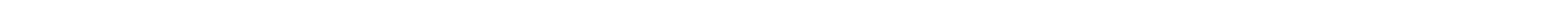
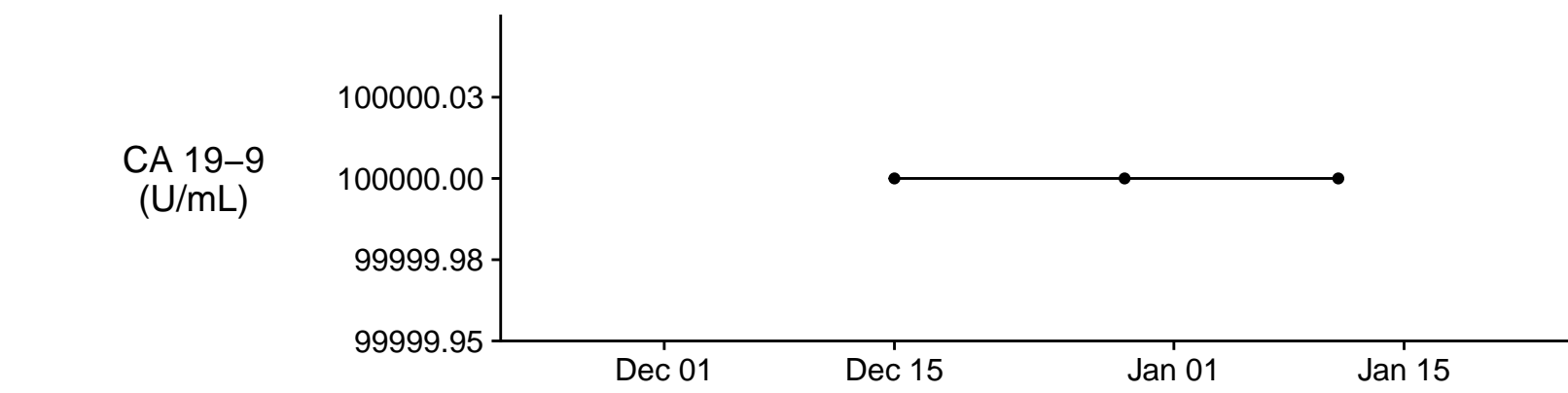
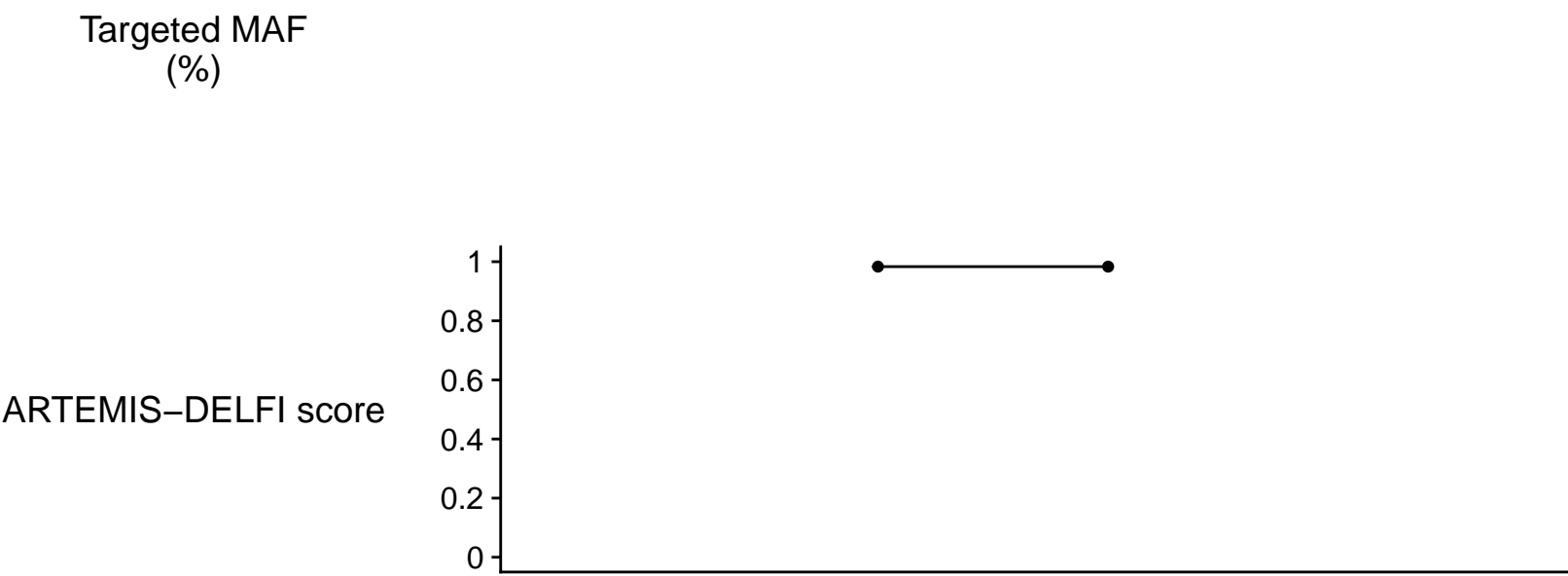
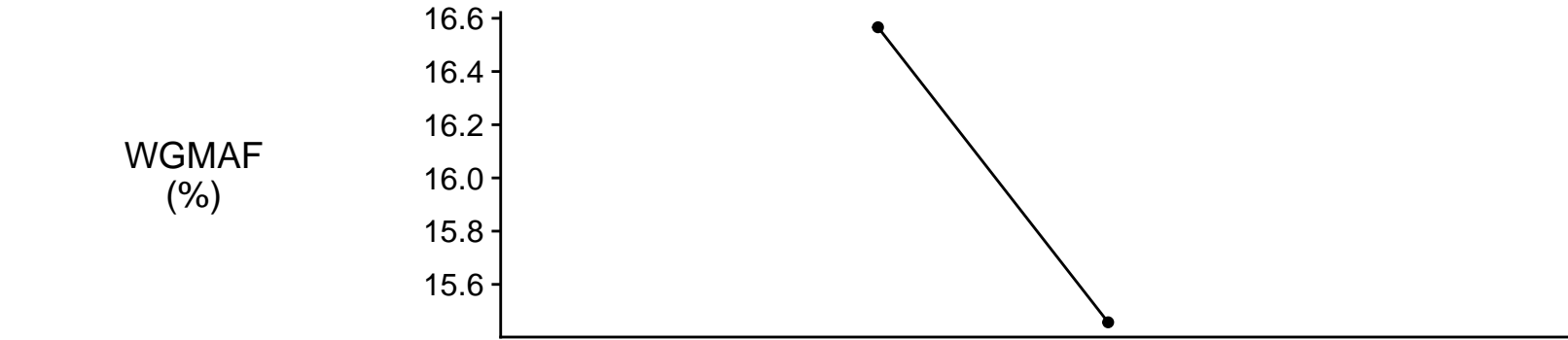
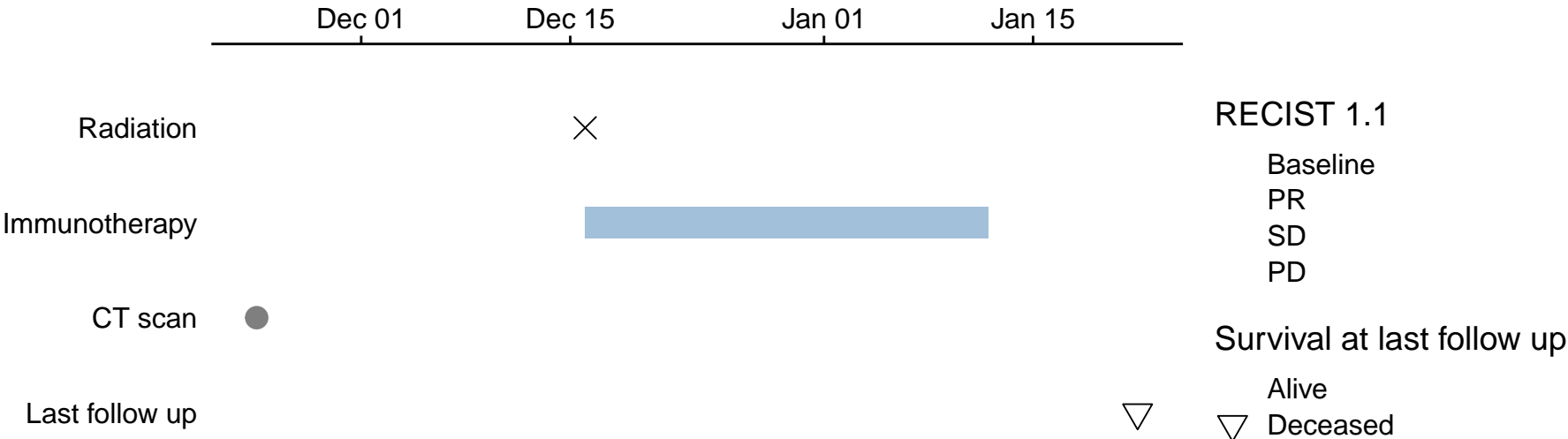
