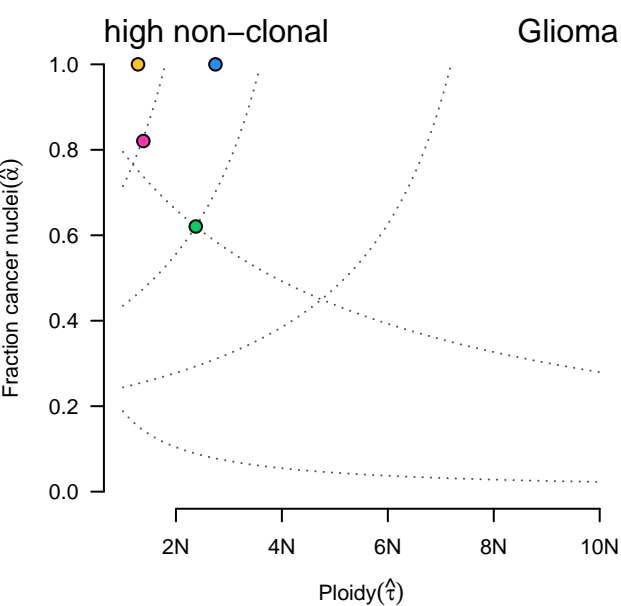
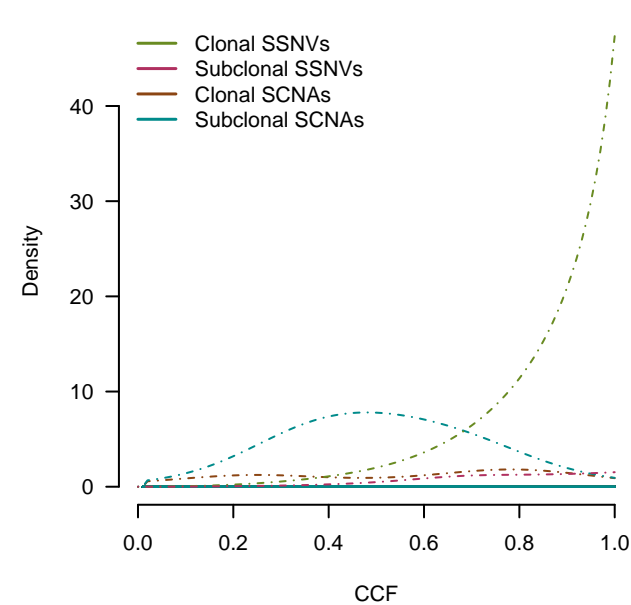
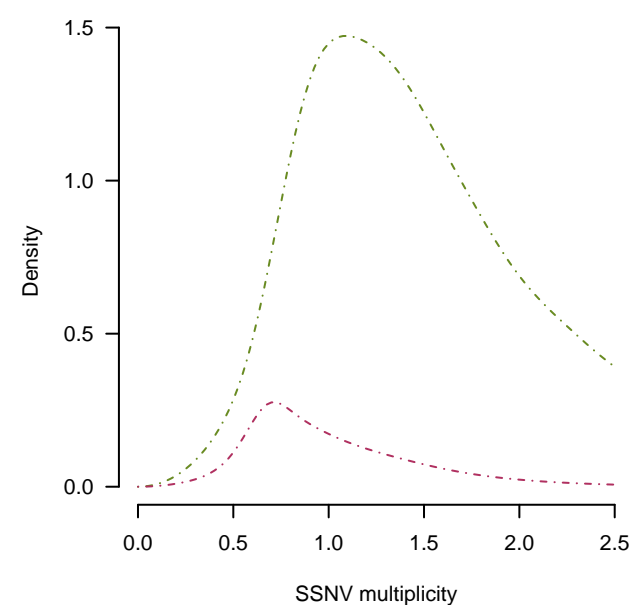
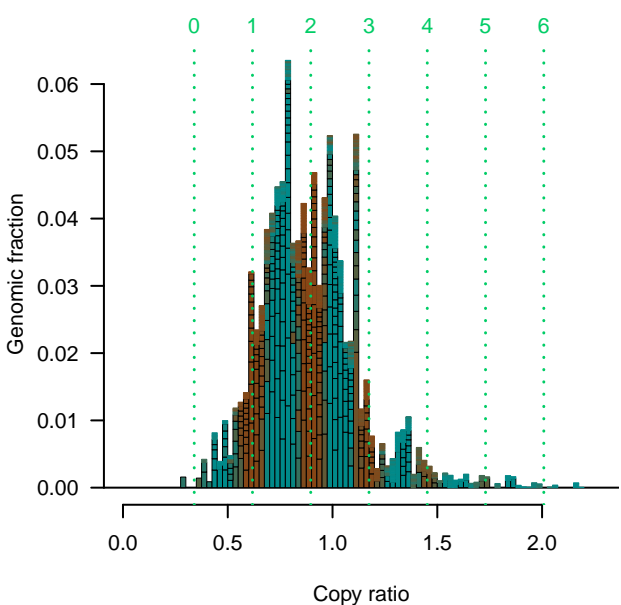


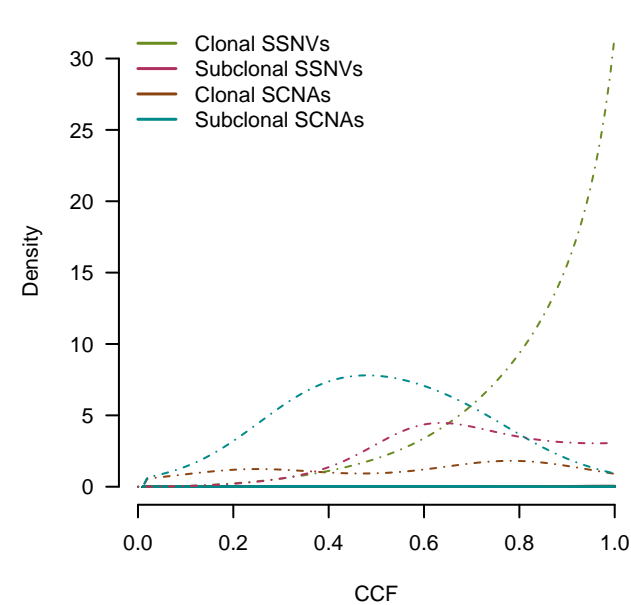
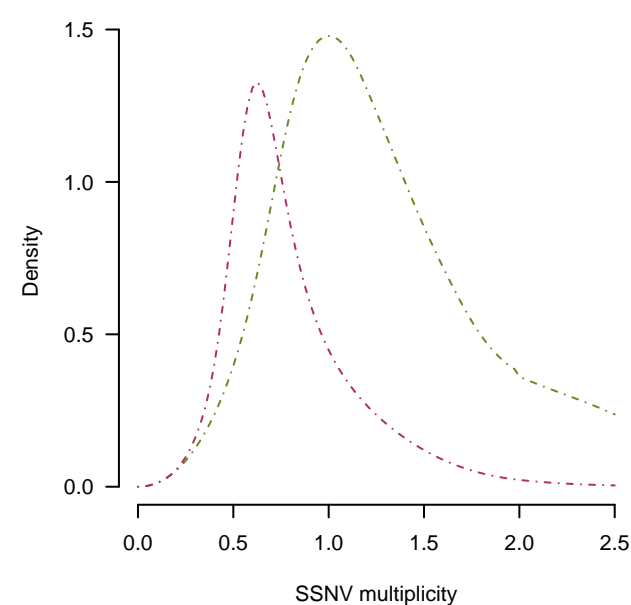
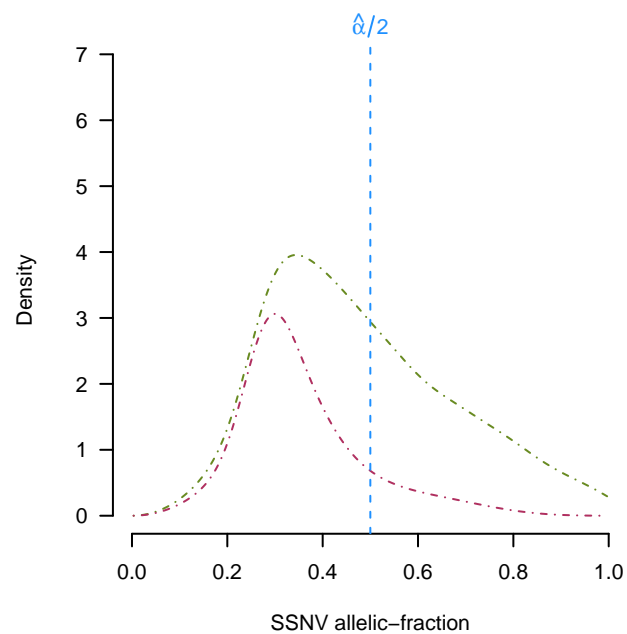
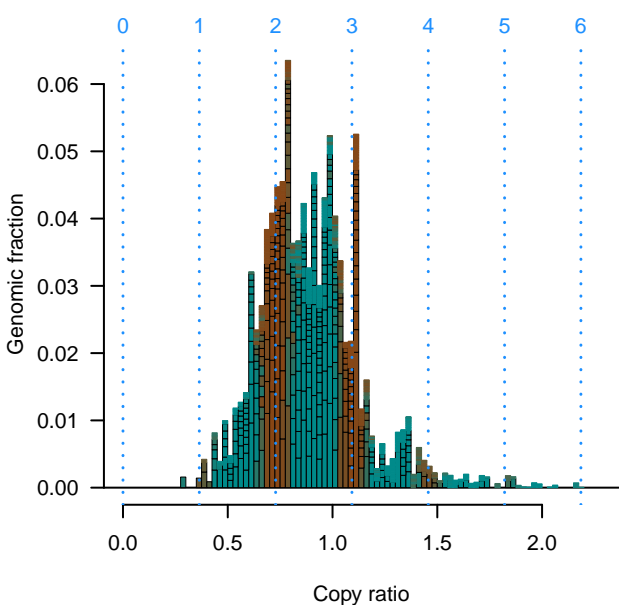
