CROP BREEDING AND APPLIED BIOTECHNOLOGY

ARTICLE

Yield, maturation cycle, and estimates of genetic parameters of Robusta coffee genotypes under irrigation in the Cerrado

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Abstract: Eighty-five Coffea canephora genotypes were evaluated for three crop seasons under irrigation in the Cerrado of the Distrito Federal, Brazil, to characterize the population regarding phenotype, yield and maturation cycle. The experiment was conducted through the Basic Model of Repeatability without Design methodology. Regarding the cycles, genotypes were divided into four groups with the following periods, in days, from resumption of irrigation to the cherry stage: very early (243-255), early (256-267), medium (268-280), and semilate (281-293). For yield, the repeatability obtained was approximately 33%, a medium value that represents the proportion of the permanent phenotypic variance in relation to the total phenotypic variance. The average repeatability obtained surpassed 59%, and accuracy, with three crop seasons, reached 77%. The selection gains obtained may be higher than 38%. There are promising genotypes for cultivation in the region under study. The mean repeatability and the accuracy obtained favor selection based on phenotypes.

Keywords: Coffea canephora Pierre ex Froehner, repeatability, phenotypic variance, crop breeding.

INTRODUCTION

The *Coffea canephora* species, though high yielding, is less planted than *Coffea arabica* in Brazil, and is practically restricted to the states of Espírito Santo and Rondônia. *C. canephora* is better adapted to the soil and climatic conditions of tropical regions of low altitudes and high air temperatures. Thus, it is not commonly grown in the Cerrado (tropical savanna) of Central Brazil, where it should be established through selection of genotypes adapted to a climate with especially low nighttime temperatures in the winter and to the cultivation system of the region, specifically to mechanized harvesting.

When cultivated at high altitudes, *C. arabica* usually performs better than *C. canephora*, and the opposite is perceived in warm lands at lower altitudes (Rodrigues et al. 2016). Partelli et al. (2011) quote several authors reporting that low air temperatures limit the geographic distribution of coffee, especially when monthly averages are below 15-16 °C; photosynthesis is already affected below 18 °C, and severe cold (chilling and daytime temperatures remaining around 15 °C and nighttime temperatures below 10 °C) has serious consequences on yield. However, the authors also clarify that the plants have a certain capacity for acclimatization to cold.

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The Cerrado has been very conducive to coffee growing, especially when using technologies such as irrigation and mechanized farming practices (Fernandes et al. 2012), especially harvest. The same authors state that a significant limitation for *C. canephora* in the Cerrado is associated with low temperatures in high altitude areas. However, due to considerable genetic diversity within this species (Fonseca et al. 2006), different mechanisms of tolerance to low temperatures are used (Partelli et al. 2011). From this, it can be inferred that selection of materials adapted to cultivation in the central region of Brazil is possible.

Because coffee is a biennial crop, coffee plants have vegetative growth and maturation of reproductive buds for the next year's harvest throughout the reproductive cycle. The planting of cultivars with a late or very late cycle is not recommended in regions with milder air temperatures, for harvest may coincide with the beginning of flowering for the next harvest, which can damage flowers (or buds), as well as risk frost on green fruit. Although frost rarely occurs in Central Brazil, the use of mechanized harvesting, widely diffused in this region, may result in yield loss for the next harvest, due to damage caused to flowers or floral buds in late maturity cultivars, especially when water stress technology is not adopted to promote uniform flowering. Thus, study and differentiation of the maturation cycle of the genotypes under assessment is of paramount importance. Furthermore, according to the same author, the highest yielding cultivars are those of medium or late cycle. Therefore, identification of high yielding materials with early cycles would be of great value, especially to reduce the risk of rainfall during harvest.

Repeatability is a genetic parameter that expresses the proportion of the total variance that is explained by the variations proportioned by the genotypes and by the permanent alterations attributed to the common environment that act on the genotypes. Through study of repeatability, the number of measurements needed to predict the real value of a genotype can be estimated. Repeatability provides the maximum value that heritability in the broad sense can achieve, expressing the proportion of the phenotypic variance that is attributed to genetic differences along with confused with the permanent effects acting on the genotype (Cruz et al. 2012). Thus, through repeatability, one can compare distinct individuals by means of data collected over time, which is quite useful when considering a highly heterozygous heterogamous species like coffee, without the need to reproduce clones of a population of many distinct genotypes. This allows evaluation of a large number of materials in a reduced space (Della Bruna et al. 2012). Besides, since it is related to the minimum number of measurements necessary for accurate selection of superior genotypes, repeatability and maximum broad-sense heritability allows quantifying the maintenance of clone superiority overtime (Rocha et al. 2015).

Perennial plant species, such as coffee, have peculiar biological aspects, for instance, a long reproductive cycle, accentuated annual oscillation of production (resulting in a biennial cycle), overlapping of generations, trait expression over several years, and differences in days to maturity and yield longevity. These characteristics lead to some consequences, such as use of the selected genotypes for several years, use of evaluations repeated in each individual over time, and reduction in the survival rate of individual plants over the time of experiment. This reduction tends to generate unbalanced data for use in estimation of genetic parameters and in prediction of additive and genotypic values. Due to these agronomic peculiarities, coffee breeding is difficult, and the use of special methods to estimate genetic parameters and to predict genetic values is recommended (Oliveira et al. 2011). Thus, the standard analytical procedure that has been recommended for studies in quantitative genetics and also for selection in perennial plants is the mixed model methodology. This approach allows accurate and unbiased prediction of the genetic values even under imbalance, and it also facilitates simultaneous use of information on the individual and the family and of measures repeated over time, providing more accurate estimates of the components of genetic variation and of individual genetic values (Pereira et al. 2013).

The objective of this study was to use repeatability to evaluate the yield and the maturation cycle of *C. canephora* genotypes irrigated in the Cerrado and determine the genetic and phenotypic parameters of the population and the different maturation cycles of these genotypes in order to select high-yielding genotypes for the region.

MATERIAL AND METHODS

The study was performed in the experimental field of Embrapa Cerrados in Planaltina (lat 15° 35′ 30,00″ S, long 47° 42′ 30,00″ W, and alt of 1007 m asl), Distrito Federal, Brazil, on a soil classified as a clayey *Latossolo Vermelho distrófico típico* (Oxisol).

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In previous years, the area had been planted to annual crops. The trial was established in April 2009, with a spacing of 3.5 m between rows and 1.0 m between plants. The 85 *C. canephora* genotypes studied were obtained from natural breeding in an isolated experimental field of the cultivar Robusta Tropical (EMCAPER 8151) from the Capixaba Research and Rural Extension Company (Empresa Capixaba de Pesquisa e Extensão Rural) - EMCAPER. A center pivot sprinkler irrigation method was adopted. The irrigation management criterion was based on monitoring the climate, and the periodicity of irrigation was every five days, according to the Cerrado Irrigation Monitoring Program (Rocha et al. 2008). To synchronize flowering, irrigation was suspended on July 1 and resumed when at least 80% of the flower buds reached the E4 stage, with a final date for return to irrigation on September 4 of each year. In the years in which the experiment was performed, the return to irrigation always occurred on the final date. The practice of subjecting plants to water deficit breaks flower bud dormancy and thus concentrates flowering and fruit maturation in each plant.

In the experimental field, there were about 3,500 genotypes planted without replication repetition. From the harvest data of the 2011/12, 2012/13, and 2013/14 crop years, 85 materials were selected with minimum production of seven liters per plant in each of the three years and production variation lower than 25%. According to Silva et al. (2018), lower biannual variation is as important as yield potential for the selection of clones with higher yield stability that maintain their superiority overtime.

These materials were used in statistical analysis, through repeatability, to obtain the genetic parameter values, phenotypic values, and determination of yield. Preliminary observations, especially those performed by Carneiro et al. (2013), allowed this pre-selection and non-selection of late maturation materials, because of the risk of a later harvest operation compromising production in the following year by causing damage to flowers and flower buds. The standard deviation and the coefficient of variation of the yield of each material during the three harvests were calculated.

The maturation cycles were determined through weekly observation of the genotypes, following the scale developed by Pezzopane et al. (2003), in which a change in level was considered only when at least 80% of the fruit was at the same maturation stage. That way, considering the cherry stage as ideal for harvest, the time between the resumption of irrigation and the cherry stage, in days, was counted, ensuring that each material reached the cherry stage. The difference in days to reach the cherry stage between the earliest material (243 days) and the latest (293 days) was 50 days. Thus, three intervals were made in the mean values (13, 25, and 38 days, approximately), forming four maturation groups, which were denominated very early (243-255 days), early (256-267), medium (268-280), and semi-late (281-293).

