CONTRIBUTION OF AGRONOMIC TRAITS TO THE COFFEE YIELD OF *Coffea canephora* **Pierre ex A. Froehner IN THE WESTERN AMAZON REGION**

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(Received: March 20, 2018; accepted: June 12, 2018)

ABSTRACT: The evaluation of morphological characters related to the hulled coffee yield subsidizes the selection of *Coffea canephora* plants that combine a set of favorable traits. The aim of this study was to evaluate the direct and indirect effects of agronomic traits on the production of hulled coffee to subside the plant selection. To this, nine morphological descriptors were evaluated of 130 clones of the botanical varieties Conilon and Robusta over two crop years in the experimental field of Embrapa, in the municipality of Ouro Preto do Oeste, state of Rondônia (RO). To quantify the genetic variability the path analysis and the Scott Knott cluster test were used. The effect of genotype x year interaction was significant for eight of the nine characteristics analyzed. The genotypes were clustered in three to five classes, subsidizing the establishment of a scale to evaluate the variability of this genetic resource. Pathway analysis indicated that the number of plagiotropic branches and the number of rosettes per productive branch were the traits that exhibited the greatest direct effect on hulled coffee yield. These results show that it is possible to select plants with complementarity traits which favor a higher production of hulled coffee.

Index terms: Conilon coffee, Robusta coffee, path analysis, genetic variability.

CONTRIBUIÇÃO DE CARACTERES AGRONÔMICOS PARA A PRODUTIVIDADE DE CAFÉ BENEFICIADO DE *Coffea canephora* Pierre ex A. Froehner NA REGIÃO AMAZONICA OCIDENTAL

RESUMO: A utilização simultânea de caracteres morfológicos associados à produtividade de café beneficiado subsidia a seleção de plantas de *Coffea canephora* que apresentem um conjunto de características favoráveis. O objetivo desse estudo foi avaliar os efeitos diretos e indiretos de características agronômicas sobre a produção de café beneficiado de *C. canephora* subsidiando a seleção de plantas. Para isso foram considerados nove descritores morfológicos avaliados em 130 clones das variedades botânicas Robusta e Conilon ao longo de dois anos agrícolas no campo experimental da Embrapa, no município de Ouro Preto do Oeste, estado de Rondônia (RO). Para quantificar a variabilidade genética, utilizou-se a análise de trilha e o teste do agrupamento Scott Knott. O efeito da interação genótipo x ano foi significativo para oito das nove características analisadas. Os genótipos foram agrupados em três a cinco classes, delimitando uma escala que pode ser utilizada para avaliar a variabilidade deste recurso genético. A análise de trilha indicou que o número de ramos plagiotrópicos e o número de rosetas por ramo produtivo foram as características que apresentaram o maior efeito direto sobre o rendimento do café beneficiado.

Termos para indexação: Café Conilon, café Robusta, análise de trilha, variabilidade genética.

1 INTRODUCTION

The *Coffea canephora* Pierre ex A. Froehner represents approximately 35% of world coffee production and it is characterized by high plant vigor and high yield (Ico, 2016). In the Western Amazon, the state of Rondônia stands out as a traditional coffee producer with approximate production of 1.93 million bags of hulled coffee in 2017 (Conab, 2017).

This species presents two botanical varieties with distinct characteristics that are commercially cultivated (Davis et al., 2006). The botanical variety Robusta is characterized by erect growth, larger leaf size, higher average sieve, late maturation, less tolerance to water deficit, and greater resistance to diseases and pests.

The botanical variety Conilon is characterized by shrub growth, early maturation, elongated leaves, greater tolerance to drought, and greater susceptibility to pests and diseases (Montagnon et al., 2012; Oliveira et al., 2018).

