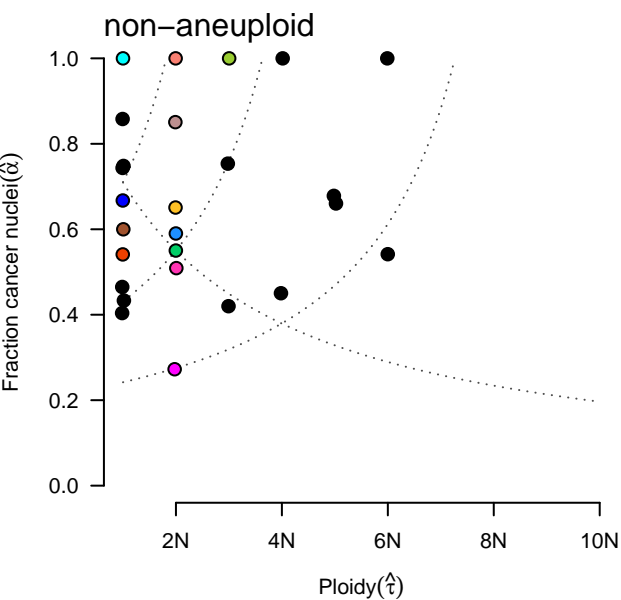
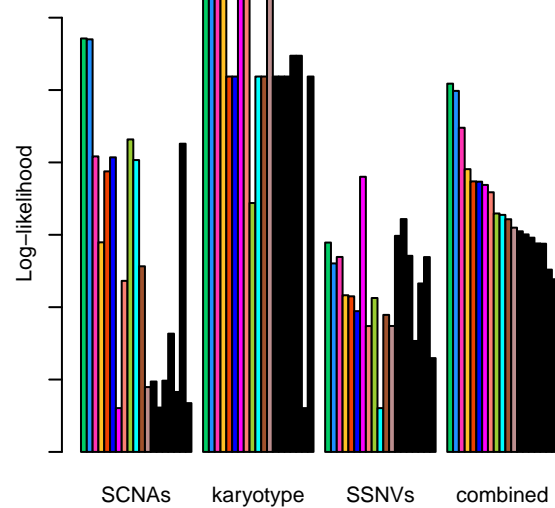


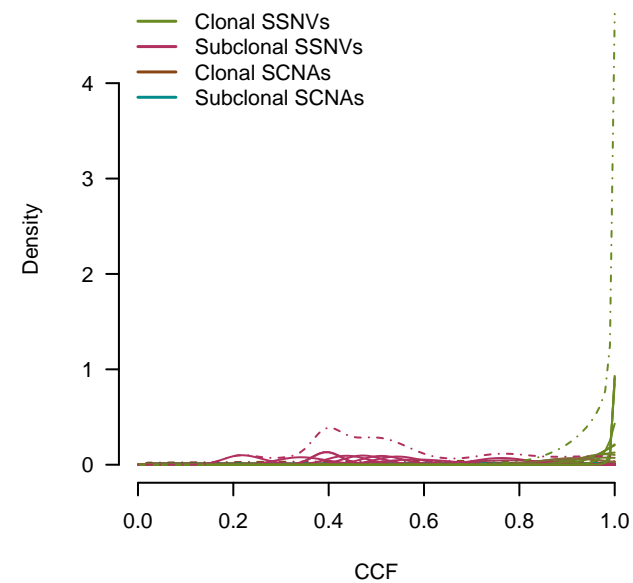
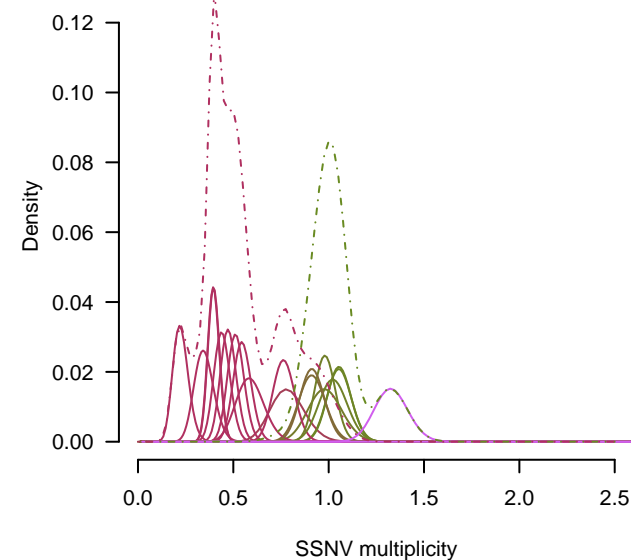
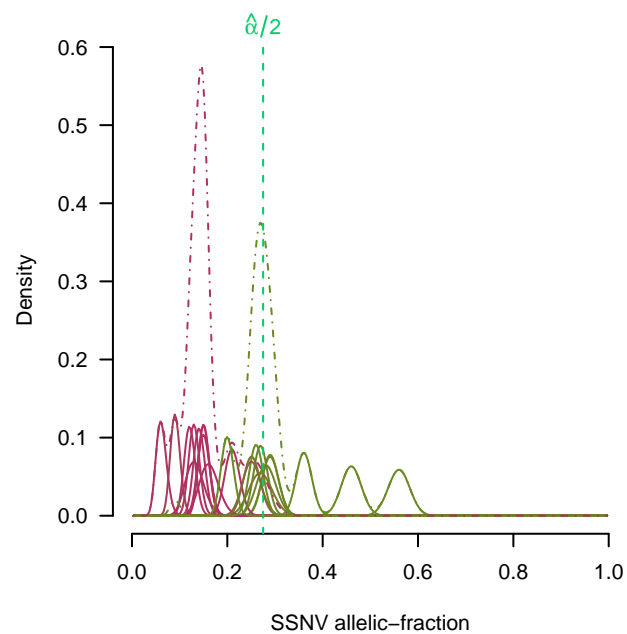
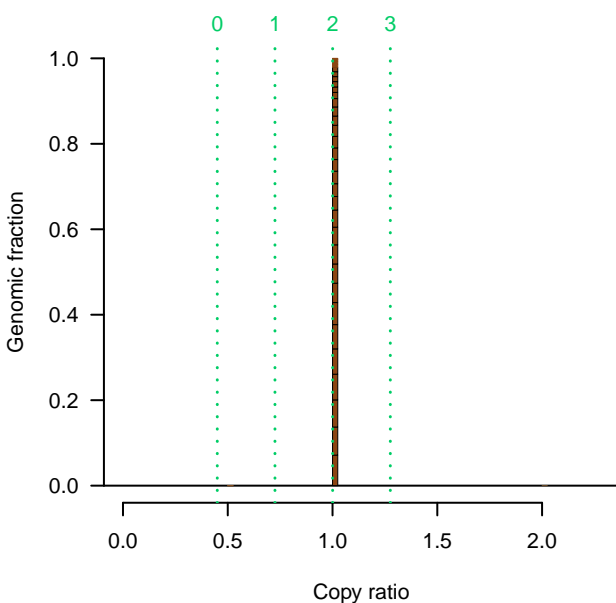
SIM\_DATA\_3.3



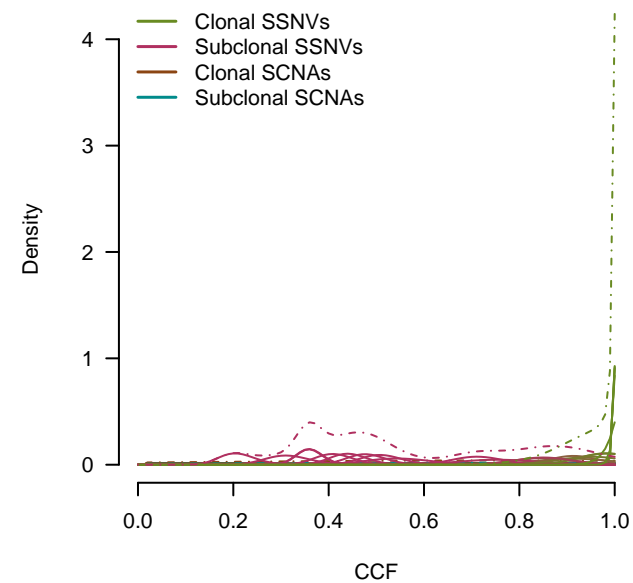
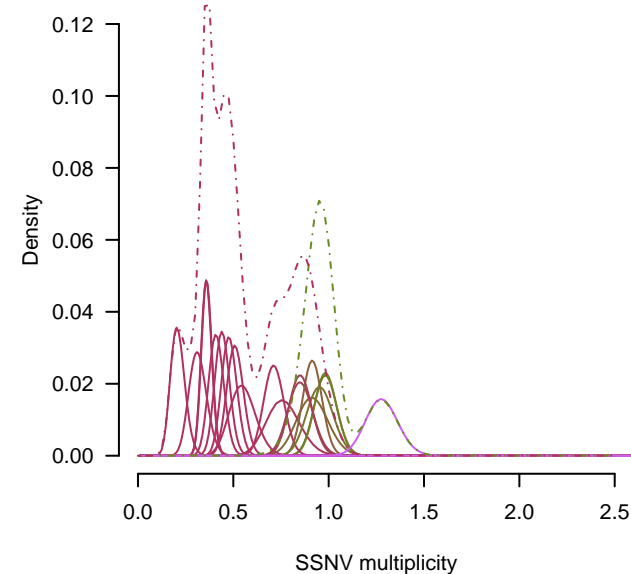
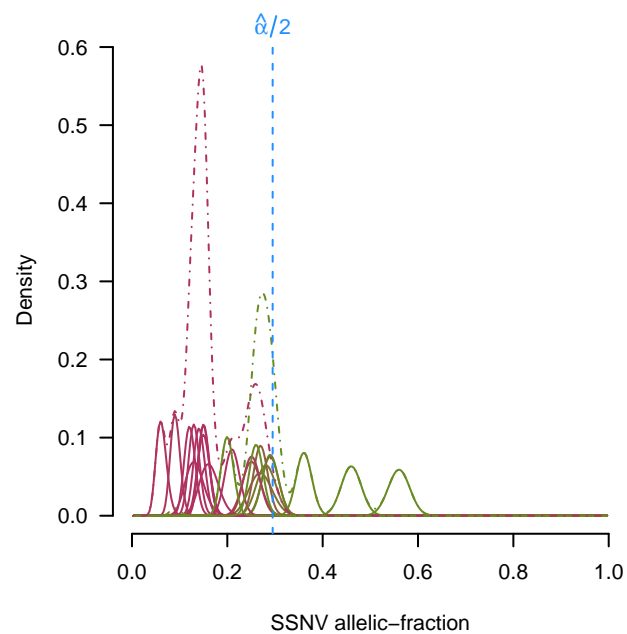
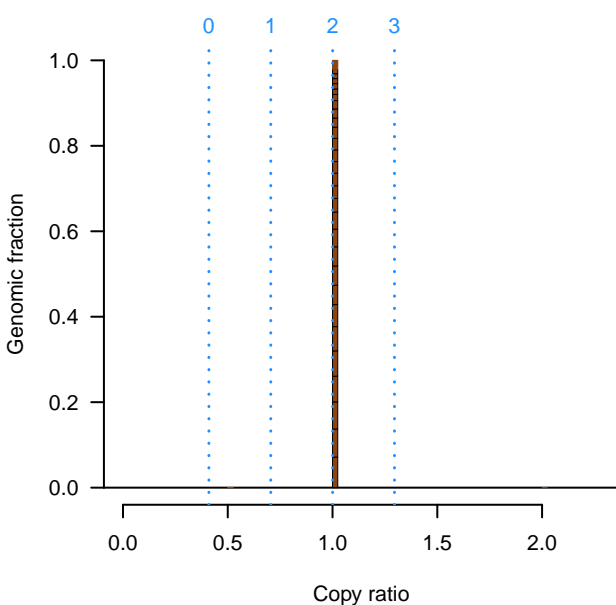
Model-based evaluation



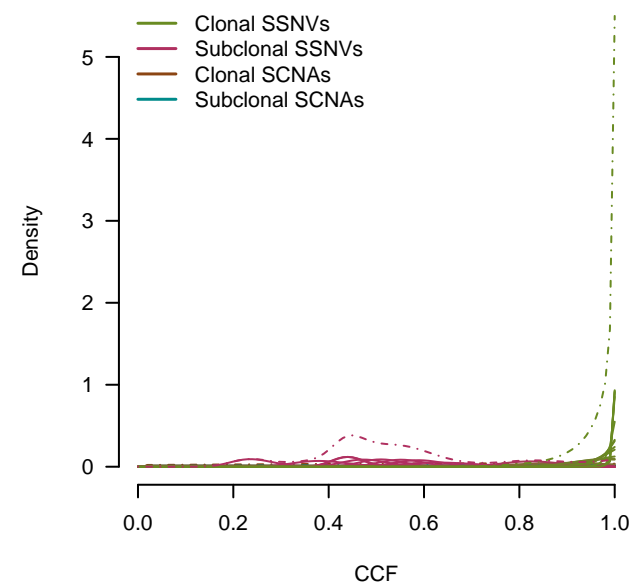
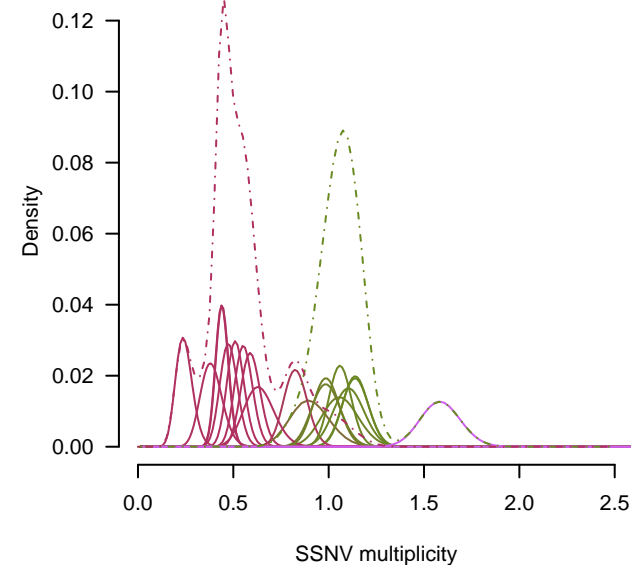
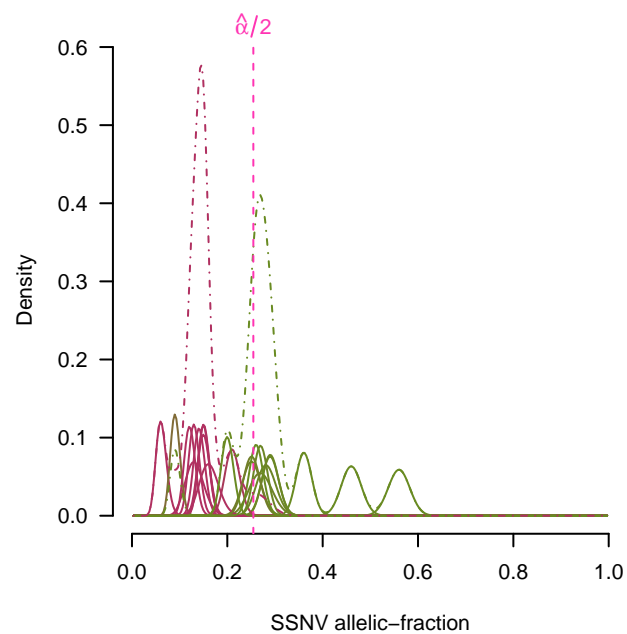
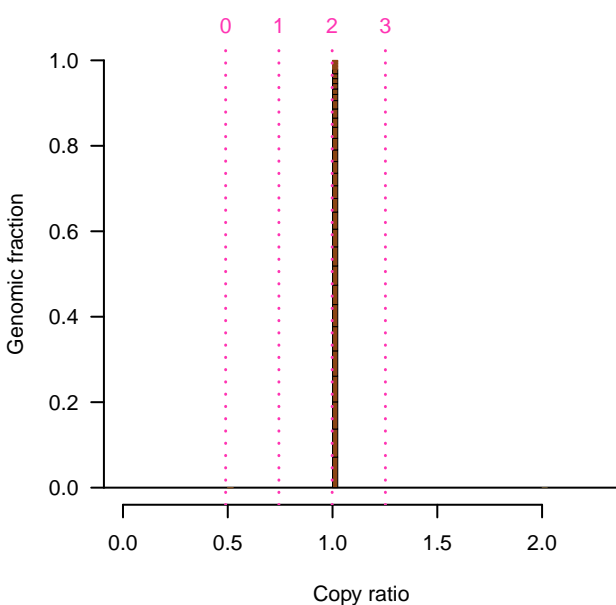