Statistical analysis was performed through the Statistical System and Computerized Genetic Selection via Mixed Linear Models, Selegen – REML/BLUP (Resende 2016), using the Basic Model of Repeatability Without Design (model 63 of the program). The Mixed Model Methodology was used to obtain the Best Linear Unbiased Prediction (BLUP) of the genotypic effects and the Restricted Maximum Likelihood (REML) process for estimation of the components of variance and the genotypic parameters.

The basic model of repeatability without design is used when repeated data are taken from individual plants with no defined experimental design. The statistical model is y = Xm + Wp + e where y is the data vector, m is the vector of the effects of measurement (assumed as fixed) added to the overall average, p is the vector of the permanent effects of plants (genotypic effects plus environment effects considered to be permanent, assumed to be random), and e is the error or residue vector (random). Capital letters represent the incidence matrices for the referred effects.

The following values were then obtained: coefficient of repeatability, which, in this case, is considered equal to heritability; permanent phenotypic variance among plants (genotypic variance plus permanent environmental variance from one crop season to the next); temporary environmental variance (environmental variance of each crop season); individual phenotypic variance; average repeatability of *m* crops or repeated measures; accuracy of selection based on the average of *m* harvests or repeated measurements; and the gain from selection for yield in liters per plant.

RESULTS AND DISCUSSION

Table 1 shows the yield of the genotypes in each of the years evaluated, their averages, coefficients of variation of yield, cycle in days from return to irrigation to the cherry stage, and cycle classification.

The highest yields in each of the harvests were obtained by genotypes 82 (20 L), 40 (20 L), and 74 (19 L) in the 2012, 2013, and 2014 harvests, respectively. The overall average yield was 10.79 L per harvest, and the average number of days to reach the cherry stage was 268.6. The average standard deviation of yield was 1.82%, while the coefficient of variation of the yield values over the three years was 16.83%.

Gen	2012	2013	2014	Days	С	CV	Gen	2012	2013	2014	Days	С	CV
5	8	11.5	12	263	E	16.9%	121	8	13	10	263	E	19.9%
6	10	11	16	270	Е	21.3%	122	10	7	7	263	Е	17.7%
7	11	8.5	8	263	Е	14.3%	132	13	8	14	270	Е	22.5%
8	15	10	12	270	Е	16.7%	135	14	19.5	11	270	Е	23.7%
14	12	15	9	270	Е	20.4%	140	10	10	9	263	Е	4.9%
16	12	18.5	11	270	Е	24.0%	144	13	10	7	252	VE	24.5%
17	7	10	10	270	Е	15.7%	145	13	12	16	270	Е	12.4%
18	7	8	10	263	Е	15.0%	146	9	9	13	270	Е	18.2%
24	10	8	11	270	Е	12.9%	147	14	10	12	263	Е	13.6%
27	12	10	17	270	Μ	22.6%	150	14	12	8	263	Е	22.0%
28	17	14	10	263	Е	21.0%	154	8	8	11	277	М	15.7%
32	7	7	11	252	VE	22.6%	164	7	12	8	270	Е	24.0%
40	13	20	15	263	Е	18.4%	165	13	11	12	263	Е	6.8%
44	14	17	9	279	Μ	24.7%	168	8	7.5	7	277	М	5.4%
46	10	10	15	270	Е	20.2%	169	7	9.5	8	263	Е	12.6%
49	11	13	15	279	Μ	12.6%	171	12	11.5	14	263	Е	8.6%
50	7	12	9	279	Μ	22.0%	176	12	13	12	263	Е	3.8%
54	14	10.5	15.5	279	Μ	15.7%	178	8	8	13	277	М	24.4%
60	13	9	13	263	Е	16.2%	181	8	11	12	263	Е	16.4%
62	13	14	10	263	Е	13.8%	183	8	9	10	279	М	9.1%
65	10	9	7	263	Е	14.4%	184	9	7.5	9	263	Е	8.3%
68	7	11	7	263	Е	22.6%	190	9	11.5	8	277	Μ	15.5%
73	8	12	11	270	Е	16.4%	191	7	7.5	9	277	Μ	10.8%
74	11	13.5	19	284	SL	23.0%	193	12	11	10	263	Е	7.4%
78	11	10	10	279	Μ	4.6%	194	16	10	11	279	Μ	21.3%
80	7	11	13	270	Е	24.1%	202	8	9	8	279	Μ	5.7%
82	20	15	18	263	Е	11.6%	205	12	9	13	263	Е	15.0%
83	13	13	19	279	Μ	18.9%	206	13	10	10	263	Е	12.9%
84	12	8	10	263	Е	16.3%	208	10	8	10	279	М	10.1%
88	8	11	7	263	Е	19.6%	209	11	10	7	263	Е	18.2%
89	8	13.5	9.5	263	Е	22.5%	213	14	9.5	8	263	Е	24.3%
90	13	9	13	263	Е	16.2%	216	8	9	12	279	Μ	17.6%
94	17	15	11	279	Μ	17.4%	218	13	12	18	263	Е	18.3%
96	11	9	11	270	Е	9.1%	219	14	8	12	263	E	22.0%
98	10	8	12	270	Е	16.3%	220	11	10	15	279	Μ	18.0%
100	14	10.5	17	270	Е	19.2%	221	13	7.5	10	279	Μ	22.1%
108	10	10.5	10	263	Е	2.3%	225	9	7.5	8	279	Μ	7.6%
109	11	7	7	263	Е	22.6%	226	12	9	8	279	Μ	17.6%
110	13	10	12	263	Е	10.7%	229	11	8.5	8	270	Е	14.3%
111	9	10	9	263	Е	5.1%	231	10	7	10	270	Е	15.7%
112	12	8	8	263	Е	20.2%	236	7	8	12	263	Е	24.0%
114	12	9	15	270	Е	20.4%	238	7	7	7	263	Е	0.0%
118	10	7	10	263	E	15.7%	Aver.	10.88	10.41	11.1	268.6	-	16.4%

