

Genetic Progress in 53 Years of the Peach Breeding Program of Embrapa: Canning Genotypes

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1 **Title Page**

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4 **genotypes**

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32 **Abstract.** Peach is a traditional crop in the south of Rio Grande do Sul State, Brazil, where 30
33 to 53 million cans of peaches in syrup are produced annually. All the raw material produced in
34 the region consists of fruits originating from the peach breeding program of the Brazilian
35 Agricultural Research Corporation (Embrapa Temperate Agriculture), which started even
36 before Embrapa at the Experimental Station of Pelotas, Ministry of Agriculture. The objective
37 was to estimate the genetic progress in phenological traits and production of canning peach
38 resulting from the peach breeding program of Embrapa Temperate Agriculture in 53 years. We
39 divided the data records considered in the estimation of genetic progress into two periods, 1964-
40 1984 and 1985-2017, totaling 53 years. The following traits: maturing period, cycle, number of
41 fruits, fruit weight, yield, and soluble solids content were evaluated. We initially tabulated data
42 and analyzed descriptive statistics. Subsequently, we conducted analysis of mixed models and
43 obtained the estimates of genetic progress through meta-analysis. Genetic gain for earliness,
44 shortening the cycle from flowering to maturation, and genetic gain for fruit yield were
45 observed.

46 **Keywords:** *Prunus persica*; genetic gain; peach breeding; fruit development period
47

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55

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60 There is no conflict of interest
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66 **Introduction**

67 The peach breeding program officially started in the municipality of Pelotas at the
68 Experimental Station of Pelotas (today part of Embrapa Temperate Agriculture) in 1963,
69 although germplasm introduction had already been conducted before this date.

70 Initially, the priority of this program was to develop cultivars for production of canning
71 peaches with the main goals of adapting fruits to the mild winter conditions and extending the
72 harvest period. Dr. Sérgio Sachs, who at the time led the program, managed to introduce a
73 valuable germplasm of American origin, followed by the introduction of hundreds of seeds by
74 the breeder Bonifácio H. Nakasu. This germplasm, added to what already existed in the country,
75 ensured the initial variability of the working collection, which has been increasing over the
76 years through other additions. However, one of the difficulties at the time was that most of the
77 material introduced consisted of table fruit producing genotypes. One of the procedures adopted
78 was to cross table and canning fruit genotypes and cross the next generation again with an
79 processing-type fruit cultivar or selection (Raseira et al., 2014). Thus, the breeders obtained
80 canning-type cultivars and selections with early or very early maturation (Raseira, 2010;
81 Raseira et al., 2015, 2018).

82 Over the years, other priorities have been added to the initial goals, such as fruit size,
83 pulp firmness, disease resistance, yield, total soluble solids content, increased pulp-stone ratio,
84 and production stability. The cycle from flowering to maturation also stirred interest since the
85 longer the cycle, the longer the fruit remains in the orchard and the greater the risk of losses
86 caused by climatic phenomena such as strong winds, hail (Raseira et al., 2014), and/or biotic
87 factors. Very early flowering, although being related to low chilling requirement, is undesirable
88 in some cultivation areas due to the occurrence of frosts (Raseira, 2010).

89 Thus, low chilling requirement and late flowering (due to high heat requirement for
90 flowering - Growing degree hours, GDH) may be ideal, as long as the short cycle with rapid
91 fruit growth does not lead to the incidence of what is known as “split stone”. The objective of
92 this work was to estimate the genetic progress in 53 years of the peach breeding program of
93 Embrapa Temperate Agriculture for fruit producing cultivars intended for processing and
94 canning.

95

96 **Material and Methods**

97 **Plant material**

98 Initially, we worked on data of 1687 genotypes obtained in 53 years, being 104 canning
99 cultivars, 298 table fruit cultivars, 613 table fruit selections, and 672 canning selections.
100 However, as many selections are preserved only for new hybridizations due to fruit quality or

101 other traits, not being included in the object of this study and not having the perspective of
102 becoming cultivars, we subsequently decided to only use cultivars that received variety
103 denomination. Nevertheless, the number of data was very large and we decided to separately
104 study table genotypes from those intended for canning (fruit produced for processing).

105 In the present study on genetic progress, we only considered canning peach genotypes
106 released by the Embrapa Temperate Agriculture program. Thus, we analyzed 52 canning
107 cultivars. During the period from 1964 to 1984, the plants were established in the current
108 Cascata Experimental Station (CEE) (31° 37' 15.93" S, 52° 31' 25.20" W, 148.6451 m
109 altitude). After this period, the same cultivars were accompanied at the headquarters of
110 Embrapa Temperate Agriculture (ETA) (31° 42' S, 52° 24' W, 57 m altitude) (1985 to 2017).

111 Collections of the two locations differ both in phytotechnical parameters, such as
112 reduction of spacing between plants from four meters (CEE) to three meters (ETC), as well as
113 in the soil type and its physical and chemical characteristics. At ETC, the soil is shallower and
114 chemically poorer (at a depth of 30 cm), being heavy and moist after this depth, from which
115 roots have difficulty penetrating. However, fertilization remained the same, based on leaf
116 analysis. The two locations also differ in altitude (150 m and 60 m), which is reflected in the
117 temperature, generally 2 °C higher at the ETC, consequently decreasing chilling accumulation
118 by 100 hours (Embrapa, 2020).

119 We analyzed genetic progress considering the genotypes listed in Table 1.

120 **Data collection**

121 The phenological monitoring of the cultivars is conducted yearly observing the dates of
122 beginning of flowering and full flowering and the beginning and end of maturation. Annually,
123 fruits from each cultivar are evaluated regarding fruit color, shape, and size, pulp firmness, total
124 soluble solids content, flavor, adherence to the endosperm, firmness, and health.

125 In the present study, we analyzed the following variables: beginning of maturation
126 (MAT), considering the number of days from January 1st to the beginning of the first harvest of
127 the cultivar in the current season; fruit development period (FDP I), number of days between
128 the beginning of flowering and the beginning of harvest, which we have agreed to name initial
129 cycle; fruit development period considered from full flowering (FDP II), corresponding to the
130 number of days between full flowering (more than 50% open flowers) and the beginning of
131 harvest; total soluble solids (TSS), which is measured in °Brix using a digital refractometer on
132 three to five fruits per cultivar and year; average fruit mass (FM) in grams, determined using
133 an analog scale and a sample of 10 to 20 fruits per cultivar and per season; number of fruits
134 (NF), obtained by counting the number of fruits per plant (unit), and production (PD) per plant,

135 obtained by multiplying the number of fruits per plant by their average mass (Kg), from $PD =$
136 $FM \times NF$.

137 We highlight that in the initial period, from 1964 to 1984, the number of fruits was
138 counted during fruit thinning. The number of fruits removed from the plant was counted and
139 the remaining fruits were counted afterwards. For this purpose, a manual counter was frequently
140 used to help record the numbers. However, fruit counting is very difficult in this period as the
141 plant presents many leaves and the fruits are green. Thus, failures may have occurred. In
142 addition, fruit may fall until harvest, which may have overestimated production at times in this
143 period. However, with the increase in the number of genotypes and reduction of human
144 resources to conduct this task, those counting became almost impossible in the second period.
145 Our option was to assign a degree of production on a scale of 1 to 5, where G1 = 50 fruits; G2
146 = 100 fruits; G3 = 250 fruits; G4 = 300 fruits; and G5 equal to or above 450 fruits. This scale
147 was attributed to be compatible with the 1964-1984 period, but it limited the largest
148 productions.

