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| --- | --- | --- | --- | --- |
| **Seed Source** | **Mother**  **Selection Type** | **Progeny Test** | **Selection and Family Numbering** | **Progeny Sel’n** |
| Native stands,  Seed acquistions | 1st gen seln | **1st gen progeny test**  Eg., 13-10-30A1, (PTEH-SAF-30A1) | ProjectNum = species (e.g., 13 = TEH)  ProvNum = provenance (e.g., 05 = San Jeronimo)  FamNum = family  Notes:   * Some species have more than one project number * Prov numbers are not unique to a species. * Family numbers are not unique to a species | 2nd gen seln |
| Thinned 1st gen progeny test, grafted CSO | 2nd gen seln | **2nd gen progeny test**  e.g., 13-63-X18A1, (PTEH-YOR-X18A1) | ProjectNum = species  F2Prov SSMMN SS=species code, MM=memcode (Camcore =0), N= number: 1, 2, … Ex 13001, 13021 (SKC only prov)  FamilyNum = family  e.g. 13-037  5/2023: new code (when/if applied) = 13-2037 and/or PTH-2037 | 3rd gen seln |
| Thinned 2nd gen progeny test | 3rd gen seln | **3rd gen progeny test**  e.g., 13-63-Y18A1, (PTEH-YOR-Y18A1) | SpeciesCode = letter code  Seln/Family Num = number code  e.g., PTH-3nnn |  |
|  |  |  |  |  |

* Options
  + PTH3-nnn
  + **PTH-3nnn**
  + PTEH-3nnn
  + PTH-Xnnn (use the X,Y,Z, coding that matches 2nd gen trials)