In modern agriculture, two main strategies are considered to increase yield per area: planting genetically superior materials and improving growing conditions (Ramalho et al., 2012; Ferrão et al., 2017). In various crops, an additive relationship between these factors has been observed, the result of the expression of superior genetic potential with improvement of growing conditions (Dubberstein et al., 2017; Rodrigues et al., 2016). In classical breeding of *C. canephora*, the aim is to select plants with superior yield potential associated with greater uniformity of ripening, greater sieve size, and smaller plant architecture (Teixeira et al., 2017).

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In addition to the long reproductive cycle, the overlapping of generations, and the vigor that is manifested in plants from divergent crosses; association among the coffee bean yield components has considerable implications for breeding of this species (Souza et al., 2017; Rocha et al., 2015). According to Cruz et al. (2014), genetic gain from direct selection is defined as changes in the population mean resulting from selection practiced on the trait itself. However, it is known that selection in one main trait may result changes that can favor or hinder the selection of plants that combine a series of favorable characteristics.

Pathway analysis, originally proposed by Wright (1923), allows the correlation coefficients to be broken down into their direct and indirect effects on a main trait. Appropriate interpretation of pathway analysis depends on the creation of a cause and effect diagram, which represents the relationship of secondary traits on a main trait when complex relationships are involved (Cruz et al., 2014).

Bikila and Sakiyama (2017) observed significant estimates of the simple correlation coefficient between yield, number of plagiotropic branches and plant height. Cilas et al. (2006) observed significant phenotypic correlations among the distance between rosettes, the number of rosettes and hulled coffee yield. Ferrão et al. (2008a) observed higher magnitude of the genotypic correlations compared to the environmental correlations, selecting plants of higher yield potential. Studies were not found in the literature that estimated the direct and indirect effects of other agronomic traits on *C. canephora* hulled coffee yield.

In this context, the aim of this study was to quantify the direct and indirect effects of morphological and yield traits on the hulled coffee bean yield of *C. canephora*, for the purpose of subsidizing the characterization and use of the genetic variability of this species.

2 MATERIALS AND METHODS

Field experiment

Over the 2013-2014 and 2014-2015 crop years, nine morphological and yield descriptors were evaluated from 130 genotypes with characteristics of the botanical varieties Robusta (37), Conilon (75), and interspecific hybrids (18): plant height (PHt), number of plagiotropic branches (NPLAG), number of rosettes per branch (NROS), length of plagiotropic branches (LPLAG), number of fruits per rosette (FROS), distance between rosettes (DROS), number of days to fruit ripening (NDAYS), hulled coffee bean yield (YLD), and mean sieve size (SIEV). The field experiment was managed according to Marcolan et al. (2009).

The assay installed in November 2011 was carried out in the experimental field of Embrapa, in the municipality of Ouro Preto do Oeste, state of Rondônia (RO). The Climate in the region, according to Köppen, is tropical rainy, with mean annual rainfall of 1939 mm/year and mean annual temperatures from 21.2°C to 30.3°C. The region is located at 10°37'03"S and 62°51'50"W, with relative humidity throughout the year around 81%. Soil in the experimental area is eutrophic red-yellow oxisol with clayey texture and flat topography, it is a deep and well-drained soil. The experimental design was randomized blocks with four replications of four plants per plot at spacing of 3.5 x 1.5 meters, between plants and between rows, respectively.

Multicollinearity diagnosis and pathway analysis

Data were subjected to analysis of variance and analysis of homoscedasticity. With the aim of clustering the clones in divergent and mutually exclusive groups, the Scott Knott mean clustering test was used at 5% probability (Cruz, 2013).

Pathway analysis was conducted by estimating the phenotypic correlations (r_f) between the traits and their significance was tested at 1% and 5% probability by the t test at (n-2) degrees of freedom (Cruz et al., 2014). Multicollinearity was diagnosed according to the condition number (CN) of the singular matrix X'X, defining the ratio between the largest and smallest eigen value of the correlation matrix. According to Montgomery & Peck, (1981), if CN<100, multicollinearity is weak and does not constitute a problem for analysis; if 100<CN<1000 multicollinearity is moderate to strong; and if CN>1000, it is severe.

To obtain estimates of the direct and indirect effects, a causal diagram was established that considers the logical, additive relationship among the primary and secondary yield components (Figure 1). The statistical analyses were done using the statistical software Genes, version 1990.2018.18 (Cruz, 2013).

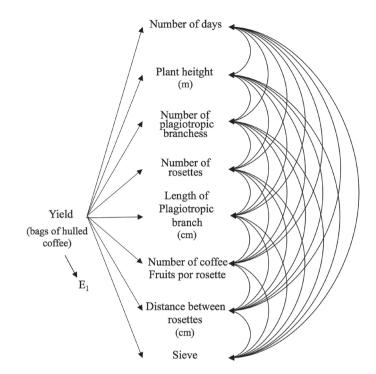


FIGURE 1 - Illustrative diagram representing in the single-headed arrows, the direct effects of eight explanatory variables on the main variable, yield. The bidirectional arrows represent the indirect effects quantified by the phenotypic correlation coefficients.

3 RESULTS AND DISCUSSION

The effect of the clone \times year interaction wasn't significant for only one of the nine traits analyzed (Table 1). The significant effect of the clones x year interaction difficult the improvement of perennial species, as it results in a decrease in the genetic progress obtained with plant selection resulting from changes in the order of clones selected from one year to the next. Expressive clone \times year interaction were also observed by Souza et al. (2013) and Montagnon et al. (2012) in genotypes evaluated in the state of Espírito Santo and in Ivory Coast.

Characterization of *C. canephora* genotypes should consider both superior performance and maintenance of superiority over time (Silva et al., 2017). Although the effect of the genotype \times year interaction was significant, except for SIEV, the repeatability estimates of the traits evaluated indicate the predominance of simple interaction (Resende; Duarte, 2007) (Table1). Simple interaction is characterized by smaller changes in ordering of the genotypes over time, unlike complex interaction, which is characterized by the absence of correlation in ordering of genotypes over time (Cruz et al., 2014). The Scott Knott cluster test at 5% probability was used with the aim of grouping the genotypes in mutually exclusive classes. This test allows the genotypes to be grouped in sets of minimum variation within groups and maximum variation among groups, making it easier to interpret results, due to the absence of ambiguity (Bhering et al., 2008). The clones evaluated were clustered in three to five mutually exclusive groups, depending on the trait (Table 2).

The grouping based in the number of days to fruit ripening (NDAYS) indicated the existence of three different times of fruit ripening, with approximately 20 days difference between them (Table 2). This grouping corroborates the variability observed in the field in which clones that mature in Abril, May, and June are called early, intermediate, and late clones, respectively. Partelli et al. (2014) and Morais et al. (2012) observed genotypes with fruit ripening in March and July called super early and super late, respectively.

For hulled coffee yield (YLD), four different yield classes were observed, with amplitude from 8.8 to 72.9 bags.ha⁻¹ in the first harvest and from 11.3 to 123.7 bags.ha⁻¹ in the second harvest (Table 2).

TABLE 1 - Summary of the F estimates of the joint analyses variance for the traits: number of days to fruit ripening (NDAYS), plant height (PHt), number of plagiotropic branches (NPLAG), number of rosettes per branch (NROS), length of plagiotropic branches (LPLAG), number of fruits per rosette (FROS), distance between rosettes (DROS), yield (YLD), and mean sieve size (SIEV) of *C. canephora* coffee, evaluated in the 2013-2014 and 2014-2015 crop years.

