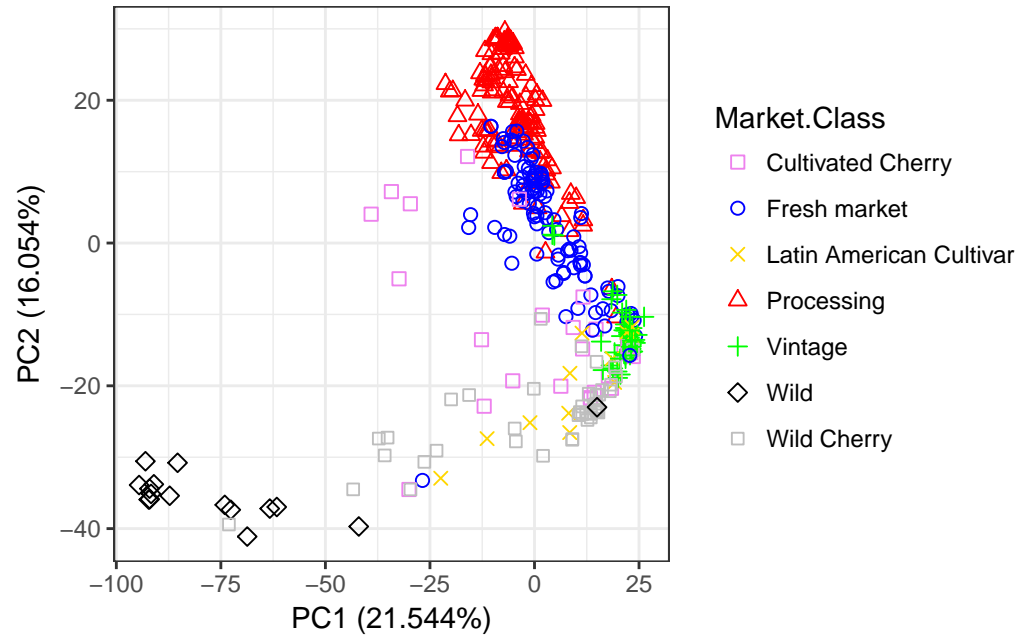Figure 1. A
PCA based on 4,393 SNP markers: All Sub−populations

Figure 1. B
PCA based on 4,393 SNP markers: 3 large fruited sub−populations