149 Data were recorded in a physical file. For analysis, they were digitized in an Excel®
150 spreadsheet.

151 **Statistical analysis**

152 Data analysis – genetic progress

153 Data were grouped from the historical series of the breeding program according to the
154 year of inclusion of the genotype in the evaluation. We initially tabulated data and analyzed the
155 descriptive statistics, considering each variable individually. Subsequently, we performed
156 analysis by mixed linear models generalized by *PROC GLIMMIX* from the SAS software. The
157 generalized linear mixed model used was:

$$158 \quad y_{ij} = m + g_i + y_j + \varepsilon_{ij} \quad (1)$$

159 where y_{ij} is the observed data of the variable in genotype i in year j , m is the intercept or
160 the general mean, g_i is the random effect of genotype $g_i \sim \text{IIDN}(0, \sigma_g^2)$, y_j is the fixed effect of
161 year j and ε_{ij} is the effect of the random error $\varepsilon \sim \text{IID N}(0, \sigma_\varepsilon^2)$.

162 The methodology for estimating genetic progress adopted was according to Breseghello
163 et al. (1999). We conducted the following procedures on the SAS software (SAS Institute, 2014)
164 with generalized mixed linear models for each variable using the following commands: PROC
165 GLIMMIX DATA = NAME; CLASS GEN YEAR; MODEL VAR = YEAR/DIST =
166 NORMAL; LSMEANS YEAR/CL COV; RANDOM GEN/SOLUTION CL; RUN;

167 After application of the generalized mixed linear model, in which genotype means for
168 each year were fitted and covariance matrices were obtained, an estimate of the average annual
169 genetic progress was conducted through analysis of the generalized linear regression. The

170 average annual genetic progress was estimated by the linear regression coefficient b_1 of \hat{Y}_K^* ,
171 given by the ratio between the b_1 (*slope*) and the b_0 (*intercept*) from the initial phase of the
172 breeding program for the year 64 (*phase 1*) and 84 (*phase 2*), where b_1 was obtained by the
173 generalized least squares method (Hoffmann and Vieira, 1987). When the value of b was
174 significant, the intercept of the regression regarding the initial theoretical value of the studied
175 period was calculated as follows:

$$176 \quad \hat{\beta} = \frac{\hat{b}_1}{\hat{b}_0} = (X'XD^{-1})^{-1}(X'D^{-1}\hat{Y}^*) \quad (2)$$

177 where \hat{b}_1 is the coefficient for average weights \hat{Y}_K^* according to the year, in order to estimate
178 the average annual genetic gain, and X is the matrix of dimension $a \times 2$, which consists of a
179 column vector of relative and \hat{b}_0 a column vector of 1, 2, ..., a related to b_1 , where a is the
180 number of years. The significance of the model was obtained by the t test.

181 Data analysis – *most important genotype*

182 Among all the genotypes released commercially, some were highlighted for traits that
183 made them more important for the breeding program or made them have a higher rate of
184 adoption by producers. These genotypes were separated in an electronic spreadsheet in order to
185 conduct another analysis according to commands from the R software (Olivoto and Lúcio,
186 2020). In this analysis, the year information was considered fixed effect and genotype a random
187 effect $N(0, \sigma_g^2)$. After obtaining BLUP values for the genotype, the confidence intervals (95%)
188 were obtained and the graphics were built using the *ggplot* package (Wickham, 2016).

189 After obtaining the BLUP values of the most important genotypes, genotypic distance
190 was analyzed using the Euclidean distance through the *factoextra* package and hierarchical
191 grouping through multi-scale *bootstrap* using the *pv_clust* package with alpha of 0.95 and *nboot*
192 100.

193

194 Results

195 We divided the results of the breeding program into the two following periods: the 1st
196 period, from 1964 to 1984, and the 2nd period, from 1985 to 2017, due to the change of location
197 (environment) of the germplasm, whose differences we mentioned in the Material and Methods
198 section. Subsequently, we conducted the stratification of gain estimates for each period
199 separately, coding them as 1964 and 1985, respectively. We divided results by period and by
200 trait, as described below.

201 The variance components of random factors obtained by the generalized linear mixed
202 models are shown in Table 2. For the variance components of traits related to the cycle, we
203 observed a greater proportion of genotypic effects compared to environment effects. On the

204 other hand, environment variance strongly influenced production variables, that is, environment
205 effects provided great variations in the expression of these traits during the evaluated period.
206 For the fixed year effect, we observed significant effect on the factor for all the traits evaluated
207 in all periods according to the F test (Table 2).

208

209 **Descriptive statistics and comparison between years of the general means**

210 **Maturation:** Among the genetic material of the Embrapa breeding program, we
211 observed from early-maturing genotypes (267 days, corresponding to the end of September) to
212 late-maturing genotypes (422 days, corresponding to February). The earliest genotype
213 presented flowering on June 6 and maturation on September 27, while the latest presented
214 flowering on August 2 and maturation on February 26, with a difference of 155 days between
215 the earliest and the latest (Table 3). Regarding the number of days for maturation, we observed
216 a small increase in the number of days in the 1964-1984 period, while the number of days for
217 maturation decreased in the 1985-2017 period, mainly from the genotypes released after 2010
218 (Figure 1 a and b).

219 **Fruit Development Period:** Regarding FDP I_1964, the genotypes with the lowest
220 values presented 100-day of FDP and the genotypes with the highest values presented 209-day
221 of FDP, that is, a difference of 109 days. On the other hand, in FDP I_1985 the minimum values
222 were 77 days and the maximum 219 days, that is, a difference of 142 days (Table 3). In the
223 analysis of FDP_I, we observed reduction in the 1964-1984 period and also observed reduction
224 between 1985-2017 (Figure 1 c and d).

225 Regarding FDP II_1964, the genotypes with the lowest values presented 89-day of FDP
226 and the genotypes with the highest values presented 199-day of FDP, that is, a difference of
227 110 days. On the other hand, in FDP I_1985, the minimum values were 66 days and the
228 maximum was 203 days, with a difference of 137 days (Table 2). For the analysis of results of
229 the FDP_II, we observed reduction in the two periods, that is, from 1964-1984 and 1985-2017,
230 with the greatest reduction in the second period (Figure 1 e and f).

231 **Number of fruits (NF):** Due the different evaluation methods between periods, it was
232 observed considerable variation between absolute values, with a higher NF in the first period
233 (Table 3). Regarding results of NF per plant, we observed increase in the number of fruits per
234 plant between 1964-1984. For the second period, 1985-2017, the NF per plant remained stable,
235 and we highlight that the values were lower than for the first period (Figure 2 a and b).

236 **Average fruit weight:** For FW_1964, we observed minimum values 30 grams lighter
237 than for the 1985 period, in which we observed minimum values of 37 grams. Maximum values
238 of 216 grams and 288 grams for the periods 1964 and 1985, respectively (Table 3) were

239 observed. Regarding FW, we observed stable weight in 1964-1984, although observing
240 variation in some years. We observed a trend of increasing FW in the second period (Figure 2
241 c and d).