Table 1. Yield, in liters per plant, for three consecutive harvests, cycle (Days) from resumption of irrigation to the cherry stage, cycle classification (C), yield averages, and coefficients of variation of 85 genotypes (Gen) of *C. canephora* under irrigation in the Cerrado

Classes of cycle classification: VE = very early; E = early; M = medium; SL = semi late.

The genotypes with the highest yield averages over the three harvests were 82, with an average of 17.67 L per harvest, and 40, with an average of 16.0 L. The main difference between the two was that genotype 82 had a more stable yield, with variation of less than 12% over the 3 years. This factor, considering the desirability of materials with low bianniality, can be of great value. Furthermore, genotype 82 was classified as early, with a cycle of 263 days from return to irrigation to the cherry stage, whereas genotype 40 took 279 days. From the aspect of yield stability, some genotypes stood out through very low variation (less than 10%), or even had no variation. The results obtained from repeatability analysis in each of the situations tested are shown in Table 2.

The permanent phenotypic variance among plants, the variance regardless of temporary environmental effects, i.e., the effects of the environment in each crop season, represented 30% of the total phenotypic variation. This factor shows the lower effect of temporary variations on the response of the materials, or greater adaptation to local conditions. Individual repeatability, which in this case can be considered equivalent to heritability, reached a value over 0.38 (Table 2). According to Resende (2009), repeatability values between 0.30 and 0.60 are considered medium. However, considering the low number of repetitions (only 3) and that coffee is a perennial species, these values of repeatability will certainly be more consistent with a higher number of repetitions, or as the cycles of selection and evaluation of genotypes progress.

In addition, when considering the average repeatability (rm), the value rises to 55.5%, a value, according to the same author, very close to the values considered high for this parameter. Ramalho et al. (2016), working with robusta coffee in the state of Rondônia, obtained coefficients of repeatability of 0.43 in an analysis of four harvests, while they obtained 0.427 for heritability in the broad sense. The proximity of these two values is noteworthy, which corroborates the idea that the two parameters may be considered equivalent. Another factor to be considered is that yield is highly influenced by environment.

Repeatability can be interpreted as the maximum value of heritability in the broad sense, since it expresses genotypic variance added to the permanent effects of environment. In this sense, the repeatability values obtained in this study are close to those perceived by Ramalho et al. (2016) working with robusta coffee in the state of Rondônia. In addition, Mistro et al. (2008) observed values for the repeatability coefficient between 0.26 and 0.63, similar to those presented here. It should be noted, however, that these authors used six and nine repetitions, respectively, always with two plants per plot, whereas here, only one repetition was used, with one plant per plot, and harvests for three consecutive years.