purity( $\hat{\alpha}$ ) = 0.55, ploidy:  $\hat{\tau} = 2$ ,  $\hat{\tau}_g = 2$   
 $\hat{\sigma}_H = 0.001$ ,  $\hat{\theta}_Z = 0$ , %Het = 0  
 SCNAs = 153.08, Kar = -4.62, SSNVs = 66.25

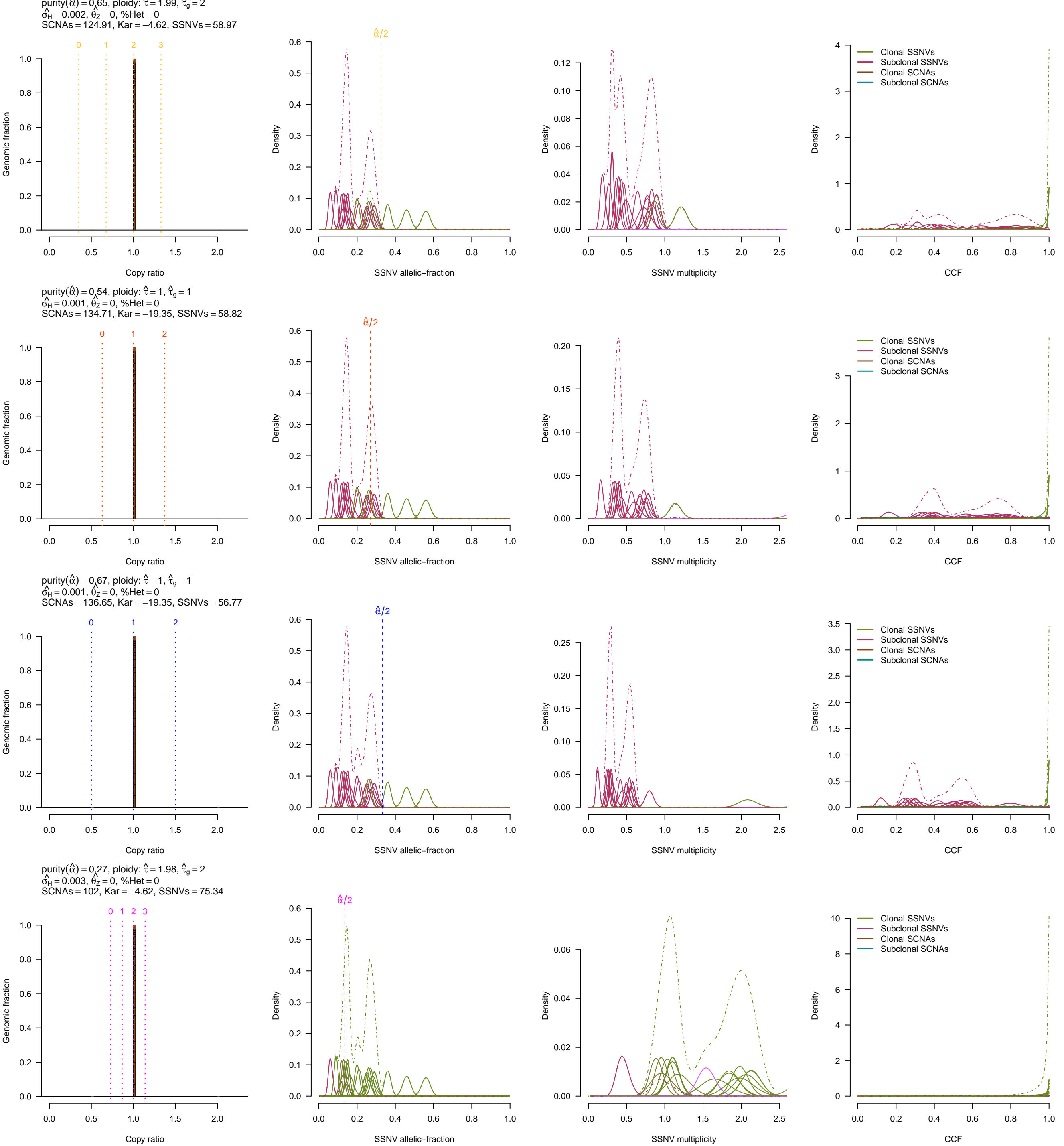


purity( $\hat{\alpha}$ ) = 0.59, ploidy:  $\hat{\tau} = 2$ ,  $\hat{\tau}_g = 2$   
