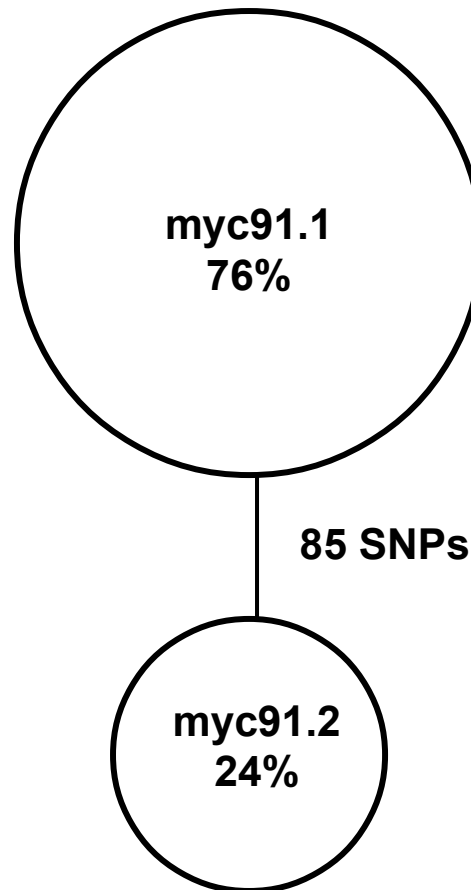
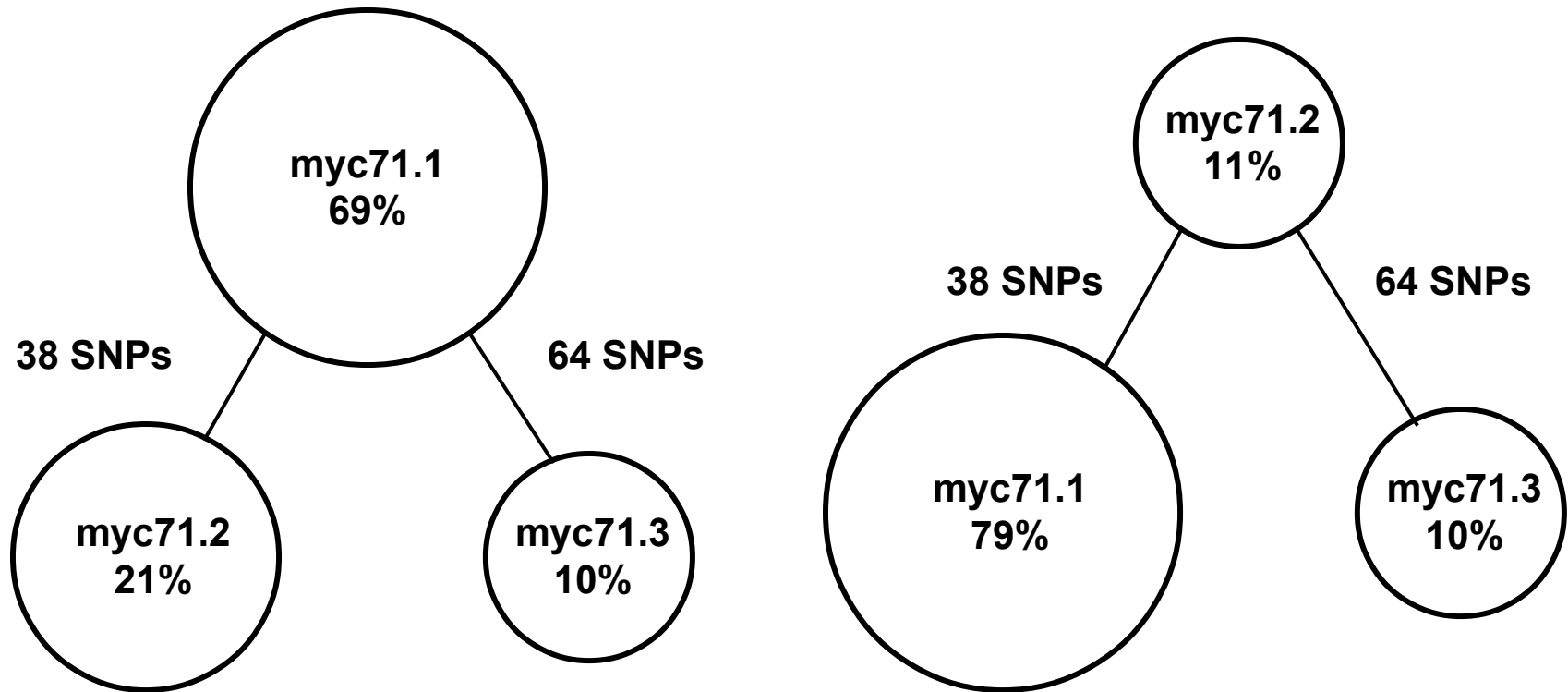


*The distribution of the minor allele frequency for myc91 can be explained by two genotypes (or strains) in the sample, with the minor one comprising 24% of the cells*

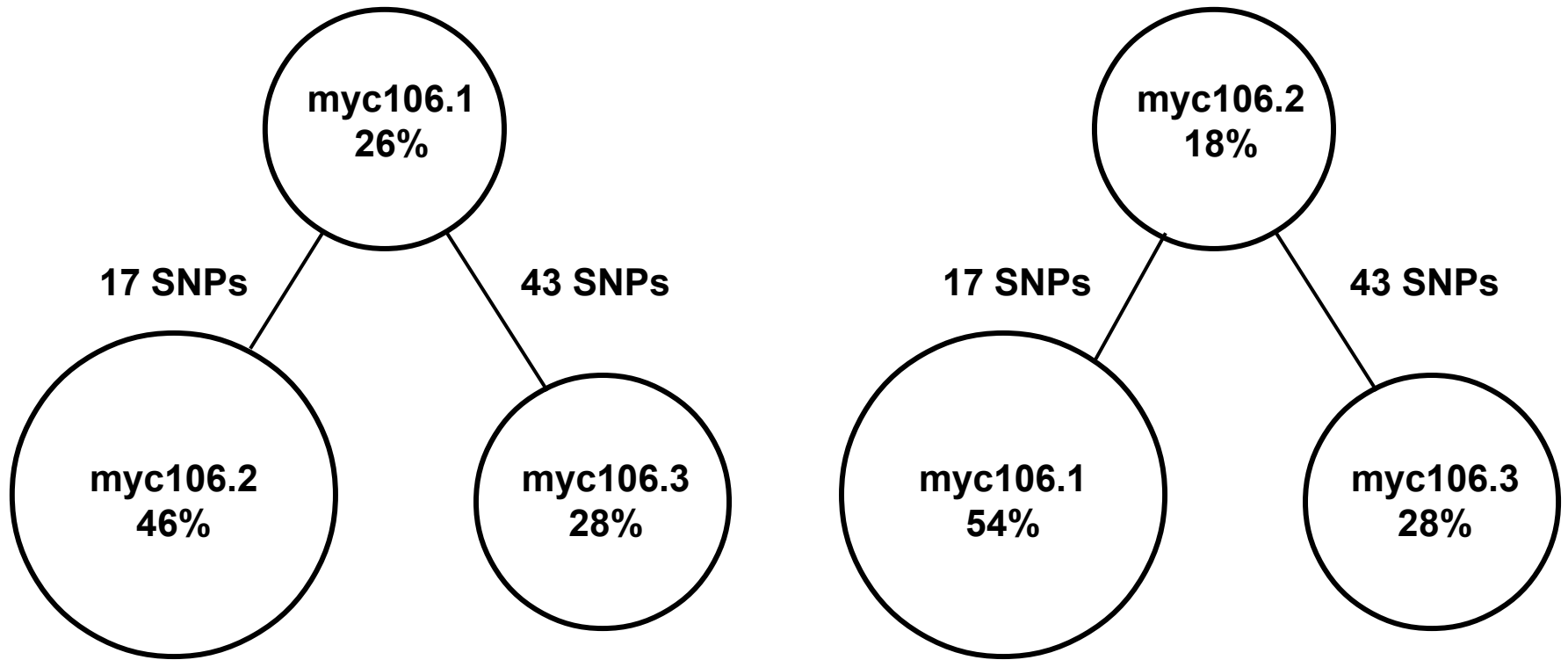


*The distribution of the minor allele frequency for myc71 can be explained by three genotypes in the sample; of the two possible trees, the one at the right seems more likely and corresponds to the tree in Fig 2*