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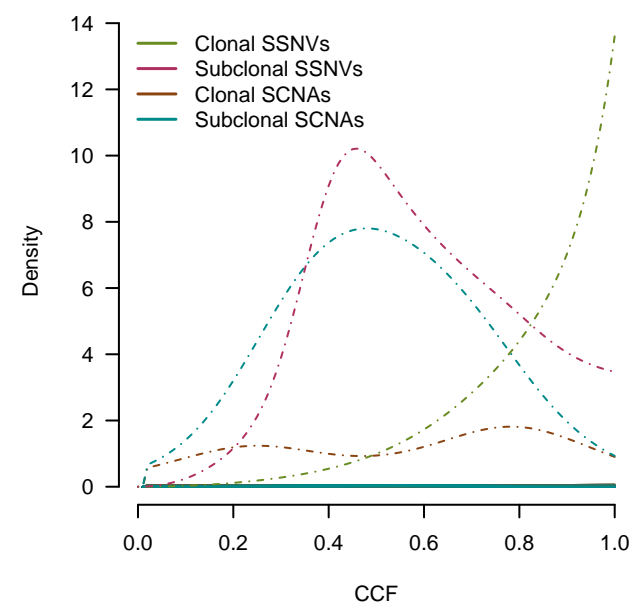
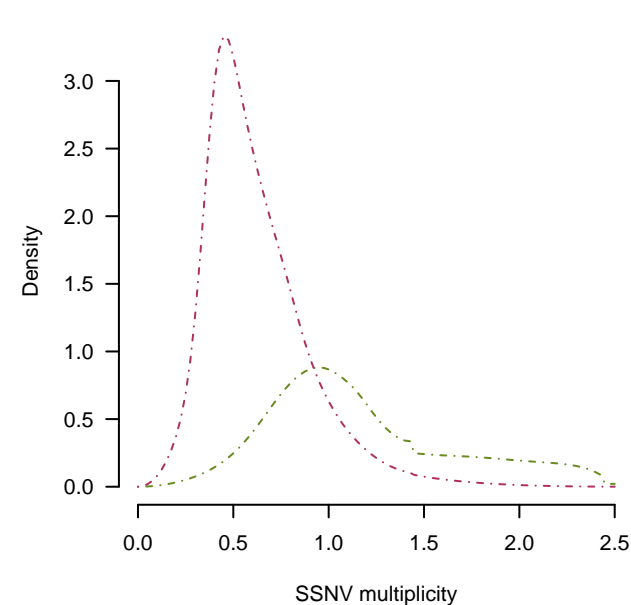
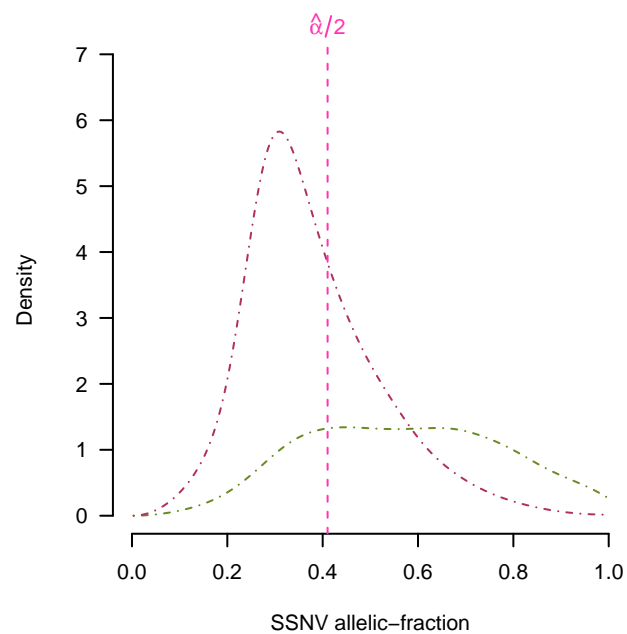
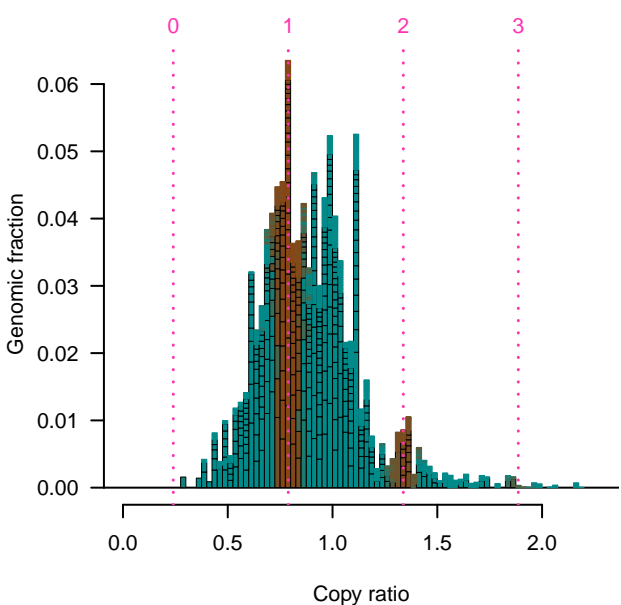
purity($\hat{\alpha}$) = 0.62, ploidy: $\hat{\tau}$ = 2.37, $\hat{\tau}_g$ = 2.02
 $\hat{\sigma}_H$ = 0.02, $\hat{\theta}_Z$ = 0.59, %Het = 58.51
