Figure 1. A
Principal component analysis (PCA) based on 4,393 SNP markers: All Sub–populations

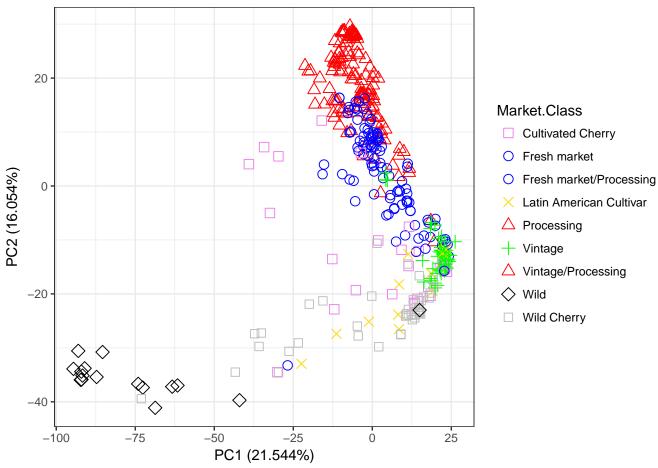
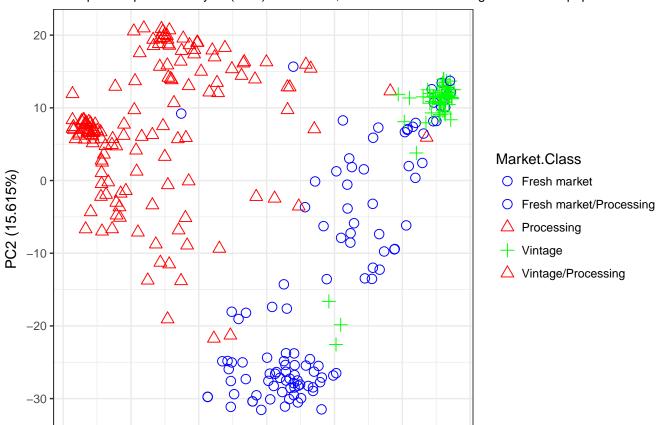


Figure 1. B
Principal component analysis (PCA) based on 4,393 SNP markers: 3 large fruited sub–populations



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-10

PC1 (21.85%)

-30

-20