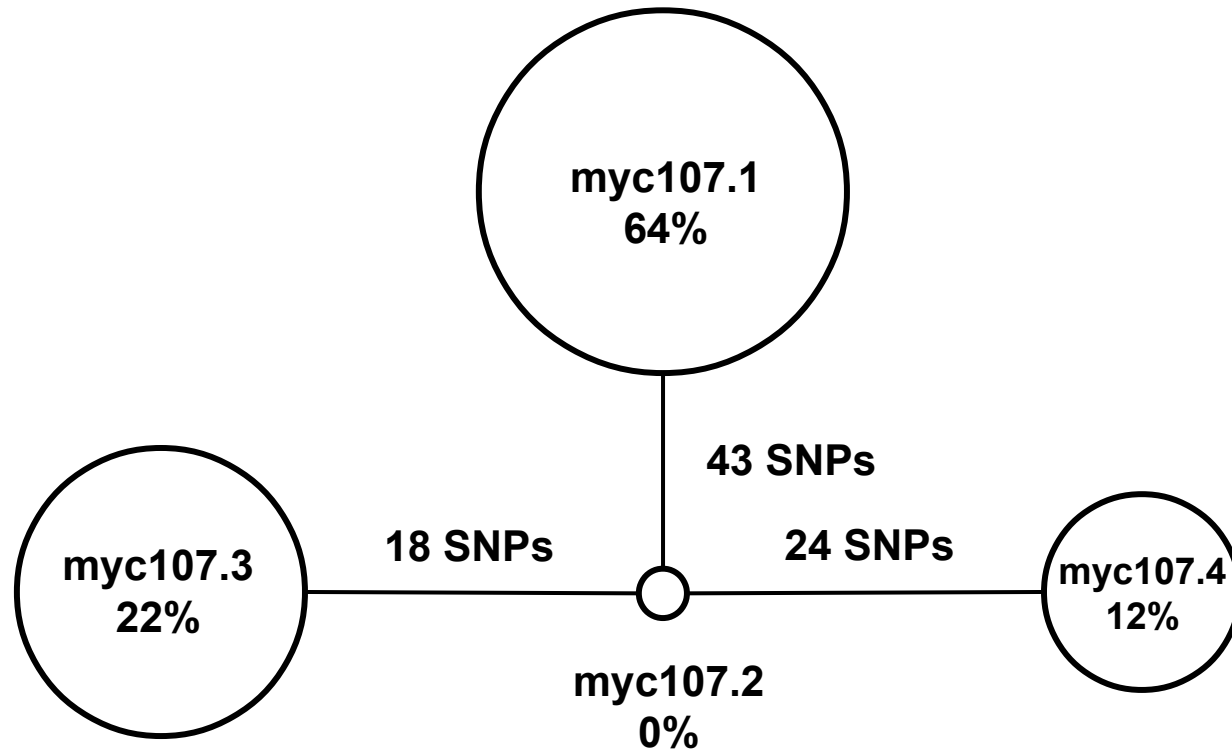
**Tree consistent  
with Fig 2**

*The distribution of the minor allele frequency for myc106 can be explained by three genotypes in the sample; of the two possible trees, the one at the left seems more likely and corresponds to the tree in Fig 2*



**Tree consistent  
with Fig 2**

*The distribution of the minor allele frequency for myc107 can be explained by either three or four genotypes in the sample; the three-genotype tree shown here seems more likely and corresponds to the tree in Fig 2; myc107.2 is an artificial genotype*



Tree consistent  
with Fig 2