CGPLPA796
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 478, 17



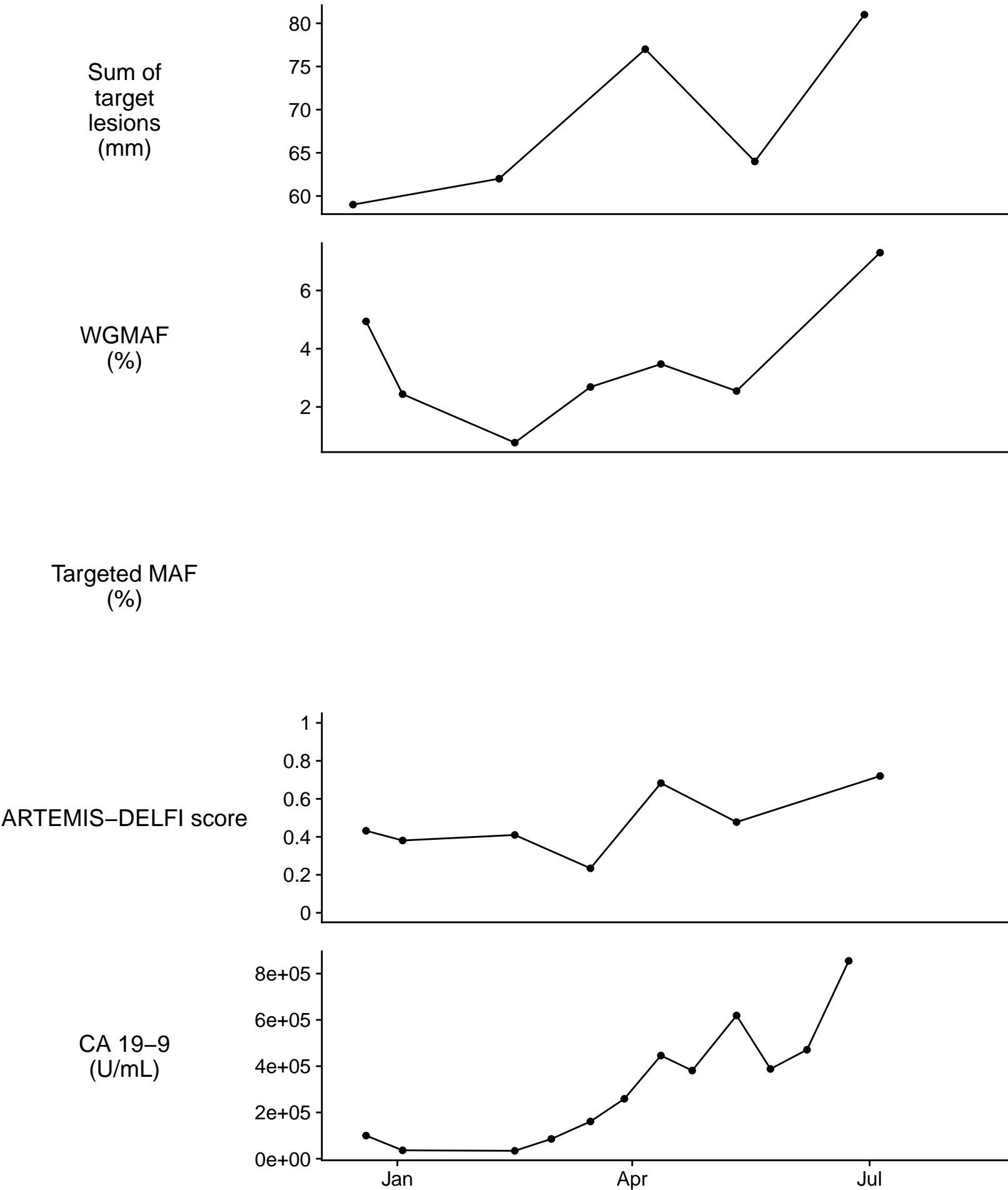
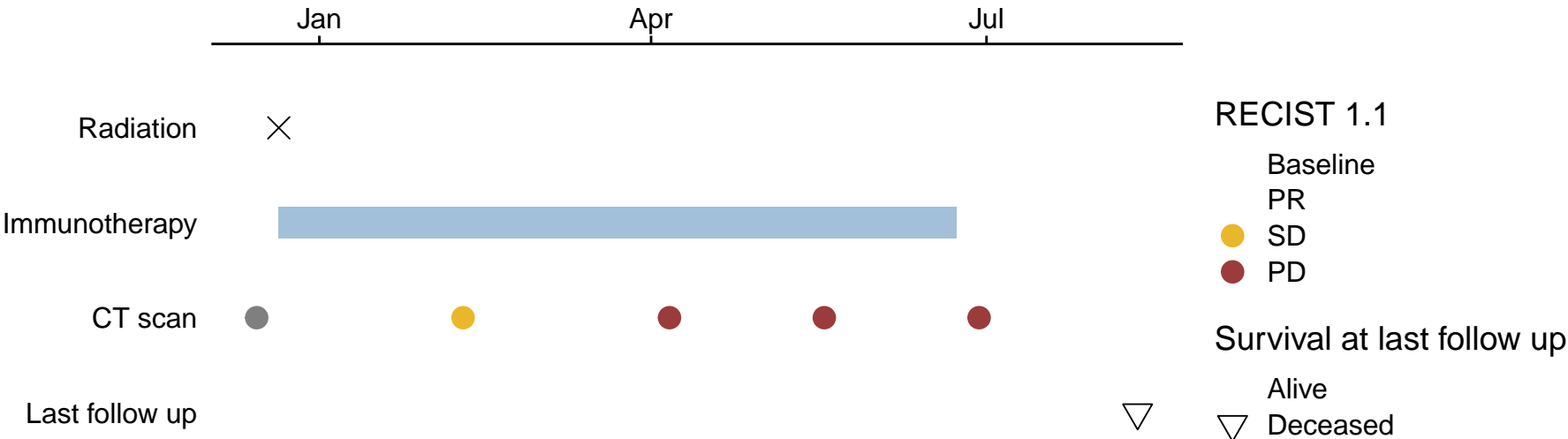
CGPLPA797
WGS data: Tumor (Yes), Normal (Yes), Plasma (8 