

TRAILS_TR067_I260038_BULK_CAPTURE_PCF_V2_HG19_LB1_411_HW57JDMXX_1, n: 0.5, p: 2, log likelihood: 22260

Tumor Fraction: 0.4762, Ploidy: 2.2

Subclone Fraction: 0.999, Frac. Genome Subclonal: 0.06, Frac. CNA Subclonal: 0.10