The values of accuracy in relation to the number of measurements are shown in Table 3. Fonseca et al. (2004) cited that four successive harvests would be sufficient for selection of robusta coffee genotypes for yield with an accuracy of 80%, a value also considered quite satisfactory by Resende (2009), for whom selective accuracy between 0.7 and 0.9 is considered high. Pereira et al. (2013) obtained estimated accuracy of 0.84 in three harvests in an experiment with four repetitions in two years. In the experiment described here, the accuracy obtained at 79% with four harvests, very close to the values obtained by the authors cited. Selective accuracy depends on the heritability and repeatability of the trait, the quantity and quality of information, and the procedures used to predict

Table 2. Values of the genetic parameter obtained from analysis
of repeatability of C. Canephora in three harvests

Genetic Parameter	Value				
Vpp	2.5479				
Vte	5.9377				
Vip	8.4049				
r	0.2935 ± 0.0938				
rm	0.5549				
Acm	0.7449				
Overall Average	10.79				

Vpp: Permanent phenotypic variance among plants; Vte: variance of temporary environment; Vip: individual phenotypic variance; $r = h^2$: individual repeatability; rm: average repeatability; Acm: average accuracy of the selection based on the average of m harvests.

Table 3. Selection accuracy in accordance with the number of repeated measurements for yield in robusta coffee

Nr. of measurements or repetitions	Accuracy			
1	0.5418			
2	0.6737			
3	0.7449			
4	0.7902			
5	0.8216			
6	0.8448			
7	0.8626			
8	0.8768			
9	0.8883			
10	0.8978			

genetic values. Selective accuracy is a measure associated with accuracy in selection, that is, it refers to the correlation between predicted genetic values and true genetic values of individuals, and the greater the accuracy in an individual assessment, the greater the confidence in the assessment and in the predicted genetic value of the individual. Accuracy is also the main element of genetic progress that the breeder can change in order to maximize genetic gain, and refers to the correlation between the true genotypic value of the genetic material and the estimated or predicted value (Lopes et al. 2018).

Order Order Gen fp u + fp Gain New Average Gen fp u + fp Gain New Average 1 82 4.088 14.874 4.088 14.874 44 121 -0.269 10.517 0.962 11.748 2 40 3.097 13.884 3.593 14.379 45 146 -0.269 10.517 0.935 11.721 3 83 2.503 13.290 3.230 14.016 46 181 -0.269 10.517 0.908 11.695 4 135 3.023 47 108 -0.368 0.881 2.404 13.191 13.810 10.418 11.668 5 74 12.993 -0.368 2.206 2.860 13.646 48 221 10.418 0.855 11.641 6 94 12.894 49 -0.467 0.828 2.107 2.734 13.521 84 10.319 11.614 7 218 2.107 12.894 2.645 13.431 50 98 -0.467 10.319 0.802 11.589 8 16 1.810 12.597 2.541 13.327 51 144 -0.467 10.319 0.777 11.564 9 100 1.810 12.597 52 24 -0.665 0.750 2.459 13.246 10.121 11.536 10 28 1.711 12.498 2.385 13.171 53 140 -0.665 10.121 0.723 11.509 1.711 54 178 -0.665 11 145 12.498 2.323 13.110 10.121 0.697 11.484 55 11.459 12 44 1.513 12.300 2.256 13.042 216 -0.66510.121 0.673 13 54 12.300 2.199 12.985 56 226 -0.665 0.649 1.513 10.121 11.435 27 57 -0.764 14 1.315 12.101 2.136 190 10.022 0.624 12.922 11.410 49 15 1.315 12.101 2.081 12.867 58 50 -0.863 9.923 0.598 11.384 16 171 1.018 11.804 2.015 12.801 59 111 -0.863 9.923 0.573 11.360 6 17 0.919 11.705 1.950 12.736 60 112 -0.863 9.923 0.549 11.336 8 0.919 11.705 1.893 208 -0.863 9.923 0.526 11.313 18 12.679 61 209 19 62 0.919 11.705 1.842 12.628 62 -0.863 9.923 0.504 11.290 20 176 0.919 11.705 63 7 -0.962 1.795 12.582 9.824 0.481 11.267 21 194 0.919 11.705 1.754 12.540 64 229 -0.962 9.824 0.458 11.244 22 11.507 65 -1.061 0.435 14 0.721 1.707 12.493 17 9.725 11.221 23 114 0.721 11.507 1.664 12.450 66 118 -1.061 9.725 0.412 11.198 24 147 0.721 11.507 1.625 12.411 67 154 -1.061 9.725 0.390 11.176 25 0.721 11.507 12.375 68 164 -1.061 9.725 0.369 165 1.588 11.155 26 220 0.721 11.507 1.555 12.341 69 183 -1.061 9.725 0.348 11.134 70 27 231 46 0.523 11.309 1.517 12.303 -1.0619.725 0.328 11.114 28 60 0.523 11.309 1.481 71 236 -1.061 9.725 0.308 11.095 12.268 29 90 0.523 11.309 1.448 12.235 72 65 -1.259 9.527 0.287 11.073 73 30 110 0.523 11.309 1.418 12.204 88 -1.259 9.527 0.265 11.052 31 0.523 11.309 1.389 74 184 -1.358 0.243 132 12.175 9.428 11.030 32 150 0.325 11.111 1.355 12.142 75 18 -1.4579.329 0.221 11.007 33 205 0.325 11.111 1.324 12.110 76 32 -1.457 9.329 0.199 10.985 34 219 0.325 11.111 1.295 12.081 77 68 -1.457 9.329 0.177 10.963 10.913 78 35 193 0.127 1.261 12.048 109 -1.457 9.329 0.156 10.942 36 206 0.127 10.913 1.230 12.016 79 202 -1.457 9.329 0.136 10.922 37 5 -0.170 10.616 1.192 11.978 80 169 -1.556 9.230 0.115 10.901 38 213 -0.17010.616 1.156 11.943 81 225 -1.556 9.230 0.094 10.880 39 73 -0.269 10.517 1.120 11.906 82 122 -1.655 9.131 0.073 10.859 40 78 -0.269 10.517 83 191 -1.754 9.032 0.051 1.085 11.871 10.837 41 80 -0.269 10.517 1.052 11.838 84 168 -1.952 8.834 0.027 10.813 42 89 -0.269 10.517 1.021 11.807 85 238 -2.249 8.537 0.000 10.786 43 96 -0.269 10.517 0.991 11.777