242 **Production per plant:** For the variable PD₁₉₆₄, the minimum value of 0.73
243 kilograms/plant, was observed which was approximate to the minimum values in 1985, 0.82
244 kilograms/plant. The maximum values were 123.30 kilograms/plant for PD₁₉₆₄ and 101.91
245 kilograms for PD₁₉₈₅ (the use of a scale of degree of production decreased the amplitude)
246 (Table 2). Regarding PD per plant, it was observed an increase in total production in the 1964-
247 1984 period, which was also observed for the 1985-2017 period, mainly from 2010, when
248 genotypes presented higher average production, considering the average for the year (Figure 2
249 e and f).

250 **Total soluble solids content in fruits:** Regarding TSS, minimum value of 5.7 ° Brix
251 and maximum value of 19.9 ° Brix were obtaining, presenting amplitude of 14.2 ° Brix (Table
252 3). The total soluble solids content was only determined in the second period, 1985-2017. We
253 observed a trend towards reduction of TSS content considering the average value of the
254 analyzed years (Figure 2 g), with great variation between years.

255

256 **Genetic Gain**

257 The results of genetic gain estimates for the 1964-1984 and 1985-2017 periods for the
258 traits studied are shown in Table 4.

259 We observed that the slope was positive for maturing in the 1964-1984 period, resulting
260 in a positive annual gain of 0.05 (%) year, that is, the result showed a trend of a later beginning
261 of maturing, with predominance of late-maturing genotypes. On the other hand, we observed a
262 negative slope for the 1985-2017 period with an estimated genetic gain of -0.05 (%) year, that
263 is, the genotypes released in this period were predominantly of early maturing.

264 In the analysis of genetic gain for the cycle, considering the beginning of flowering, we
265 observed a negative slope, as well as an annual genetic gain of -0.09 and -0.08% for the 1964-
266 1984 and 1985-2017 periods, respectively. If we compare this result with a previous study
267 (Corrêa et al., 2019) considering a period of 16 years starting in 2000 and 24 canning genotypes,
268 nine of which are cultivars and the others advanced selections from the program, we can assume
269 that the fruit development period was more strongly reduced (-12, 7%) between 2000 and 2015.

270 Regarding the cycle from full flowering, we observed that the number of days until
271 flowering reduced in both periods with genetic gains of -0.02 and -0.14%, respectively, for the
272 first and second period.

273 For the number of fruits, we observed a high genetic gain of 7.97% during the first
274 period, that is, breeding was directed towards production of genotypes with a greater number
275 of fruits per plant. On the other hand, we also observed a gain of 0.39% in the second period,
276 although being well below the first period, which means that breeding worked towards other
277 traits of interest.

278 In the analysis of fruit weight, we observed a negative genetic gain of -0.13% from
279 1964-1984, most likely due to the increase in the number of fruits per plant. For the 1985-2017
280 period, a positive genetic gain of 0.38%, was achieved, which may be aligned with a smaller
281 number of fruits and consequently heavier fruits, and also with the fact that cultivars obtained
282 from the 1980s have greater potential for fruit size. This gain was considerably higher in the 24
283 genotypes studied between 2000 and 2015 (5.3%), possibly because only the most important
284 genotypes were selected for the study (Corrêa et al., 2019).

285 Regarding fruit yield, a gain of 19.38% was observed in the first period, 1964-1984, and
286 also a positive genetic gain of 1.23% for the second period, 1985-2017.

287 In the evaluation of TSS content obtained for the second period, 1985-2017, we
288 observed a reduction of genetic gain in the genotypes, corresponding to -0.11% per year. We
289 expected this reduction due to the negative association with the days until maturing, which was
290 negative in this period as already highlighted.

291

292 **Highlighted genotypes**

293 Regarding the evaluation of genotypes highlighted for their importance for the peach
294 breeding program of Embrapa or that were the most planted by fruit growers and are still
295 currently found in the area (even if in a few orchards), Best Linear Unbiased Prediction (BLUP)
296 results for each trait are shown individually in Figure 3.

297 For beginning of harvest (Figure 3 a), great differences were observed between
298 genotypes. For example, the genotype `BRS Libra´ presented 290 days until maturity
299 (corresponding to October 18), while genotypes `Magno´ and `BR 6´ reach maturity after
300 approximately 380 days (corresponding to January 26th of the year following the beginning of
301 counting), that is, approximately 90 days of difference. The mean BLUP value was 340 days
302 from January 1 until maturity.

303 Regarding the initial cycle (Figure 3 b), the genotypes `BRS Libra´ and `Pepita´
304 presented a cycle of approximately 100 days from the beginning of flowering until harvest. On
305 the other hand, the genotypes `Cerrito´ and `Turquesa´ presented a mean cycle of 180 days,
306 with the mean BLUP value of approximately 140 days.

307 Regarding the date of full flowering (Figure 3 c), great differences were observed
308 between genotypes, with the earliest flowering being observed for cultivars `Pepita` and `BRS
309 `Libra`, 100 days, and the latest full flowering date being observed for cultivar `Cerrito` and
310 `Turquesa`, approximately 180 days. The mean BLUP value was of 128 days from January 1st
311 until full flowering.

312 In the evaluation of total soluble solids content (Figure 3 d), genotypes `BR 6`,
313 `Olimpia`, and `Cerrito` presented the highest content. On the other hand, genotypes
314 `Precocinho` and `BRS Citrino` presented the lowest TSS contents. The mean BLUP value was
315 12.5° brix.

316 Regarding the analysis of the most relevant genotypes for the average fruit weight
317 (Figure 3 e), the heaviest weights were observed for cultivars Âambar, Agata, and Granada, with
318 approximately 140 g/fruit. The lightest average fruit weights were observed in the early
319 cultivars BRS_Libra, Precocinho, and Pepita, with approximately 80 g/fruit. The general mean
320 of the most relevant genotypes was approximately 120 g/fruit.

321 In the yield evaluation (Figure 3 f), cultivars Aldrighi and Cerrito presented averages
322 with values of approximately 38 kg/plant and were more productive. Cultivars BRS Libra and
323 Granada with productions in BLUP values close to 21 kg/plant presented the lowest yields. The
324 general mean of the BLUP value was approximately 30 kg/plant.

325 **Genetic diversity**

326 BLUP values for cycle, yield, and brix traits were submitted to cluster analysis by
327 Euclidean distance. Afterwards, considering this group of traits, four groups were formed
328 (Figure 4). The largest group was formed by eight genotypes, cultivars Maciel, Eldorado, Onix,
329 Jubileu, Leonense, Jade, Esmeralda, and Âambar. The smallest group was formed by three
330 genotypes, cultivars BRS Libra, Pepita, and Precocinho.