 $\hat{\sigma}_H = 0.001$ ,  $\hat{\theta}_Z = 0$ , %Het = 0  
 SCNAs = 152.95, Kar = -4.62, SSNVs = 63.36

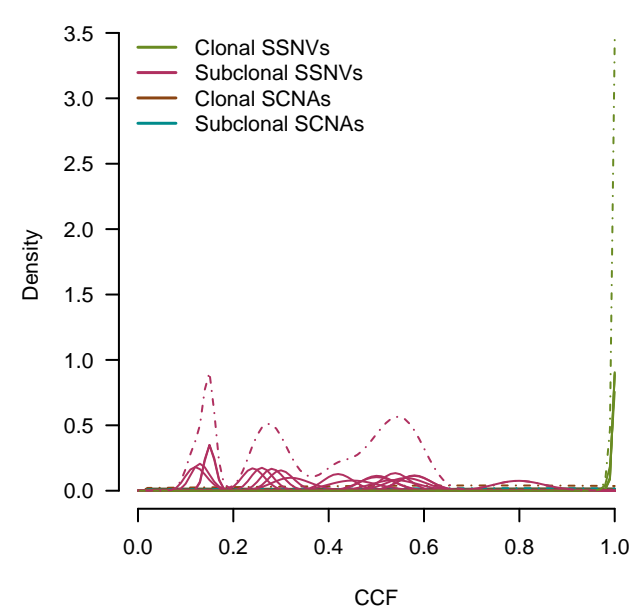
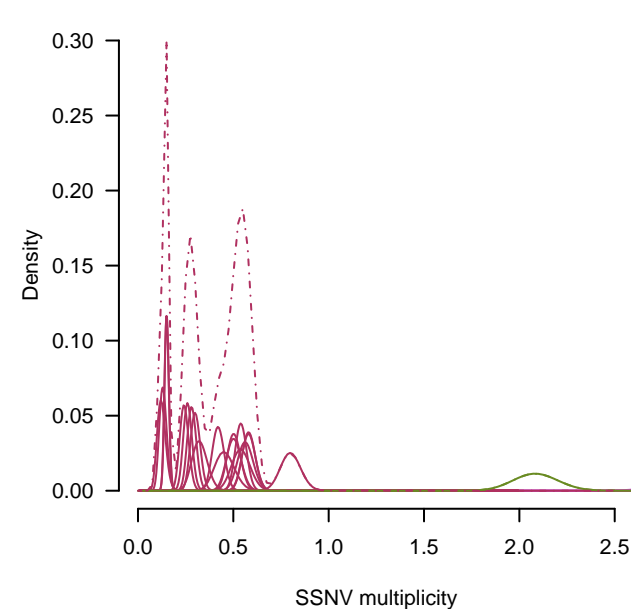
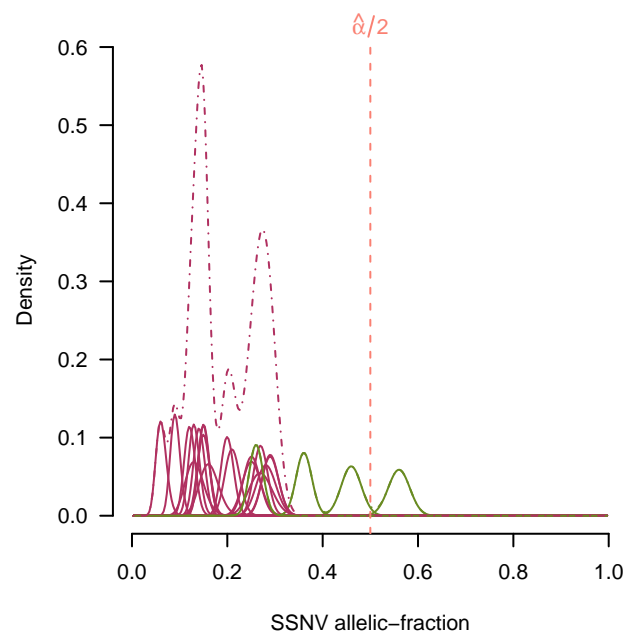
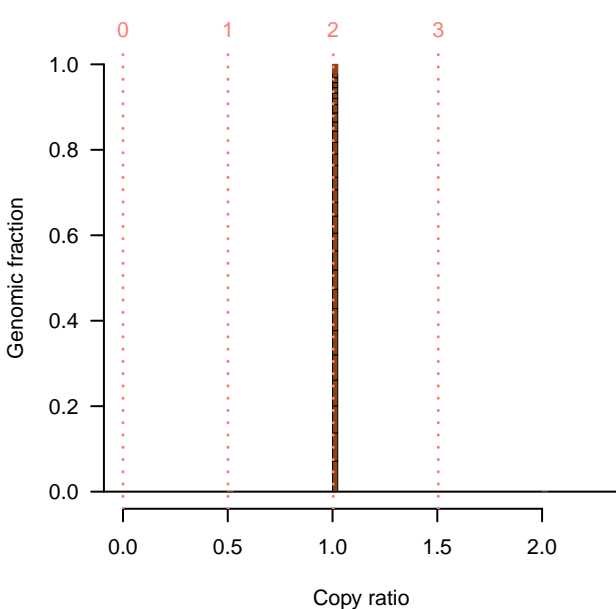


purity( $\hat{\alpha}$ ) = 0.51, ploidy:  $\hat{\tau} = 2.01$ ,  $\hat{\tau}_g = 2$   
