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Prediction of selection gains in Coffee canephore based on factorial scores

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ABSTRACT - The technique of factor analysis in the simultaneous selection of traits and prediction of genetic gains was evaluated in Coffea canephora var. conilon. Fourteen traits in 40 assessed genotypes were evaluated at two sites. The technique was used aiming at the structuring and simplification of the data, without information loss and with biological interpretation. The experimental design was of randomized blocks in four replications, each plot containing two useful plants. The technique was efficient for the data simplification and structuring. Moreover, the estimates of the predicted gains in the traits involved in the factors showed magnitude near the direct selection gain, attesting the suitability of the technique and its use in improvement programs of the species.

Key words: coffee, genetics and plant improvement, multivariate analysis.

INTRODUCTION

Coffea canephora is a species that has been commercially exploited in Brazil since the 50s, when soluble coffee came up. The species was used in blends of roasted and ground coffee with *Coffea arabica* (Malta 1986) and gives these a greater industrial output with lower sales prices, and consequently a higher competitiveness on the market.

The programs of genetic improvement of *C. canephora* have focused especially on achieving productive clonal or

synthetic varieties with enhanced beverage quality (Charrier and Berthaud 1988, Ferrão et al. 2000).

According to Cruz and Carneiro (2003), a way to achieve a genetic superior material in improvement programs, is to deal, simultaneously, with a set of favorable attributes that lead to a comparatively higher productivity, meeting all requirements of consumers as well as producers. The authors mention that selection based on one or few traits has proven inadequate.

C. canephora breeding programs with the objective of simultaneously selecting several attributes require special

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attention, since many of the analyzed traits are highly correlated and can therefore obtain indirect favorable gains with positive and negative correlations, according to the desired objective. However, undesirable indirect gains are also possible, causing serious practical problems, owing to the inconvenient positive and negative correlations. The high magnitudes of these correlations can cause difficulties at obtaining reliable estimates of genetic parameters, due to the multicolinearity.

The breeding strategies for the crop must be reasonably applied, using adequate techniques. Factor analysis is, according to Cruz and Carneiro (2003), a multivariate technique that aims at structuring and simplifying the original data, so that a relatively large number of traits can be represented by fewer, conserving a maximum of information. So theoretically, the technique can be used to minimize the problem of the practice of simultaneous selection in *Coffea canephora*.

This study aimed at evaluating the possibility of using factor analysis to help solve the problem of selection with several attributes in *Coffea canephora* var. *Conilon* improvement programs predicting selection gains, assessing 14 traits in 40 genotypes and in two environments.

MATERIAL AND METHODS

Forty *Coffea canephora* var. *Conilon* genotypes were studied, which are part of the improvement program of the INCAPER of the Espírito Santo (ES) State. Thirty-five of the clones had their origin in the phenotypic selection of mother plants from agricultural properties in several counties across the northern region of the ES; three were elite clones (ES 01 classified as of early maturation, ES 23 of intermediate, and ES 36 of late maturation); a cultivar of the seminal population type from the INCAPER improvement program, and a polyclonal commercial cultivar.

Data from two experiments conducted on experimental farms of the INCAPER, in Marilândia and Sooretama, counties whose soil and climate are representative of the region in the north of the state, where the species is intensely cultivated, were collected in the harvests of 1996, 1998, 1999, 2000, and 2001 and analyzed.

In the county of Marilândia (19⁰ 24' lat S, 40⁰ 31' long W, and altitude 150 m asl) the soil of low fertility is prevailingly crystalline; the annual rainfall precipitation 1200 mm; the mean annual temperature 23 0 C; relative humidity 80%; and the topography is strongly undulated. The county of Sooretama (15⁰ 47' lat S, 43⁰ 18' long W, and altitude 40 m asl) has a sandy soil (80% sand), yellow red dystrophic latosol of low fertility; with an annual rainfall precipitation of 1100 mm; mean annual temperature of 24 0 C; relative humidity of 80%; and the topography is plain with predominant south wind.