S.V.	NDAYS	PHt	NPLAG	NROS	LPLAG	FROS	DROS	SIEV	YLD
Blocks									
Clones	6.78**	5.52**	5.92**	1.79**	4.94**	1.51*	2.72**	4.57**	3.64**
Years	2.10**	1387.98**	169.47**	68.43**	14.96**	26.33**	24.67**	0.51^{NS}	148.38**
Clones xYears	2.94**	4.43**	2.29**	3.11**	2.70**	8.12**	7.84**	2.96 ^{NS}	3.90**
Residue									
Mean	302.0	1.58	81.34	10.33	0.79	17.01	4.91	15.09	43.31
Mean _{1st measure}	302.7	1.32	67.95	9.44	0.77	18.09	5.06	15.12	33.70
Mean_2nd measure	301.2	1.84	94.72	11.22	0.80	15.92	4.76	15.06	52.91
Repeatability	85.25	81.89	83.13	44.38	79.83	33.68	63.23	78.14	72.51
CV	3.25	6.65	26.91	19.09	9.43	14.03	7.25	5.43	29.72

*and ** = significant at 1% and 5% probability by the F test, respectively. r = repeatability; CV = coefficient of variation.

NDAYS 2013-2014		NDAYS 2014-2015		PHt 2013-2014		PHt 2014-2015		NPLAG 2013-2014		NPLAG 2014-2015	
Classes	f	Classes	f	Classes	f	Classes	f	Classes	f	Classes	f
271 288	34	262 286	28	0.9 1.3	59	1.4 1.6	32	13 71	79	13 73	31
295 314	71	288 314	78	1.3 1.5	52	1.6 1.8	27	75 100	32	74 102	44
317 332	25	317 349	24	1.5 1.6	14	1.8 1.9	44	104 121	16	103 128	34
-	-	-	-	1.6 1.9	5	2.0 2.4	27	136 143	3	129 206	21
LPLAG		LPLAG		NROS		NROS		FROS		FROS	
2013-2014		2014-2015		2013-2014		2014-2015		2013-2014		2014-2015	
Classes	f	Classes	f	Classes	f	Classes	f	Classes	f	Classes	f
0.50 0.77	70	0.50 0.78	60	5.7 7.6	21	5.7 11.4	74	9 17	52	10 15	75
0.78 0.86	38	0.79 0.86	41	7.7 9.5	48	11.5 13.3	34	18 22	66	16 19	35
0.87 0.94	16	0.87 0.94	17	9.6 11.2	40	13.5 14.4	14	23 27	10	20 23	12
0.96 1.00	6	0.95 1.12	12	11.3 14.5	21	14.7 22.0	8	30 35	2	23 30	8
DROS		DROS		YLD		YLD		SIEV		SIEV	
2013-2014		2014-2015		2013-2014		2014-2015		2013-2014		2014-2015	
Classes	f	Classes	f	Classes	f	Classes	f	Classes	f	Classes	f
3.6 4.3	18	3.4 4.3	34	8.8 20.8	36	11 46	54	19 18	10	19 18	11
4.4 5.4	81	4.4 5.4	50	21.8 46.8	66	47 70	45	17 16	33	17 16	33
5.5 5.9	20	5.1 5.9	10	47.5 56.5	20	72 81	21	15 14	65	15 14	67
6.0 7.2	11	6.0 7.5	36	58.4 72.9	8	85 123	10	13 12	18	13 12	17
-	-	-	-	-	-	-	-	11 10	4	11 10	2

TABLE 2 - Grouping by interval of classes of the C. canephora coffee yield traits. Embrapa, RO, Brazil.

f = simple absolute frequency; NDAYS = number of days to fruit ripening; PHt = plant height; NPLAG = number of plagiotropic branches; LPLAG = length of plagiotropic branches; NROS = number of rosettes; DROS = distance between rosettes; FROS = number of fruits per rosette; YLD = hulled coffee yield; SIEV = mean sieve size.

Hulled coffee yield is a quantitative trait of continuous distribution expression, influenced both by the genotype and by the environment (Cruz et al., 2014). The establishment of mutually exclusive groups ordered according to the variability of the trait, allowed establishment of scales that differentiate plants with low, medium, high, and very high yield (Table 2).

