Figures S4.1-S4.4 STRUCTURE bar plots using SSR or SNP loci of *Armillaria cepistipes* genotypes in four subpopulations. Each bar represents the average estimated individual membership probability (ordinate) of an individual to belong to a specific cluster (indicated by specific color). Bar plots are supported by a scatterplot (below) with mean log-likelihood values (± standard deviation) for different numbers of clusters (K)

Figure S4.1 Loci: 17 SSRs

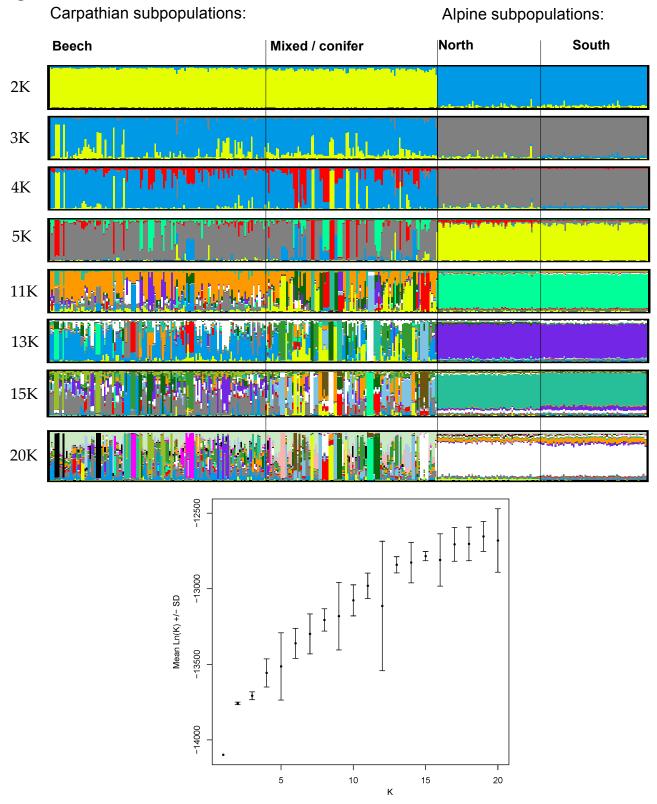


Figure S4.2 Loci: 24 SNPs

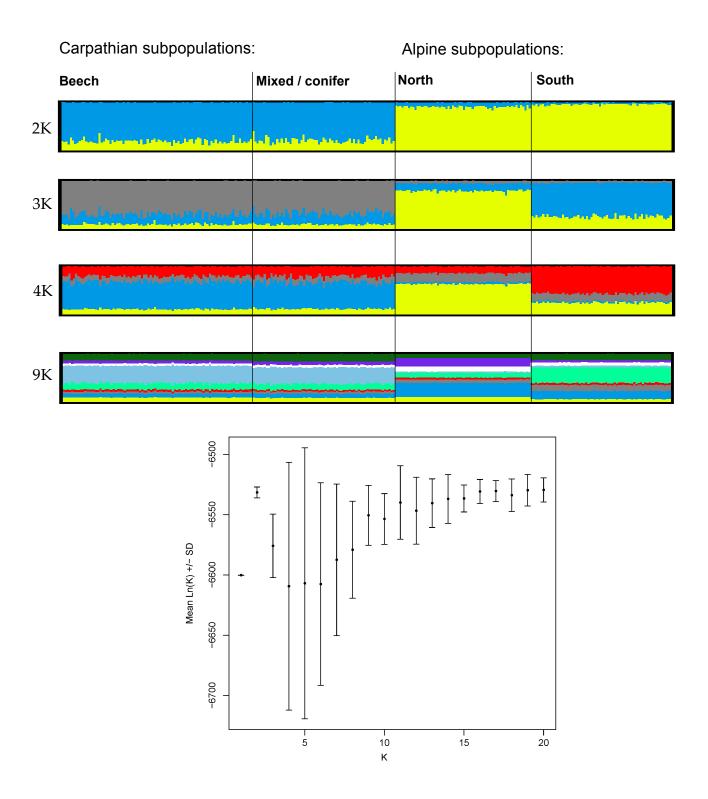


Figure S4.3 Loci: 19 SNPs (excluded 5 loci presumably under natural selection pressure)

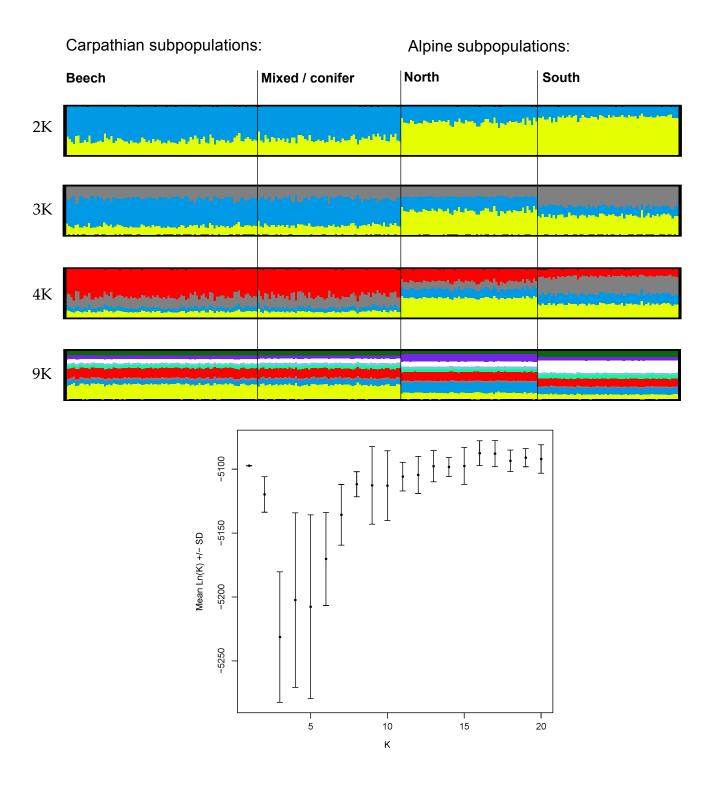


Figure S4.4 Loci: 24 SNP loci

Alpine subpopulations:

