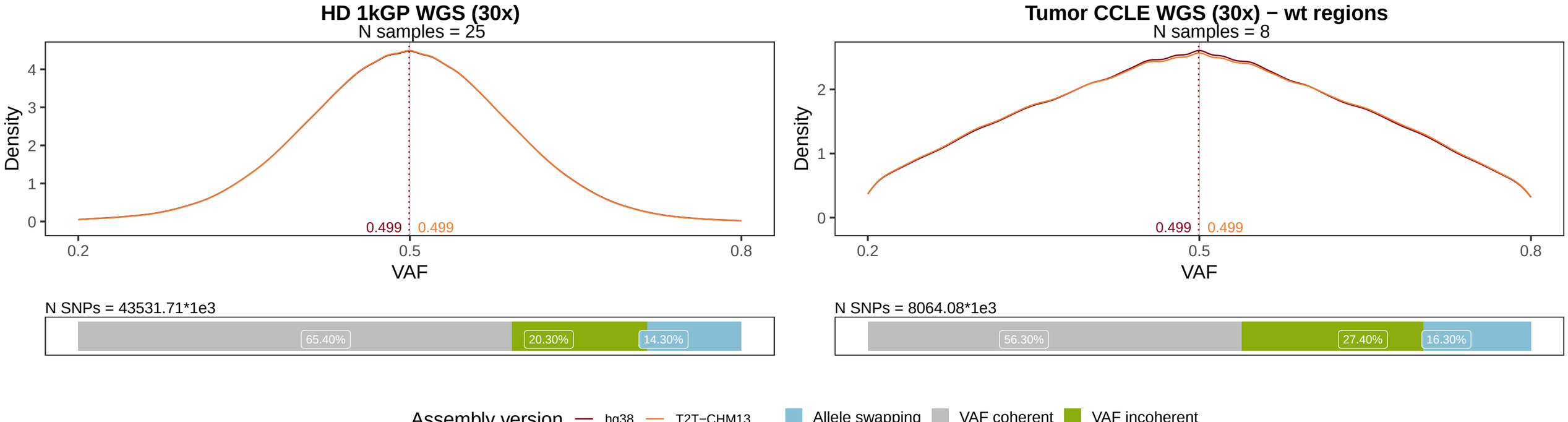
HD 1kGP WGS (30x)
N samples = 25
Density
VAF
0.499
0.499
N SNPs = 43531.71*1e3
65.40%
20.30%
14.30%
Tumor CCLE WGS (30x) − wt regions
N samples = 8
Density
VAF
0.499
0.499
N SNPs = 8064.08*1e3
56.30%
27.40%
16.30%
Assembly version
hg38
T2T−CHM13
Allele swapping
VAF coherent
VAF incoherent