331

332 **Discussion**

333 The priority goals at the beginning of the breeding program were to offer alternatives
334 for cultivars adapted to the south of Brazil and to extend the period of harvest season, which by
335 then, began in late December or in the first half of January, with the harvest of fruits from
336 cultivar Aldrighi (Medeiros and Raseira, 1998). Until the mid-1970s, this was achieved with
337 the development of late-maturing cultivars. It was only at the end of this decade that the first
338 early-maturing cultivar were obtained, being emphasized after 1984 and mainly in the last 10
339 years (Figure 1), when it is possible to observe a reduction in the number of days until
340 maturation. The cultivar Diamante was the first to allow significant anticipation of the harvest,
341 which occurred approximately 20 days earlier (Medeiros and Raseira, 1998) than cultivar

342 Aldrighi, the peach cultivar planted for the canneries, when the breeding program started. The
343 plant of cv. Diamante was selected from seedlings of the progeny obtained from crossing cv.
344 Convenio, with pollen of at he selection from the Estação Experimental de Taquari/RS, named
345 Pelotas 77. `Convenio´ was the result of open pollination of a cross between the cvs. Amsdem
346 and Abóbora, while Pelotas 77 originated from open pollination of seedlings of the cross
347 between the fresh peach cultivar Cardeal and a canning peach selection that is, (´Cardeal´ ×
348 Aldrighi´) P1 (Table 1). Thus, it is proved that in order to obtain the first very early-maturing
349 cultivars, hybridizations with table peach genotypes was necessary, and in order to achieve the
350 desirable traits for processing, in general, a new hybridization with a canning genotype or
351 another generation is necessary (Byrne et al., 2012).

352 The earliness of maturation period developed by the Embrapa program was an important
353 step in making fruits available for a greater number of weeks and even months. The beginning
354 of the crop season in the region changed from December to October.

355 We did not present the data of flowering periods. However, we used it for calculating
356 the cycles. We observed that the beginning of flowering in some cultivars occurred at the
357 beginning of July or even at the end of June (as in cv. Cerrito) until the end of August (as in
358 Santa Áurea). As, in a simplified way, several breeders estimate the chilling requirement of a
359 cultivar for its flowering period, that is, the chill accumulated up to the flowering date of a
360 cultivar would be its chilling requirement (Rawandoozi et al., 2021), we can observe that
361 amplitude regarding the chilling requirement of the cultivars has been increased, although most
362 of them are among those of low chilling requirement.

363 The result for the number of fruits partly reflects the evaluation system of the two
364 periods studied, since we consider pre-determined intervals and not the actual number of fruits
365 when evaluating degrees. In addition, in the new area - ETC - winter chilling accumulation is
366 lower (Embrapa, 2020) and sometimes insufficient for some cultivars, which leads to decreased
367 and uneven flowering intensity, besides the fact that the soil is shallow and physically and
368 chemically poor, therefore providing less vigorous plants.

369 Similarly to the comparison by Woyann et al. (2019) while working with wheat, the
370 yield gain in the present study for the 1985-2017 period was 1.23%, that is, an increase of 1.23
371 kg for every 100 kg. Extrapolating this idea to 1 ha, which produces an average of 15 tons, that
372 would represent 184 kg more per hectare and year, and between five and six tons more per ha
373 in the 1985-2017 period.

374 It is interesting to note that we only observed a significantly reduction in production
375 (Figure 3 f) in early or very early-maturing cultivars. However, the most recently released
376 cultivars, such as BRS Citrino and BRS Jaspe, did not differ significantly from the late-maturing

377 and more productive cultivars, showing an evolution of the breeding program in the
378 development of early-maturing genotypes.

379 Regarding weight, the price differences between fruit considered of first and second
380 type class in the canneries, encourages the search for production of larger fruits, since yield and
381 profit is higher for the producer and yield in the production line is advantageous for the industry.
382 However, the so-called first quality fruit means being over 5.7 cm in diameter, and any
383 measurement above that has the same value. Diameter is highly correlated with fruit weight,
384 which means that there is no advantage in producing larger fruits beyond a certain average
385 weight, with advantages only for greater numbers of fruit (Raseira et al., 2020). The 5.7cm
386 diameter corresponds to an average weight of approximately 100g per fruit in most cultivars.

387 The emphasis on early maturity negatively affected average fruit weight in the first
388 period, since it is difficult to achieve a large size in very early-maturing and/or short-cycle
389 cultivars. However, although the period from 1985 had greater emphasis on obtaining early-
390 harvest cultivars, maturing in October or early November, the efforts were successful as we still
391 observed positive gain in fruit size. Cultivars like BRS Bonão, Granada, and Maciel certainly
392 contributed to this gain (Raseira et al., 2014).

393 The temperature and light conditions preceding the harvest of early cultivars are not
394 ideal for sugar accumulation, with cloudy and generally cool days (Byrne et al., 2012) in
395 September. We observed in general from figure 3 that the longest cycle cultivars have a higher
396 TSS content. Cultivars that mature in October in southern Brazil experience relatively few hot
397 and sunny days when compared to those whose fruits mature in December or January. The
398 reduction in cycle means that their period for transformation of carbohydrates into sugars
399 decreases. However, this does not mean that quality regarding taste has been reduced.
400 According to literature data (Cirilli et al., 2016), although soluble solids content and sugar are
401 correlated, this index ranges between 0.33 to 0.72 depending on the presence or not of other
402 active compounds, such as pectins, salts, and organic acids (Byrne et al., 1991). On the other
403 hand, there are variations in soluble solids content even among fruits of the same plant,
404 depending on their position, maturation point (Borsani et al., 2009; Lo Bianco and Rieger, 2002;
405 Lopresti et al., 2016).

406 It is interesting to observe the position of cv. Aldrighi in Figures 3 (a) to 3 (f), which
407 was considered the most important clone at the beginning of the program that emphasized
408 canning peach genotypes. Regarding the beginning of harvesting, we observed average
409 anticipation of 80 days or more. The fruit development period was shortened by 50 to 60 days
410 and regarding average fruit weight, only four of the genotypes were inferior to `Aldrighi`. There
411 is a greater number of genotypes with lower soluble solids content than fruits produced by cv.

412 Aldrighi corresponding to cultivars maturing in October or early November, when climatic
413 conditions that anticipate their harvesting in the south of the Rio Grande do Sul State correspond
414 to cool and cloudy days. On the other hand, regarding the difference in production, it is largely
415 a consequence of the anticipation of harvest and shorter cycle.

416

417 **Conclusions**

418 We observed genetic gain for earliness, with the harvest period extended to more than
419 three months.

420 There was anticipation of the beginning of the harvest and shortening in the cycle from
421 flowering to maturation, which did not affect fruit size as new cultivars became available from
422 1985.

423 A genetic gain for both periods of the peach breeding program of Embrapa was achieved
424 for fruit yield, highlighting a genetic gain of 184 kg ha⁻¹year⁻¹ for 1985-2017, corresponding to
425 1.23%.

426

427 **References**

- 428 Borsani, J., Budde, C.O., Porrini, L., Lauxmann, M.A., Lombardo, V.A., Murray, R., Andreo,
429 C.S., Drincovich, M.F., Lara, M. V., 2009. Carbon metabolism of peach fruit after harvest:
430 Changes in enzymes involved in organic acid and sugar level modifications. *Journal of*
431 *Experimental Botany* 60, 1823–1837. <https://doi.org/10.1093/jxb/erp055>
- 432 Breseghello, F., Rangel, P.H.N., De Morais, O.P., 1999. Ganho de produtividade pelo
433 melhoramento genético do arroz irrigado no Nordeste do Brasil. *Pesquisa Agropecuaria*
434 *Brasileira*.
- 435 Byrne, D.H., Nikolic, A.N., Burns, E.E., 1991. Variability in Sugars, Acids, Firmness, and
436 Color Characteristics of 12 Peach Genotypes. *Journal of the American Society for*
437 *Horticultural Science* 116, 1004–1006. <https://doi.org/10.21273/JASHS.116.6.1004>
- 438 Byrne, D.H., Raseira, M.B., Bassi, D., Piagnani, M.C., Gasic, K., Reighard, G.L., Moreno,
439 M.A., Pérez, S., 2012. Peach, in: Badenes, M.L., Byrne, D.H. (Eds.), *Fruit Breeding*.
440 Springer US, Boston, MA, pp. 505–569. https://doi.org/10.1007/978-1-4419-0763-9_14
- 441 Cirilli, M., Bassi, D., Ciacciulli, A., 2016. Sugars in peach fruit: a breeding perspective.
442 *Horticulture Research* 3, 15067. <https://doi.org/10.1038/hortres.2015.67>
- 443 Corrêa, E.R., Nardino, M., Barros, W.S., Raseira, M.D.C.B., 2019. Genetic progress of the
444 peach breeding program of embrapa over 16 years. *Crop Breeding and Applied*
445 *Biotechnology* 19, 319–328. <https://doi.org/10.1590/1984-70332019v19n3a44>
- 446 Embrapa, 2020. Horas de Frio: Maio - Setembro/2020.

447 Hoffmann, R., and Vieira, S. (1987). *Análise de regressão: uma introdução à economia* (2a.
448 ed.). Piracicaba, SP: Hucitec.

449 Lo Bianco, R., Rieger, M., 2002. Partitioning of Sorbitol and Sucrose Catabolism within Peach
450 Fruit. *Journal of the American Society for Horticultural Science*. American Society for
451 Horticultural Science 127, 115–121. <https://doi.org/10.21273/JASHS.127.1.115>

452 Lopresti, J., Goodwin, I., Stefanelli, D., Holford, P., McGlasson, B., Golding, J., 2016.
453 Understanding the factors affecting within-tree variation in soluble solids concentration in
454 peaches and nectarines. *Acta Horticulturae* 249–256.
455 <https://doi.org/10.17660/ActaHortic.2016.1130.37>

456 Medeiros, C.A.B., Raseira, M.D.C.B., 1998. A cultura do pessegueiro. Embrapa - Centro de
457 Pesquisa Agropecuária Clima Temperado, Pelotas, RS.

458 Olivoto, T., Lúcio, A., 2020. Metan: an R package for multi-environment trial analysis. .
459 11:783-789 doi:10.1111/2041-210X.13384. *Methods Ecology Evolution* 11, 783–789.
460 <https://doi.org/10.1111/2041-210X.13384>

461 Raseira, M.C.B., Franzon, R.C., Pereira, J.F.M., Scaranari, C., 2015. The first peach cultivars
462 protected in Brazil. *Acta Horticulturae* 1084, 39–44.
463 <https://doi.org/10.17660/actahortic.2015.1084.3>

464 Raseira, M.D.C.B., 2010. Pêssego Cultivar BRS Libra. *Revista Brasileira de Fruticultura* 32,
465 961–1296.

466 Raseira, M.D.C.B., Franzon, R.C., Feldberg, N.P., Scaranari, C., Pereira, J.F.M., 2020. BRS
467 Jaspe: A processing peach cultivar for low chill areas. *Crop Breeding and Applied*
468 *Biotechnology* 20, 3–6. <https://doi.org/10.1590/1984-70332020v20n1c5>

469 Raseira, M.D.C.B., Franzon, R.C., Pereira, J.F.M., Scaranari, C., Feldberg, N.P., 2018. Peach
470 cultivar BRS citrino. *Crop Breeding and Applied Biotechnology* 18, 234–236.
471 <https://doi.org/10.1590/1984-70332018v18n2a34>

472 Raseira, M.D.C.B., Pereira, J.F.M., Carvalho, F.L.C., 2014. *Pessegueiro*, 1st ed. Embrapa,
473 Brasília, DF.

474 Rawandoozi, Z., Hartmann, T., Byrne, D., Carpenedo, S., 2021. Heritability, Correlation, and
475 Genotype by Environment Interaction of Phenological and Fruit Quality Traits in Peach.
476 *Journal of the American Society for Horticultural Science* 146, 56–67.
477 <https://doi.org/10.21273/JASHS04990-20>

478 SAS Institute, 2014. *Statistical Analyysi System*. Version 9.4 STAT/SAS Software.

479 Wickham, H., 2016. *ggplot2 -Positioning Elegant Graphics for Data Analysis*, Springer, Use
480 R! Springer International Publishing, Cham. <https://doi.org/10.1007/978-3-319-24277-4>

481 Woyann, L.G., Zdziarski, A.D., Zanella, R., Rosa, A.C., de Castro, R.L., Caierão, E., Toigo,

482 M.D.C., Storck, L., Wu, J., Benin, G., 2019. Genetic gain over 30 years of spring wheat
483 breeding in Brazil. *Crop Science* 59, 2036–2045.
484 <https://doi.org/10.2135/cropsci2019.02.0136>

Table 1: List of the 52 genotypes analyzed in 53 years and their pedigree, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021

Cultivar	FP	PP	FPFP	FPPP	PPFP	PPPP
Abóbora	Introd. José melo, colonia Pelotas					
Agata	68201104XC2R19T182	PL	682011041	C2R19T182	Both unknown for being PL	
Aldrighi	Introd. Arthur Kaster-Santa Eulalia-PEL/RS					
Alpes	Aldrighi	Tapes	unkown		Lake City	Intermediário (S 56-32)
Âmbar	Esmeralda	Conserva555	Alpes	RR37201	Ambr.Perret x Cerrito	
Ametista	AlpesxRR31-201	PL	Alpes	RR31-201	Both unknown for being PL	
Atenas	Jade	PI	Alpes	RR53272	Both unknown for being PL	
Bolinha	Prov. Aldrighi					
Bonão	Conserva 594	Pepita	Capdeboscq Madrugador Aldrighi?	x PL	Precocinho	PL
Br2	Aldrighi	Cerrito		?	LakeCityXIntermediario	
Br4	Conserva 162	PL	Aldrighi	Cerrito	unkown	unk.
Br6	Amb. Perret	Tapes	unkown	Intr. cityIntermediario	unkownLake lake city	Intermediario
Brandolasco	intr. Unkown					
Brilhante	Convenio	Pelotas 76	AmsdemXAbobora	PL	Aldrighix Taq. Precoce	unk.
Capdeboscq	Lake cityx Intermediario	Lake City	Intermediari		unk.	unk.
Cerrito	LakeCityXIntermediario	PL	LakeCity	Intermediário	Both unknown for being PL	
Citrino	Conserva I 104	Leonense	Eldorado	Sensação	(Provavel) Brilhante	NJC97 X PL
Convênio	AmsdemxAbobora	PL	Amsdem	Abobora	unk.	unk.
Corisco	Aldr. X Taq Prec.	PL	Aldrighi	Taq Precoce	unk.	unk.
Diamante	Convenio	Pelotas 77	AmsdemxAbobora	PL	(CARDEAL X ALDRIGHI)	PL
Eldorado	Gaudéri	Serrano	Del.xInterludio	PL	unk.	unk.
Esmeralda	Alpes	RR37201	Aldrighi	TAPES	American introd.	
Farrapos	Edmundo Perretx Aldr.	PL	Edmundo Perret	Aldrighi	unk.	unk.
Granada	Granito	PL	Alpes	Cons 102	unk.	unk.
Granito	Alpes	Cons 102	Aldrighi	Tapes	intr USA	
Jade	Alpes	RR 53272	Aldrighi	Tapes	intr. USA	
Jubileu	Bolinha	sel.intr.7-28	Aldrighi	PL	Introd. from Mexico	
Leonense	Brilhantex NJC97	PL	Brilhante	NJC 97	unk.	unk.

