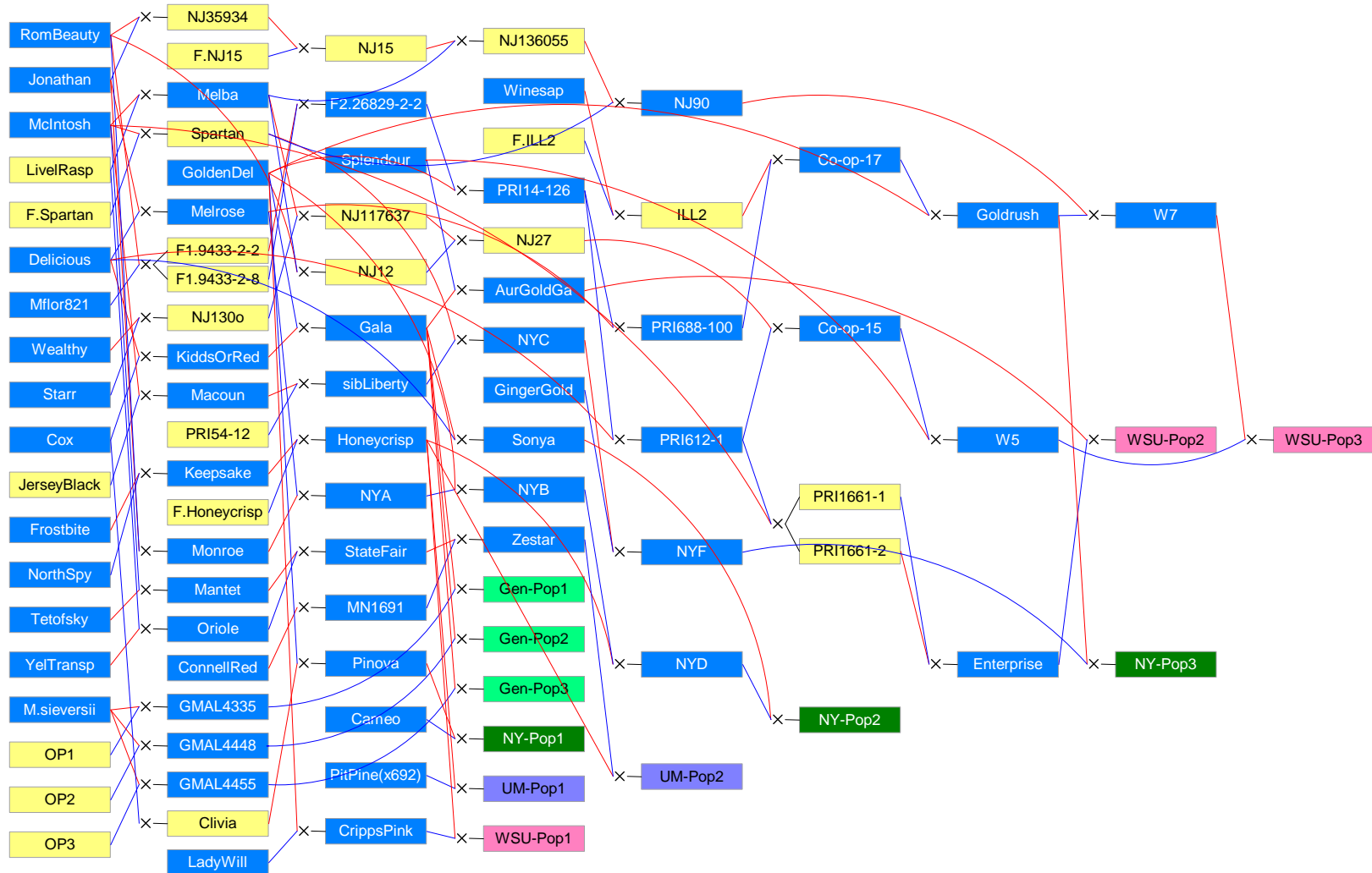
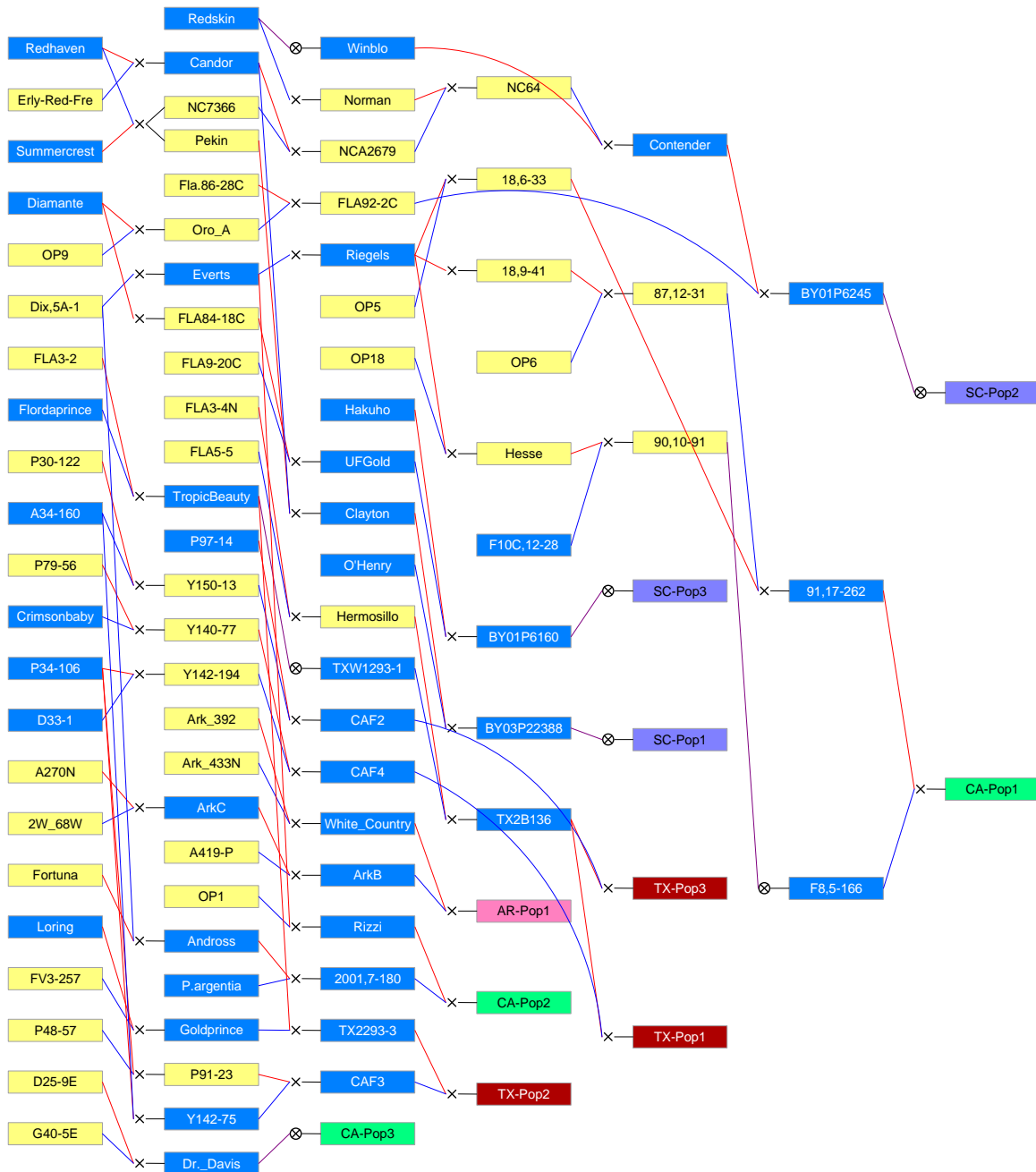