The soil was fertilized according to a previous soil analysis for a target yield of 80 sacks of processed coffee per ha (340 kg ha⁻¹ N, 55 kg ha⁻¹ P₂O₅, and 170 kg ha⁻¹ K₂O) in three applications. The management and cultural treatments during the experiments were carried out following the requirements and technical recommendations for the crop, without irrigation.

The experiment was established in randomized complete blocks in four replications, each plot represented by two plants in a 3.0 x 1.5 m spacing and plant density of 2222 plants ha⁻¹. The treatments were evaluated in five samples 24, 48, 60, 72, and 84 months after planting. The following traits were assessed: 1) cycle - number of days between the main flowering and harvest; 2) productivity - processed bean yield in kg ha⁻¹, with 14% humidity; 3) yield in kg of cherry/fruit dried (CfCo); 4) yield in kg of cherry fruit/processed beans (CfPb); 5) yield in kg dried cherry/processed beans (CoPb); 6) percentage of fruits with peaberry beans; 7) percentage of flat beans; 8) percentage of peaberry beans; 9) percentage of humidity; 10) percentage of beans retained in a screen coarser than 15 (S 17); 11) percentage of beans retained in a 15 screen (S 15); 12) percentage of beans retained in a 13 screen (S 13); 13) percentage of beans retained in a 11 screen (S 11); and 14) screen mean (Sm).

The applied factor analysis model was the one proposed by Cruz and Carneiro (2003):

$$X_{j} = I_{j1}F_{1} + I_{j2}F_{2} + ... + I_{jm}F_{m} + \varepsilon_{j}$$

where

 X_i is the j^{th} variable with j = 1, 2, ..., v;

- I_{jk} is the factorial loading for the j^{th} variable associated to the k^{th} factor, and k = 1, 2, ..., m
- F_k is the k^{th} common factor and
- ε_i the specific factor of the j^{th} variable.

Similarly to Abreu et al. (1999), an initial definition of a number of common factors equal to the number of eigenvalues that are higher than the unit and the model of the orthogonal, i.e., not correlated factors were chosen.

After establishing the number of common factors, the communalities (ζ_j^2) were calculated by the expression: $\zeta_j^2 = \sum_{k=1}^m l_{jk}^2 = 1 - \sigma_{ej}^2$, where σ_{ej}^2 is the variance of the specific factor associated to the *j*th variable, and the initial factor loadings of these factors. After the orthogonal rotation based on the varimax algorithm (Mardia et al. 1979), the final loadings were obtained. Each factor was interpreted and the factorial scores obtained by the equation: $F^* = [(G G G)^{-1} G'X]$



 $F^* = 1 \text{ x m}$ vector of dimension of the factorial scores; G = p x m matrix of dimension of the final rotated factor loadings; $X = p \times 1$ vector of dimension of the trait means of the kth genotype. Hence, for each genotype, there will be m new traits, which will correspond to different factors.

The score values were considered representative of a new variable or complex, used as selection criterion in order to obtain simultaneous gains in the original variables. Thereafter, the variance of these new complexes was analyzed and the genetic parameters and selection gains estimated for both sites.

To estimate the proportional selection gain of the original variables in consequence of the selection realized on the factorial scores of the complex, the following expression was adopted:

where

 DS_{ji} = selection differential obtained in trait j by the selection realized on the complex i

 H_i^2 = genotypic determination coefficient of trait j;

 X_i genotypic mean of trait j.

The statistical analyses were run on the software Genes (Cruz 1997).

RESULTS AND DISCUSSION

Factor evaluation

Table 1 presents the final factor loadings and the communalities obtained by the factor analysis with data of

the experiments in Marilândia and Sooretama. Based on the final factor loadings, the possibility of establishing representative factors of the biological complex was evaluated. These were determined by the greatest values of these loadings and their signals. Traits with the same target gain in *C. canephora* var. *Conilon* improvement programs must have identical signals and those with opposed objectives opposite signals, and the magnitude and signal of the final factor loadings of the trait yield must be verified. Final factor loadings with values above 0.7 were considered high, and values above 0.64 were adopted for the communality.