Libra	Cons 594	Pepita	(Capx Madrugador)	PL	Precocinho	unk.
Lord	Ab'boraxTaq.Precece	PL	Abóbora	Taq.Precece	unk.	unk.
Maciel	Conserva171	Conserva334	Aldrighi	Pelotas76	712128 Introduçãode Rutgers	
Madrugador	AbóboraxTaq. Precoce	PL	Abóbora	Taq. Precoce	unk.	unk.
Matutino	Aldrighix Taq.Precece PL		Aldrighi	Taq Precoce	unk.	unk.
Magno	AmbrosioPerret	TAPES	Introdução		LAKECity	Intermediário (S 56-32)
Morro Redondo	Lake Cityx Intermediario	PL	Lake City	Intermediario	unk.	unk.
Olímpia	Bolinha	intr 7-28	Prov. Aldrighi			introdução
Ônix	Farrapos	Diamante	Edmundo PerretxAldr.	PL	Convenio	Pel 77
Pepita	Precocinho	PL	Convenio	Pelotas 77	unk.	unk.
Piratini	Ambrosio Perret	Tapes	Intr	Lake city	Intermediario	
Ponteiro	Capdeboseq	Mimoso	Lake city	Intermediário	Leader	Prelúdio
Precocinho	Diamante	PL	Convenio	Pelotas77	Both unknown for being PL	
Real	Intr. De Campinas SP					
Rio Grandense	BrilhantexNJC97	PL	Brilhante	NJC97	unk.	unk.
Safira	Ambrosio Perret	Cerrito	Intr. unk.	LakecityxInt.	unk.	unk.
Santa Aurea	Cerrito	NJC 88	LakeCityXIntermediario	PL	Intr USA	*
Santa Helena	Prov. Intr. Da Colonia Sta Helena					
Sensação	Granito	PL				
Tarumã	Aldrighi	AmsdemxAbobora)PLunk		unk	AmsdemxAbobora	Unk
Topazio	Convenio	Pelotas 76	AmsdemxAbobora	PL	Aldr.xTaq Precoce	PL
Turmalina	Cons 334	Cons.594	Intr.71.21.28		Cap x Madrugador	PL
Turquesa	ConvenioxCerrito	PL	Convenio	Cerrito	unk	unk.
Vanguarda	AlpesxTT53.272	PL	AldrighiTapes		Intr.	

FP: mother female parent; PP: father male parent; FPPF: female parent of the mother; FPPP: male parent of the mother; PFPF: female parent of the father; PPPP: male parent of the father.

Table 2: Estimates of variance components for random effects and significance of the fixed effect (year) of the generalized linear mixed model for each variable and period, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021

Trait	n° genotype [#]	n° year [*]	Covariance Parameter Estimates		Tests of fixed effects	
			Genotype± SE	Residual± SE	Year - F value	Pr>F
MAT (64 -84)	288	16	523.42 ± 125.4	43.68 ± 3.6	27.47	< 0.0001
MAT (85-17)	732	32	622.60 ± 127.4	62.39 ± 3.3	14.85	< 0.0001
FDP I (64 - 84)	276	16	457.40 ± 114.4	79.50 ± 6.8	4.60	< 0.0001
FDP I (85-17)	740	33	463.98 ± 95.2	107.12 ± 5.6	10.32	< 0.0001
FDP II (64-84)	261	16	468.52 ± 115.6	54.23 ± 4.7	6.62	< 0.0001
FDP II (85-17)	654	33	477.97 ± 100.9	88.37 ± 4.9	8.97	< 0.0001
NF (64-84)	181	13	17180.00 ± 6716.8	40314 ± 4292.5	8.18	< 0.0001
NF (85-17)	918	29	3588.28 ± 848.8	12136 ± 566.7	19.00	< 0.0001
FM (64-84)	272	16	319.17 ± 100.47	497.20 ± 42.92	7.28	< 0.0001
FM (85-17)	774	33	401.48 ± 89.32	590.14 ± 29.98	6.20	< 0.0001
PD (64-84)	165	12	304.77 ± 119.42	569.72 ± 63.9	7.99	< 0.0001
PD (85-17)	519	27	22.46 ± 8.39	189.91 ± 11.78	8.88	< 0.0001
BRIX (85 -17)	576	30	4.12 ± 0.91	2.19 ± 0.13	7.79	< 0.0001

[#]number of genotypes evaluated during the period for the variable. ^{*} number of years considered during the period for the variable. SE: standard error

Table 3: Results of descriptive statistics for the variables evaluated, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021

Statistics (unit)	Min	q ₁	mean	q ₃	sd	max
MAT_1964 (n° days)	293.0	345.0	362.5	380.0	23.6	422.0
MAT_1985 (n° days)	267.0	324.0	341.2	361.0	25.4	401.0
FDP I_1964 (n° days)	100.0	138.0	156.0	172.0	22.1	209.0
FDP I_1985 (n° days)	77.0	121.0	138.7	156.0	23.9	219.0
FDP II_1964 (n° days)	89.0	127.0	145.2	162.0	21.7	199.0
FDP II_1985 (n° days)	66.0	110.0	128.4	147.0	23.9	203.0
NF_1964 (n° fruit)	10	140.0	364.61	550.0	262.24	1010.0
NF_1985 (n° fruit)	6.0	146.0	247.9	350.0	147.4	698.0
FW_1964 (g)	30.0	88.0	107.16	125.0	30.11	216.0
FW_1985 (g)	37.0	89.5	113.3	132.0	33.7	288.9
TSS (°Brix)	5.7	11.0	12.8	14.7	2.6	19.9
PD_1964 (kg)	0.73	13.44	39.28	66.73	31.19	123.20
PD_1985 (kg)	0.82	14.02	26.24	37.80	16.64	101.91

MAT_1964 & MAT_1985: Fruit maturation in the 1964-1984 and 1985-2017 periods (respectively); FDP I_1964 & FDP I_1985: fruit development period (considering the beginning of flowering and maturing) in the 1964-1984 and 1985-2017 periods (respectively); FDP II_1964 & FDP II_1985: fruit development period (considering full flowering and maturation) in the 1964-1984 and 1985-2017 periods (respectively); NF_1964 & NF_1985: number of fruits in the 1964-1984 and 1985-2017 periods (respectively); FW_1964 & FW_1985: fruit weight in the 1964-1984 and 1985-2017 periods (respectively); TSS: soluble solids content in the fruits; PD_1964 & 1985: fruit weight in the 1964-1984 and 1985-2017 periods (respectively); Min: minimum observed value; q₁: first quartile; Mean: arithmetic mean; q₃: third quartile; sd: sample standard deviation and Max: maximum observed value. Fruit maturing in the 1964-1984 and 1985-2017 periods, MAT_1964-1984 and MAT_1985-2017.