Table 4. Classification, genotype (Gen), phenotypic value (fp), sum between phenotypic and average value (u + fp), selection gain (in liters per plant), and new average of the population of 85 genotypes of C. canephora under irrigated growing in the Cerrado

Working with the repeatability coefficient in peach, Della Bruna et al. (2012) obtained a variation of the coefficient of repeatability between 0.22 and 0.29 for the fruit yield per plant trait, with estimates of accuracy (Acm) between 60.0 and 75.0 %. The repeatability (0.30) and the accuracy (75%) obtained in this experiment are therefore within the values observed for other fruit plants. Pedro Junior et al. (2013), working with lemon tree cultivation, obtained coefficients of individual repeatability (r) around 0.1, reaching 0.35 in the average of 5 harvests. The accuracy obtained by these authors was 58% in one harvest and 84% in five harvests, values near those shown here, 54.2% and 82.1% in one and five harvests, respectively. It is noted, therefore, that an increase in the number of measurements allows an increase in the accuracy of selection. However, these increases become smaller and less significant each year, and it can be inferred that from the fifth harvest on, increases in accuracy would not justify the necessary input of resources.

The variability of the population allows expressive gains from selection of superior genotypes, up to approximately 40% of the mean of the population under study. Ramalho et al. (2016) obtained estimates of selection gains of 43.8% in the first year, working with robusta coffee in the state of Rondônia. Therefore, the results presented here are also close to those found by those authors.

The prediction of permanent phenotypic value is extremely relevant for it allows selection of individuals that will be cultivated in an environment similar to the one under evaluation, and thereby the additive, dominance, and permanent effects of environment are capitalized. In addition, it is possible to select genotypes for vegetative propagation outside the environment in which they were evaluated, and thus the additive and dominance effects are capitalized (Della Bruna et al. 2012). The phenotypic values of the individuals, as well as the selection gain inherent to each of them, are shown in Table 4.

Thus, there is genetic variability within the population studied in relation to yield and maturation cycle. Promising genotypes were found for production under irrigation within the population of *Coffea canephora* studied in the Cerrado region of Brazil. Materials have been identified for utilization in crosses to maximize heterotrophic effects and gene complementarity within a breeding program for robusta coffee under irrigation management.

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