Four factors were obtained from the experiment in Marilândia. The first factor (F1) was related to the traits retained in screens S.17, S.15, S.13, S.11. The Sm presented the greatest final factor loadings and signals according to the objectives of coffee improvement programs. This factor was interpreted as the complex called SCREEN. The second factor (F2) presented highest factor loadings with adequate signals for the traits cycle, CfCo, CfPb, and CoPb. It was the most influenced by processing and cycle, which explains the complex designated YIELD1. The third factor (F3) was most linked to the type of bean quality, with the highest final factor loadings for the traits flat and peaberry, which explains the complex called BEAN TYPE. The fourth factor (F4) was related to the malformed fruits, with highest final factor Schading of the trait malformed, which explains the complex denominated EMPTY.

Table 1. Final loadings factor and communality (C) values obtained by the factor analysis of the traits evaluated in genotypes of *Coffea* canephora var. conilon, of the harvests 1996 and 1998 to 2001

		Marilândia				Sooretama				
Variables	Final loading factor				Final loading factor					
	F1	F2	F3	F4	С	F1	F2	F3	F4	С
cycle	-0.1998	0.7959	-0.3015	-0.0404	0.7659	0.3527	0.4679	-0.3013	-0.2110	0.4786
Product.	-0.4733	0.0961	-0.1458	-0.5681	0.5773	0.2334	-0.1359	-0.0094	0.6688	0.5203
CfCo	0.0686	0.8791	0.0428	-0.2331	0.8337	-0.1716	0.0201	0.0942	0.8438	0.7508
CfPb	0.0474	0.9294	-0.0171	0.2146	0.9123	-0.1886	0.9287	-0.0712	0.1728	0.9330
CoPb	0.1161	0.7189	-0.1046	0.5784	0.8757	-0.1115	0.9237	-0.1038	-0.2159	0.9231
Empty	0.0307	0.0364	0.1145	0.8861	0.8005	0.0840	0.8168	0.1031	-0.2510	0.7479
Flat	-0.2148	-0.1079	0.9469	0.0743	0.9599	0.2301	-0.1219	0.9411	0.0422	0.9553
peaberry	0.1879	0.0856	-0.9559	-0.0675	0.9609	-0.2271	0.1174	-0.9415	-0.0446	0.9538
moisture	-0.3886	0.1457	-0.1721	0.2884	0.2850	-0.2227	-0.6257	0.2453	-0.2002	0.5413
S 17	-0.8186	-0.1713	0.2543	0.0030	0.7641	0.8970	0.0198	0.1841	-0.0178	0.8393
S 15	-0.8926	0.1010	0.0347	-0.1249	0.8237	0.9550	0.0142	0.0760	0.0574	0.9212
S 13	0.8985	0.1210	-0.1728	-0.0862	0.8593	-0.8577	0.0117	0.0469	-0.0081	0.7381
S 11	0.8777	-0.1070	-0.1020	0.2881	0.8752	-0.8534	-0.0391	-0.2380	-0.0380	0.7880
Sm	-0.9619	-0.0396	0.1895	-0.1118	0.9752	0.9688	0.0329	0.1746	0.0161	0.9704

*Values in bold refer to the loadings factor of the traits that compose the factors and their communalities

Three factors were obtained from the experiment in Sooretama. The first factor (F1) was related to the traits bean size. This factor explains the complex called SCREEN, as in Marilândia. The second factor (F2) was more related to the yield, presenting highest final factor loadings and appropriate signals for the traits CfPb, CoPb, and malformed, so this factor explains the complex designated YIELD2. The third factor (F3) was more influenced by the quality, since it presented highest final factor loadings and adequate signals for the traits flat and mocha, which explains the complex BEAN TYPE. In this experiment, no fourth factor (F4) was obtained, since the trait CfCo presented a factorial loading of 0.84 and a high communality of 0.75, but the signal of the final factorial loading was identical to that of the productivity. This last trait had a relative high final factorial loading (0.66), which was inadequate for the goal of a Coffea canephora improvement program, where the ratio CfCo is being diminished and the productivity increased, which would not occur if this factor were used. It was therefore not considered a factor of biological interest.