 SCNAs = -1446.63, Kar = -51.74, SSNVs = 161.79



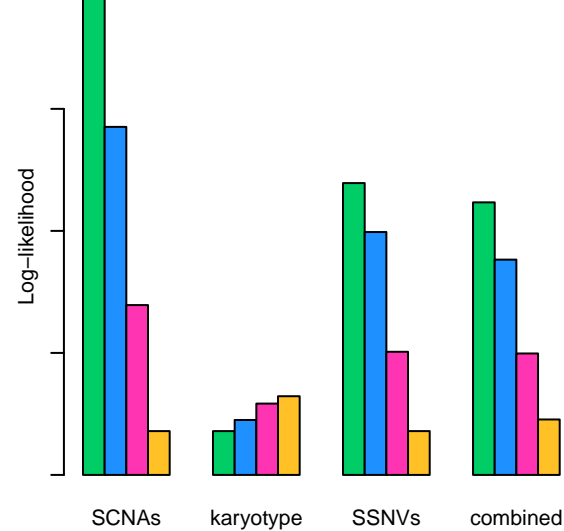
purity($\hat{\alpha}$) = 1, ploidy: $\hat{\tau}$ = 2.75, $\hat{\tau}_g$ = 2.5
 $\hat{\sigma}_H$ = 0.02, $\hat{\theta}_Z$ = 0.6, %Het = 60.48
 SCNAs = -1556.44, Kar = -42.64, SSNVs = 121.66



purity($\hat{\alpha}$) = 0.82, ploidy: $\hat{\tau}$ = 1.38, $\hat{\tau}_g$ = 1.19
 $\hat{\sigma}_H$ = 0.02, $\hat{\theta}_Z$ = 0.73, %Het = 73.35
 SCNAs = -1702.46, Kar = -29.19, SSNVs = 23.5



Model-based evaluation



purity($\hat{\alpha}$) = 1, ploidy: $\hat{\tau} = 1.28$, $\hat{\tau}_g = 1.1$
 $\hat{\sigma}_H = 0.02$, $\hat{\theta}_2 = 0.76$, %Het = 76.29
SCNAs = -1805.76, Kar = -23.15, SSNVs = -41.48