PediMap image of *Malus* germplasm representing U.S. breeding programs from which RosBREED Crop Reference Sets (CRS) and Breeding Pedigree Sets (BPS) will be selected for SNP genome scans. Individuals tentatively included in the CRS are colored in blue, and CRS populations (“Pop”) located at WSU, NY-Cornell, NY-USDA/ARS, and UM, are color coded in pink, dark green, light green and purple, respectively. Those parents/ancestors/founders for which DNA or phenotypes will probably not be available are colored in yellow.



PediMap image of *P. perisca* germplasm representing U.S. breeding programs from which RosBREED Crop Reference Sets (CRS) and Breeding Pedigree Sets (BPS) will be selected for SNP genome scans. Individuals tentatively included in the CRS are colored in blue, and CRS populations (“Pop”) located at AR, CA, SC, and TX are color coded in pink, green, purple and red, respectively. Those parents/ancestors/founders for which DNA or phenotypes will probably not be available are colored in yellow. For ease in visualization, this image just represents the latest generations in the CRS ancestry, the maximum generation number in the peach ancestry is 11.







PediMap image of *Fragaria* germplasm representing U.S. breeding programs from which RosBREED Crop Reference Sets (CRS) and Breeding Pedigree Sets (BPS) will be selected for SNP genome scans. Individuals tentatively included in the CRS are colored in blue, and CRS populations (“Pop”) located at Oregon, Michigan and California are color coded in pink and green respectively. Those parents/ancestors/founders for which DNA or phenotypes will probably not be available are colored in yellow. For ease in visualization, this image just represents the latest generations in the CRS ancestry, the maximum generation number in the strawberry ancestry is 12.