The productivity trait was not included in complexes, suggesting the absence of correlation between this trait and the factors, as demonstrated by their final factor loadings below 0.7 and communality lower than 0.64.

Variance analysis and genetic parameters

Significant values (P < 0.01) of the genotype effects were verified by the F test for all complexes and for productivity at both sites, which would give proof of the existence of genetic variability, required for selection, making the identification of superior genetic material possible (Table 2).

There were differences between the estimates of the experimental precision (VC%) (Table 2) of the evaluated complexes. This indicates a weaker influence of the environment on YIELD1 in Marilândia and, on bean type in Sooretama, which also indicates a weaker effect of the environment on these traits. There was a stronger influence for EMPTY in Marilândia and for SCREEN in Sooretama.

Table 3 presents the phenotypic, environmental, and genotypic variances, the genotypic determination coefficient among progenies, the percentage of the genotypic variation coefficient (VCg), and the ratio of the genetic and environmental VC (VC_g/VC_e) in Marilândia and Sooretama. In Marilândia, these variation values were higher than 1.04 and lower than 2.57, and in Sooretama, they were higher than 1.54 and lower than 5.10. According to Vencovsky (1987), values of this index equal to or higher than 1.0 indicate a favorable situation for selection. The complexes as well as the productivity presented a genotypic determination coefficient (H^2) equal to or higher than 81.29%, attaining 96.36% for SCREEN in Marilândia. In Sooretama, all complexes and the productivity presented a genotypic determination coefficient (H^2) equal to or higher than 90.50%, attaining 99.05%, for SCREEN. The predominance of the genetic variability in relation to the environmental was characterized at both sites, for all complexes and for

Table 2. Summary of the variance analyses, general mean and VC% of the evaluated traits in genotypes of *Coffea canephora* var. *conilon*, for the harvests 1996 and 1998 to 2001

	Mean squares of the variation sources							
** • • •	Blocks ¹	Genotypes ¹	Error ¹	General	VC			
Variables	(3)	(39)	(117)	Mean	%			
			Marilândia					
productivity	93247.85	1663294.88**	178025.76	2990.34	14.11			
SCREEN	0.27	4.00**	0.15	-5.29	7.22			
YIELD1	0.40	4.00**	0.32	21.71	2.60			
BEAN TYPE	1.41	4.00**	0.35	7.35	8.10			
EMPTY	3.29	4.00**	0.75	1.06	81.33			
			Sooretama					
productivity	131398.65	1040166.58**	62825.91	2393.37	10.47			
SCREEN	0.04	4.00**	0.04	0.40	48.16			
YIELD2	1.56	4.00**	0.38	7.36	8.37			
BEAN TYPE	0.24	4.00**	0.27	11.13	4.63			

** Significant at P<0.01 by the F test; productivity: bean yield in five harvests (kg ha⁻¹); SCREEN: S 17, S 15, S 13, S 11, and Sm; YIELD1: ratios CfCo, CfPb, CoPb, and cycle; YIELD2: ratios CfPb, CoPb, and malformed; BEAN TYPE: percentage of flat and peaberry beans; EMPTY: percentage of malformed beans. ¹value in brackets expresses the degree of freedom