Plant height (PHt) was grouped in four different classes, with amplitude from 0.92 to 1.87 m in the first harvest and from 1.36 to 2.39 min the second; a mean increase of 0.90 m between the first and second year of evaluation was observed. Bergo et al. (2008) observed genotypes of *C. canephora* that after four years of growth in the field exhibited mean height of 3.55 m in comparison with the cultivars Icatu and Catuaí of *C. arabica*, which exhibited 3.33 m and 2.71 m height, respectively. According to Ferrão et al. (2008b), this trait is predominantly under genetic control, with heritability estimates higher than 0.60.

The NPLAG and LPLAG traits were grouped in four different classes (Table 2). For NPLAG, an increase in the mean value from 26 to 47 plagiotropic branches was observed between the first and second year; and for LPLAG, a mean increase of 120 cm was observed between the first and the second year (Table 2). This classification allows to identify plants of greater yield potential, which stand out through a higher number and greater length of productive branches.

The NROS, FROS, and DROS traits were also grouped in four different classes. The NROS exhibited an increase in the mean number of rosettes over time, from 8.8 to 16.3, unlike FROS, which exhibited a similar mean performance over time of 18.09 in the first year and 15.92 in the second year of evaluation (Table 2). For DROS were identified groups with minimal, small, intermediate, and large distance between rosettes (Table 2).

In relation to mean sieve size (SIEV), distribution of clones was observed in five classes, with amplitude from 10 to 19, with a small variation over time of 14.82 in the first year and 14.75 in the second year. Ramalho et al. (2016) observed greater amplitude in genotypes of the Conilon botanical variety, from 13.6 to 17 and a mean sieve size of 15.4.

The study of the direct and indirect effects of the plant traits on coffee yield allows to identify the degree of importance of the explanatory variables to the main variable (Sureshkumar et al., 2013). The pathway diagram shows the relationship between the primary components and hulled coffee yield (Figure 1). To estimate the direct and indirect effects of pathway analysis, it is necessary that the X'X matrix not have high levels of multicollinearity, since in the presence of multicollinearity, the variances associated with the estimators of the pathway coefficients can reach excessively high values (Cruz et al., 2014). According to Montgomery & Peck (1981), the estimates of phenotypic correlations showed weak multicollinearity, and it was not necessary to discard any of the traits analyzed (Table 3).

The coefficient of determination of the pathway analysis (R^2), which measures the fraction of total variation explained by the variables under study, was estimated at 68% in the first harvest and 55% in the second harvest (Table 3). In the second year the residual effect exceed the coefficient of determination. According to Cruz et al. (2014), the values observed allow to interpret the cause and effect relations since a representative fraction of the variability is explained by the traits of the diagram.

In evaluation of the direct effects of the plant traits on the main variable (YLD), the highest magnitude was found for NPLAG that was the most important for hulled coffee yield, followed in smaller proportion by NROS in the first and second year of evaluation (Table 3). Positive associations between development of plagiotropic branching and hulled coffee yield were also observed by Dalcomo et al. (2017) in *C. canephora*.

Different associations can facilitate or hinder the selection process. The effects of DROS were negative in relation to the FROS trait and positive in relation to SIEV, since a lower distance between rosettes was associated with a lower number of fruits per rosette and with a lower mean sieve size. Cilas et al. (2006) observed that a higher number of fruits per rosette were also associated with a greater distance between them.

The small direct effect of the PHt trait on yield indicates the possibility of selecting plants of smaller size and of superior yield (Table 3). On the other hand, Bikila and Sakiyama (2017) observed a positive association between plant height and yield. This association was not observed in the germplasm evaluated, which has genotypes of the Conilon botanical variety that stand out for lower height and good yield potential. Rocha et al. (2014) also observed a positive correlation between plant height and distance between rosettes.