Table 4: Genetic gain (mean) of each variable for maturing period, fruit development period, number of fruits, and production per plant in the 1964-1984 and 1985-2017 periods, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021

Parameter	B ₀	B ₁	<i>t</i> calc-value	<i>t</i> tab-value	<i>Pr</i> > <i>t</i>	Ggen (annual %)	Ggen (full %)
MAT (64 -84)	359.14	0.19	1.97	2.1448	0.06895	0.05	0.83
MAT (85-17)	349.87	-0.19	-5.60	2.0423	0.00000	-0.05	-1.71
FDP I (64 - 84)	154.55	-0.14	-1.11	2.1448	0.28570	-0.09	-1.49
FDP I (85-17)	145.70	-0.12	-2.85	2.0395	0.00770	-0.08	-2.70
FDP II (64-84)	142.52	-0.04	-0.32	2.1448	0.75369	-0.02	-0.41
FDP II (85-17)	136.56	-0.18	-4.59	2.0395	0.00007	-0.14	-4.34
NF (64-84)	225.66	17.99	3.69	2.2010	0.00356	7.97	103.6
NF (85-17)	231.80	0.90	2.23	2.0518	0.03426	0.39	11.22
FW (64-84)	108.60	-0.14	-0.45	2.1448	0.65960	-0.13	-2.06
FW (85-17)	104.51	0.40	4.05	2.0395	0.00032	0.38	12.56
PD (64-84)	16.15	3.13	5.14	2.2281	0.00044	19.38	232.57
PD (85-17)	22.71	0.28	3.83	2.0595	0.00077	1.23	33.29
BRIX (85-17)	13.18	-0.01	-1.91	2.0484	0.06643	-0.11	-3.24

B₀: intercept, B₁: slope; Ggen (annual%): annual genetic gain and Ggen (full%): total genetic gain. MAT_196 & MAT_1985: Fruit maturing in the 1964-1984 and 1985-2017 periods (respectively); FDP I_1964 & FDP I_1985: fruit development period (considering the beginning of flowering and maturing) in the 1964-1984 and 1985-2017 periods (respectively); FDP II_1964 & FDP II_1985: fruit development period (considering full flowering and maturing) in the 1964-1984 and 1985-2017 periods (respectively); NF_1964 & NF_1985: number of fruits in the 1964-1984 and 1985-2017 periods (respectively); FW_1964 & FW_1985: fruit weight in the 1964-1984 and 1985-2017 periods (respectively); TSS: soluble solids content in the fruits; PD_1964 & 1985: fruit weight in the 1964-1984 and 1985-2017 periods (respectively);

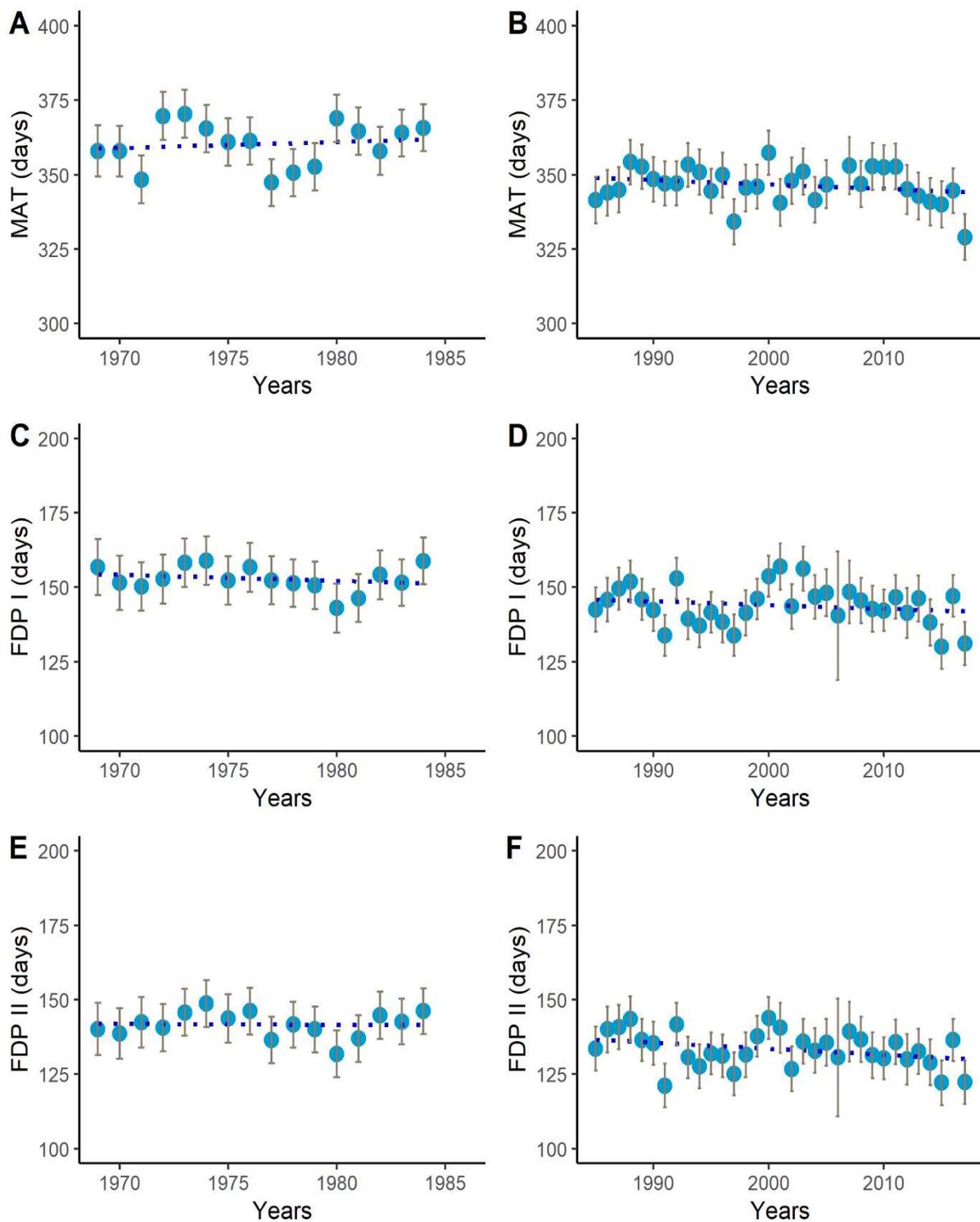


Figure 1: General means of the genotypes in each year studied. (A) Peach fruit maturing during the period 1964-1984 period. (B) Peach fruit maturing during the 1985-2017 period. (C) Cycle considering the beginning of flowering during the 1964-1984 period. (D) Cycle considering the beginning of flowering during the 1985-2017 period. (E) Cycle considering the beginning of full flowering during the 1964-1984 period. (F) Cycle considering the beginning of full flowering during the 1985-2017 period. The circles represent the general and annual means and the vertical bars represent the confidence intervals for the means, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021.