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 1623, 524



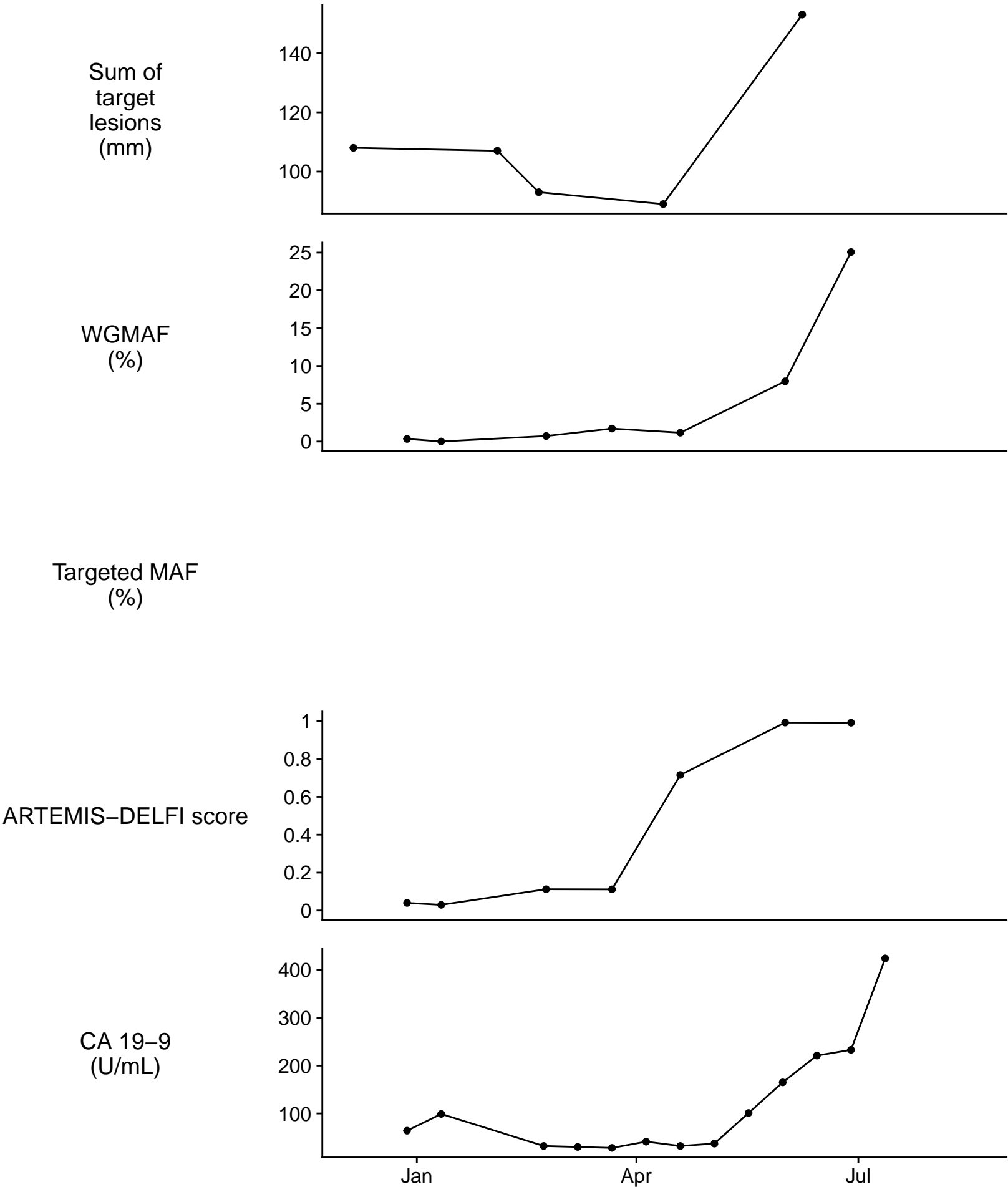
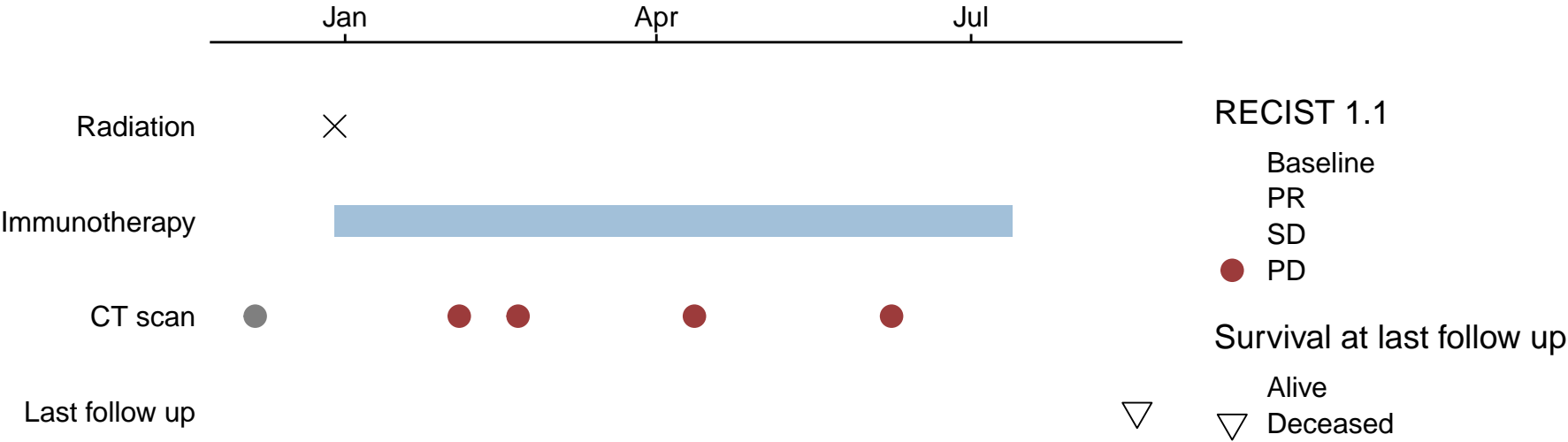
CGPLPA766
WGS data: Tumor (Yes), Normal (Yes), Plasma (2 