1st measurement - 2013-2014Indirect effect..... Direct Total Trait Effect effects SIEV PHt LPLAG **NPLAG** NROS DROS FROS **NDAYS**Hulled coffee yield..... -0.009 0.131 0.003 -0.052 -0.062 0.001 -0.001 0.004 0.015 PHt NPLAG 0.614 -0.002 0.002 0.023 0.068 0.010 -0.007 -0.018 0.691 NROS -0.002 0.133 -0.028 -0.003 -0.001 -0.006 -0.009 0.336 0.252 LPLAG -0.165 -0.003 -0.087 0.002 -0.086 -0.007 0.005 0.104 -0.237 DROS -0.011 -0.002 -0.105 0.001 -0.107 -0.057 -0.208 0.003 0.250 FROS 0.031 -0.001 0.066 -0.001 0.352 -0.003 -0.029 0.213 -0.202 NDAYS 0.039 0.002 -0.106 -0.001 -0.023 -0.019 -0.003 0.041 -0.072 SIEV 0.103 -0.109 -0.036 -0.009 -0.053 -0.001 -0.001 -0.016 0.015

TABLE 3 - Estimates of the direct and indirect effects of the secondary traits of number of days to fruit ripening
(NDAYS), number of plagiotropic branches (NPLAG), number of rosettes (NROS), length of plagiotropic branches
(LPLAG), number of fruits per rosette (FROS), distance between rosettes (DROS), and mean sieve size (SIEV) on
the primary trait of hulled coffee yield (YLD). Bold print highlights the associations of greatest relevance.

	2nd measurement - 2014-2015										
Trait	Direct effect	Indirect effect									
		PHt	NPLAG	NROS	LPLAG	DROS	FROS	NDAYS	SIEV	effects	
• • • • • • • • •			Hulled coffee yield								
PHt	0.107		0.013	0.044	0.008	-0.046	-0.035	0.001	0.023	0.115	
NPLAG	0.424	0.003		0.133	-0.004	0.041	0.007	-0.003	-0.021	0.581	
NROS	0.293	0.016	0.193		0.003	0.028	-0.006	-0.001	-0.004	0.521	
LPLAG	0.018	0.047	-0.092	0.056		-0.052	-0.028	0.004	0.018	-0.029	
DROS	-0.125	0.039	-0.139	-0.065	0.007		-0.037	0.003	0.265	-0.005	
FROS	-0.111	0.034	-0.025	0.169	0.005	-0.342		0.004	0.021	-0.245	
NDAYS	0.017	0.001	-0.070	-0.024	0.004	-0.020	-0.028		0.026	-0.093	
SIEV	0.098	0.025	-0.093	-0.012	0.003	-0.021	-0.024	0.005		-0.019	
R ²	0.55										
Residual effect	0.58										

. . .

2014 2015

R^2 : Coefficient of determination

 \mathbb{R}^2

Residual

effect

0.68

0.49

Yield potential is related to the indirect effects of NROS and LPLAG on hulled coffee yield (Table 3). Although mean sieve size (SIEV) is important for better beverage quality, the small direct and indirect effects of this trait on yield indicate that hulled coffee yield was more related to the capacity of the plant to produce fruits than to the variability of coffee bean size.

These results reveal that it is possible to select plants that aggregate a set of favorable

traits associated with higher hulled coffee yield, such as higher number of productive branches, higher number of rosettes per branch, and lower distance between rosettes. The complementarity of the Conilon and Robusta traits favors selection of hybrid genotypes that express the best traits of each breeding population.

4 CONCLUSIONS

Although the effect of the genotype \times year interaction was not significant for only one trait, the repeatability estimates indicate predominance of simple interaction over the years. The establishment of mutually exclusive groups, ordered according to the variability of the traits, subsidizes the establishment of a scale to evaluate the variability of this genetic resource. Pathway analysis indicated that the number of plagiotropic branches and the number of rosettes per productive branch were the traits that exhibited the greatest direct effect on hulled coffee yield.

5 ACKNOWLEDGEMENTS

The authors thank to the Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq) for the scholarship and research grant. We also gratefully acknowledge the Consórcio Pesquisa Café for financial support.

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