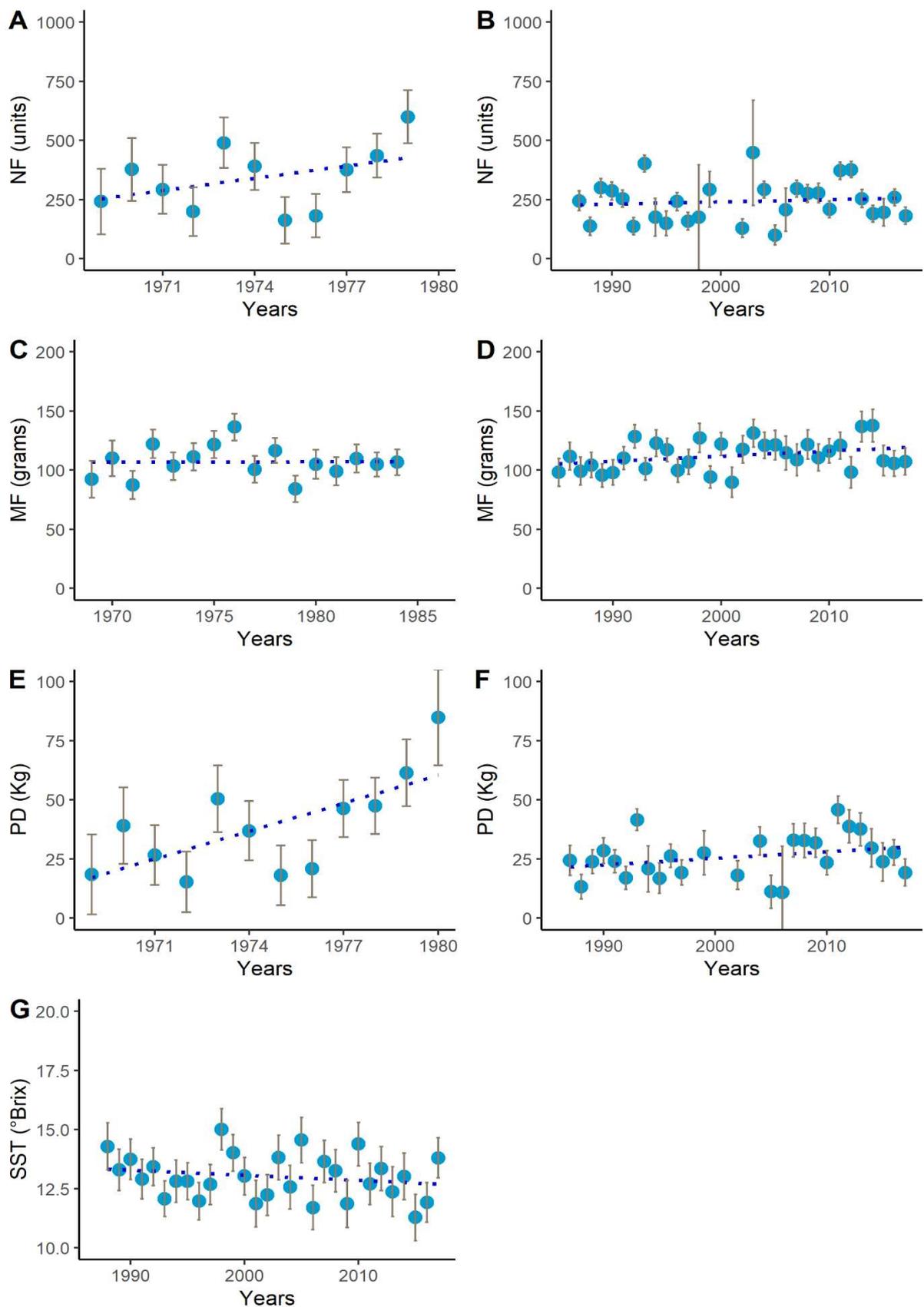


Figure 2: Comparison between general means of the canning peach genotypes in each year. (A) Number of fruits during the 1964-1984 period. (B) Number of fruits during the 1985-2017 period. (C) Fruit weight during the 1964-1984 period. (D) Fruit weight during the 1985-2017 period. (E) Production during the 1964-1984 period. (F) Production during the 1985-2017 period. (G) Mean total soluble solids content in ° Brix during the 1985-2017 period. The circles represent the general and annual means and the vertical bars represent the confidence intervals for the means, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021.

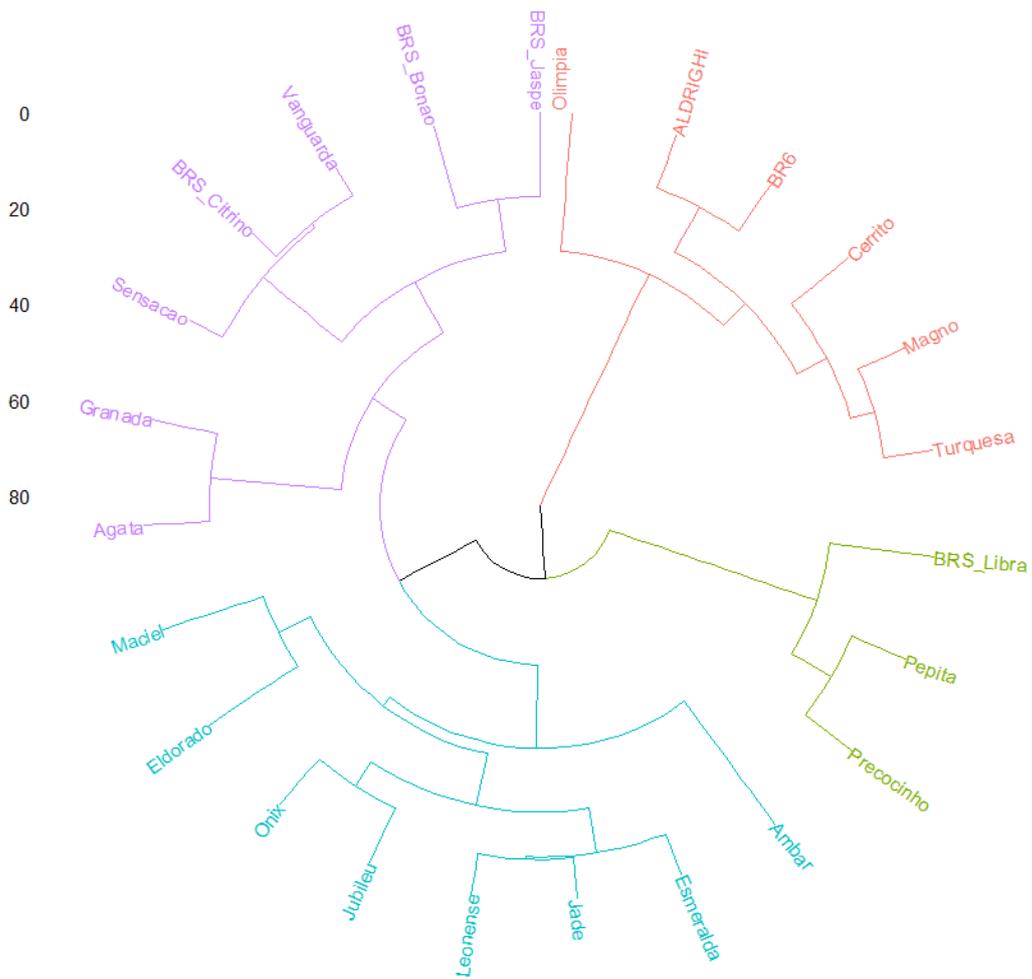


Figure 4: Dendrogram of the currently most planted and cultivated peach cultivars, Embrapa Temperate Agriculture, Pelotas/RS-Brazil, 2021.