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 5133, 654



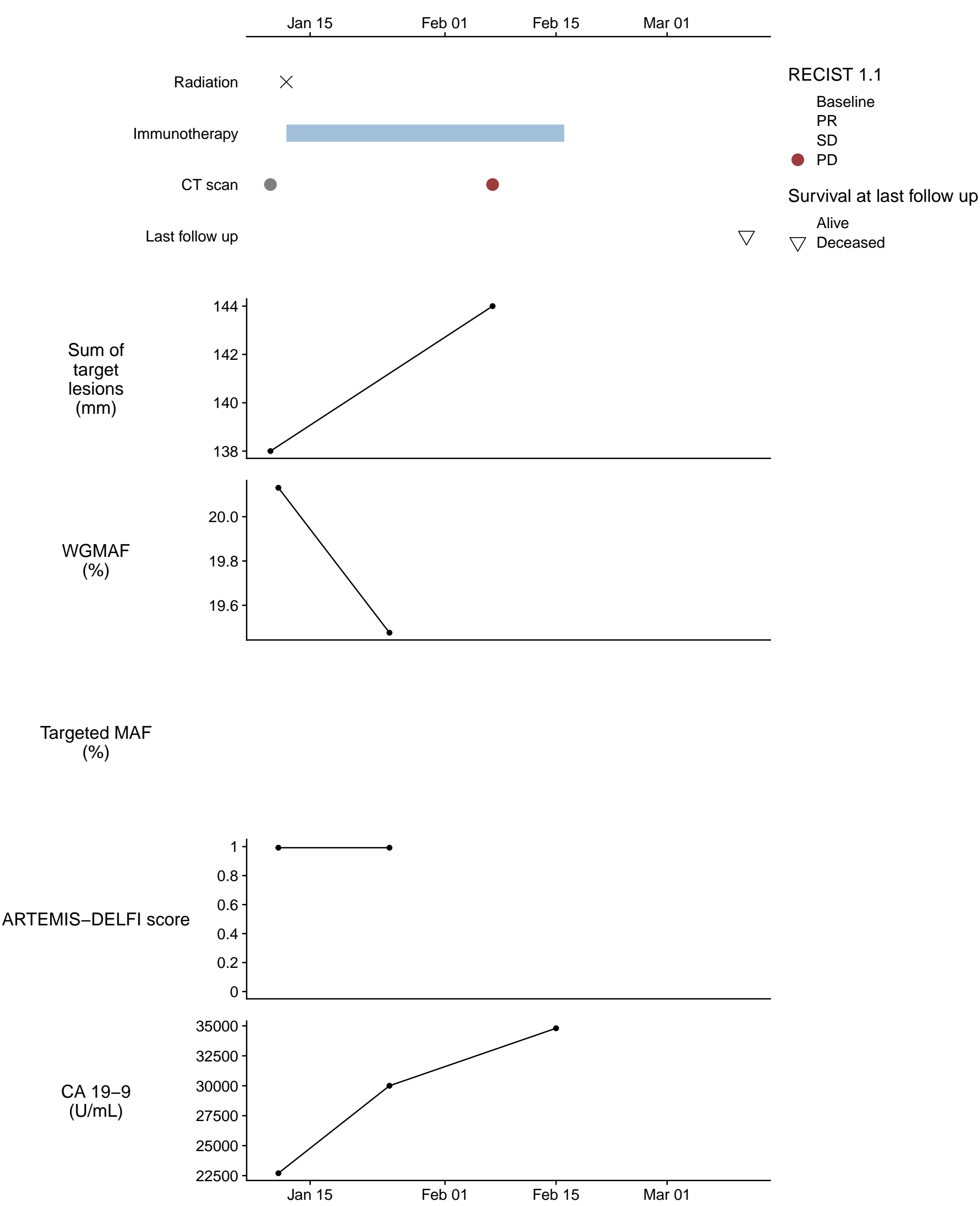
CGPLPA767
WGS data: Tumor (Yes), Normal (Yes), Plasma (7 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 1367, 423



CGPLPA768
WGS data: Tumor (Yes), Normal (Yes), Plasma (7 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 2067, 856



CGPLPA769
WGS data: Tumor (Yes), Normal (Yes), Plasma (2 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 10597, 2972



CGPLPA221
WGS data: Tumor (Yes), Normal (Yes), Plasma (3 timepoints)
Number of SBS mutations by WGS in tumor tissue (unfiltered, filtered): 4798, 4038

