

PCA plot of 1000 random SNPs showing genetic differentiation between three populations: NC (blue), NW (green), and CAC (red). The x-axis is PC1 (15.1%) and the y-axis is PC2 (10.2%). Points are labeled with accession numbers and coordinates. A color scale at the bottom indicates the proportion of each population's genetic contribution to each individual, ranging from 0.0 (blue) to 1.0 (red).

● NC_071985.2	● NC_072001.2
● NC_071988.2	● NC_072002.2
● NC_071990.2	● NC_072003.2
● NC_071991.2	● NC_072004.2
● NC_071992.2	● NC_072005.2
● NC_071994.2	● NC_072007.2
● NC_071995.2	● NC_072008.2
● NC_071996.2	● NC_072009.2
● NC_071997.2	● NC_085928.1
● NC_071998.2	● NC_085929.1
● NC_071999.2	● NW_026946739.1
● NC_072000.2	● NW_026946741.1