 $\hat{\sigma}_H = 0.001$ ,  $\hat{\theta}_Z = 0$ , %Het = 0  
 SCNAs = 136.78, Kar = -4.62, SSNVs = 64.26

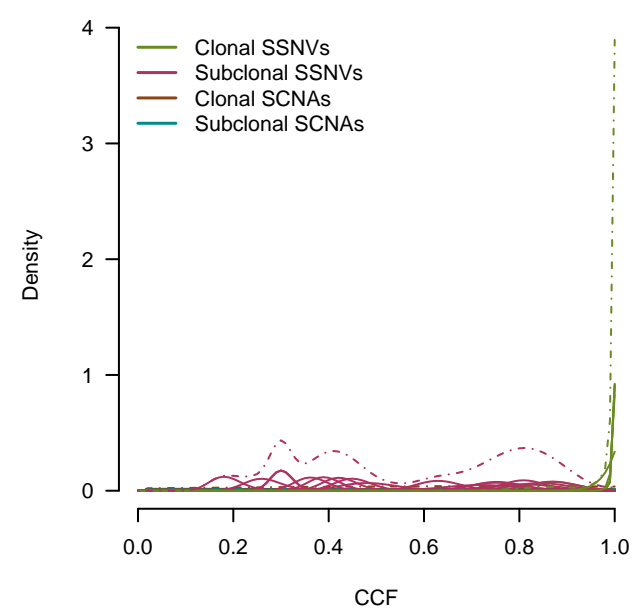
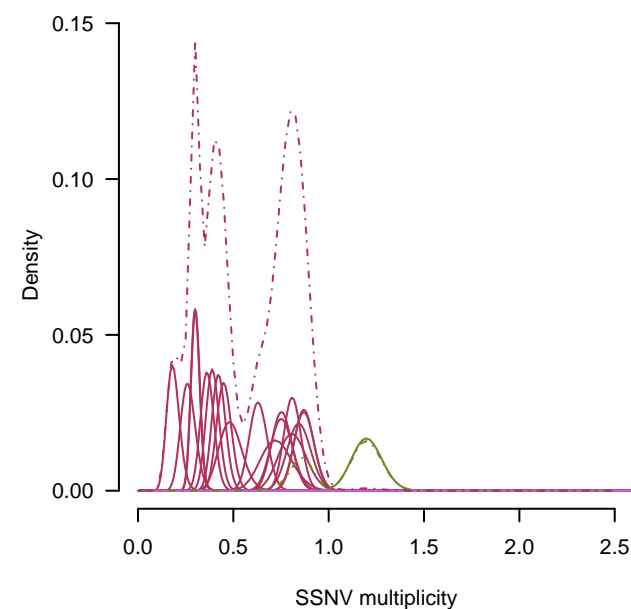
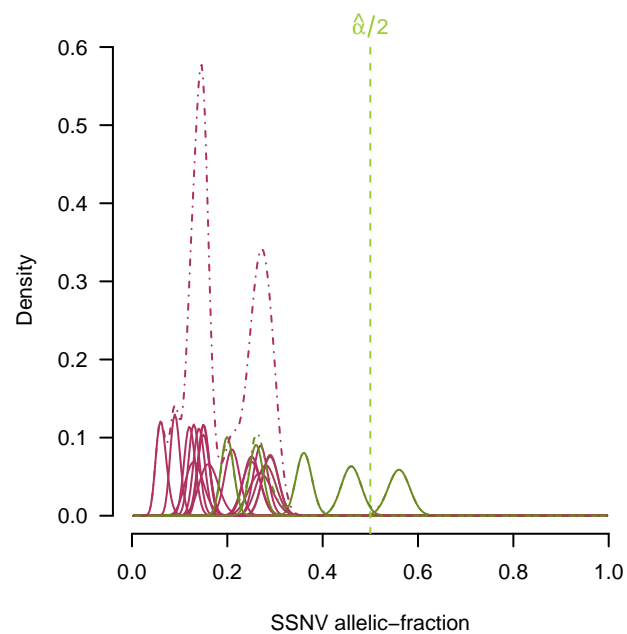
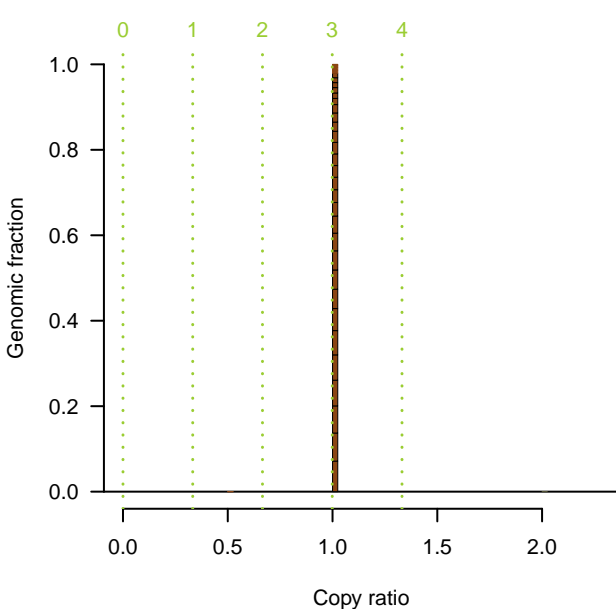




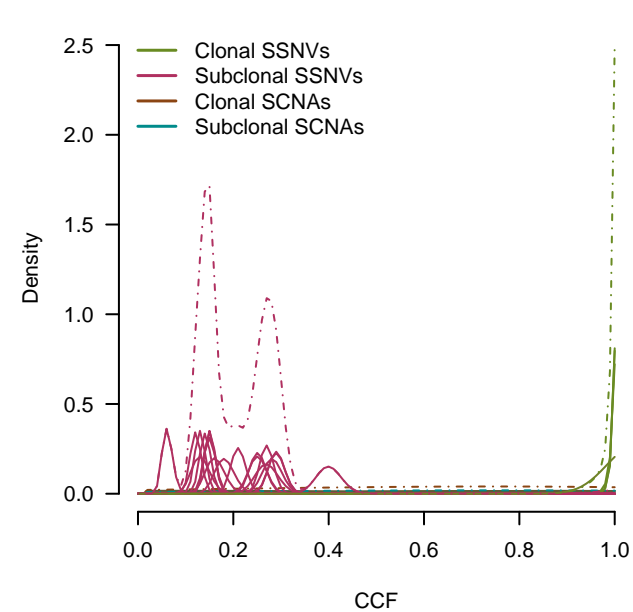
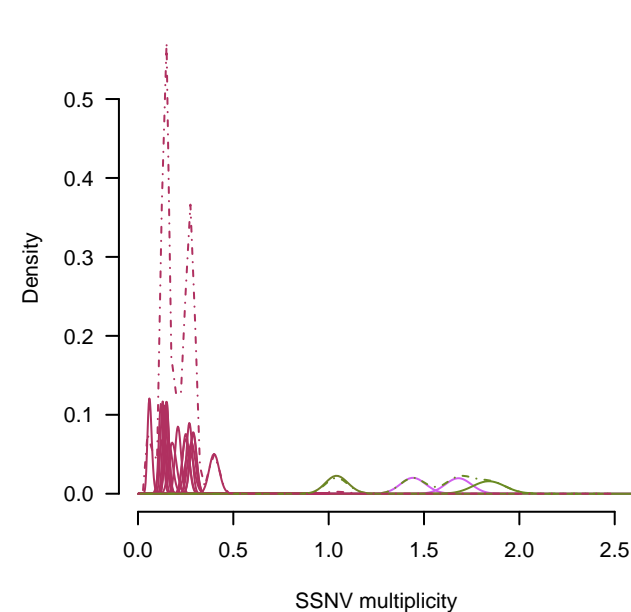
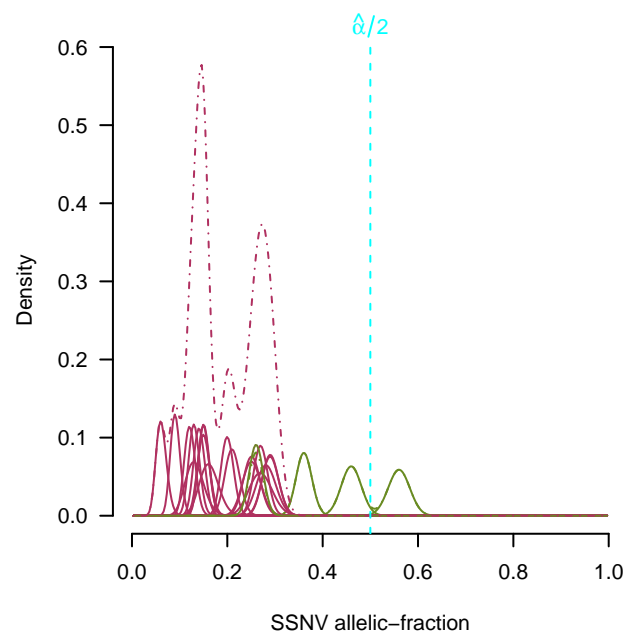
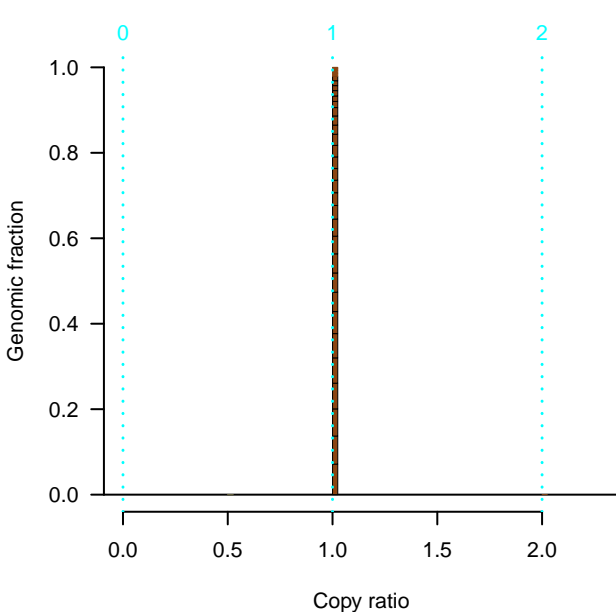
purity( $\alpha$ ) = 1, ploidy:  $\hat{\tau} = 1.99$ ,  $\hat{\tau}_g = 2$   
 $\hat{\sigma}_H = 0.003$ ,  $\hat{\theta}_Z = 0$ , %Het = 0  
 SCNAs = 119.59, Kar = -4.62, SSNVs = 54.71



$\text{purity}(\hat{\alpha}) = 1$ , ploidy:  $\hat{\tau} = 3$ ,  $\hat{\tau}_g = 3$   
 $\hat{\sigma}_H = 0.001$ ,  $\hat{\theta}_Z = 0$ , %Het = 0  
 SCNAs = 139.13, Kar = -36.83, SSNVs = 58.59



$\text{purity}(\hat{\alpha}) = 1$ ,  $\text{ploidy: } \hat{\tau} = 1, \hat{\tau}_g = 1$   
 $\hat{\sigma}_H = 0.001, \hat{\theta}_Z = 0, \% \text{Het} = 0$   
 $\text{SCNAs} = 136.28, \text{Kar} = -19.35, \text{SSNVs} = 43.37$



purity( $\hat{\alpha}$ ) = 0.6, ploidy:  $\hat{\tau} = 1.01$ ,  $\hat{\tau}_g = 1$   
 $\hat{\sigma}_H = 0.002$ ,  $\hat{\theta}_Z = 0$ , %Het = 0  
 SCNAs = 121.59, Kar = -19.35, SSNVs = 56.25