	Variances of the variation sources of genetic-statistical parameters								
Variables	$\hat{\sigma}_{f}^{2}$	σ ²	$\hat{\sigma}_{g}^{2}$	H^2	VC _g	VC_g/VC_e			
				%	%				
			Marilândi	a					
productivity	415823.72	44506.44	371317.28	89.30	20.38	1.44			
SCREEN	1.00	0.04	0.96	96.36	18.56	2.57			
YIELD1	1.00	0.08	0.92	92.02	4.42	1.70			
BEAN TYPE	1.00	0.09	0.91	91.15	12.99	1.60			
EMPTY	1.00	0.19	0.81	81.29	84.75	1.04			
			Sooretam	a					
productivity	260041.64	15706.48	244335.17	93.96	20.65	1.97			
SCREEN	1.00	0.01	0.99	99.05	245.74	5.10			
YIELD2	1.00	0.09	0.91	90.50	12.92	1.54			
BEAN TYPE	1.00	0.07	0.93	93.37	8.68	1.88			

Table 3. Estimates of the genetic and environmental parameters of the complexes, established by the factor analysis, based on the linear combinations of the traits evaluated in *Coffea canephora* var. *conilon*, genotypes of the harvests of 1996 and 1998 to 2001 in Marilândia and Sooretama

productivity: bean yields in five harvests (kg ha⁻¹); SCREEN: S 17, S 15, S 13, S 11 and Sm; YIELD1: ratios CfCo, CfPb, CoPb, and cycle; YIELD2: ratios CfPb, CoPb, and malformed; BEAN TYPE: percentage of flat and mocha beans; EMPTY: percentage of malformed beans; $\hat{\sigma}_{f}^{2}$: Mean phenotypic variance; $\hat{\sigma}_{g}^{2}$: mean variance; σ_{g}^{2} : mean genotypic variance; H^{2} : Herdability; $VC_{g}(\%)$: coefficient of the genetic variation; VC_{e} : environmental variation coefficient

productivity. However, the results suggested a favorable condition for improvement, because H^2 is a relative parameter of reliability by which the phenotypic value represents the genotypic value.

Table 4 displays the values of the original mean, selected mean, selection differential, genotypic determination coefficient, and estimates of the predicted gain by direct selection. All factors provided an adequate combination of expected gains in the considered traits, as is desirable in improvement programs of the species according to Ferrão et. al (1999) and Fonseca et al. (2001), as for example increase in the traits S 17 and S 15, decrease in the percentage of peaberry beans, and increase in the percentage of flat beans (Tables 5 and 6). The direct gains for the traits were higher than the

Table 4. Estimates of the predicted gains with direct selection (SG), genotypic determination coefficient (H^2), original mean (X_0), selection mean (X_s), and differential selection (DS) in the traits of *Coffea canephora* var. *conilon* genotypes, in the harvests 1996, 1998, 1999, 2000, and 2001 in Marilândia and Sooretama

Variables	\mathbf{X}_{0}	X _s	DS	\mathbf{H}^2	SG	SG
				<u> </u>		%
			Marilândia			
productivity	2990.34	3759.26	768.92	89.30	686.62	22.96
SCREEN	-5.29	-6.44	-1.15	96.36	-1.10	20.87
YIELD1	21.71	20.48	-1.23	92.02	-1.13	-5.22
BEAN TYPE	7.35	8.41	1.06	91.15	0.97	13.15
EMPTY	1.06	-0.23	-1.29	81.29	-1.05	-98.54
			Sooretama			
productivity	2393.37	3040.70	647.33	93.96	608.24	25.41
SCREEN	0.40	1.62	1.20	99.05	1.20	296.09
YIELD2	7.36	6.32	-1.04	90.50	-0.94	-12.82
BEAN TYPE	11.13	12.25	1.12	93.37	1.04	9.38

SG (%): selection gain based on the selection differential; productivity: beans yields in five harvests (Kg ha⁻¹); SCREEN: S 17, S 15, S 13, S 11, and Sm; YIELD1: ratios CfCo, CfPb, CoPb, and cycle; YIELD2: ratios CfPb, CoPb, and malformed; BEAN TYPE: percentage of flat and malformed beans; EMPTY: percentage of malformed beans

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Variables	Selection gain via complexes	Direct selection gain in the traits	Variables	Selection gain via complexes	Direct selection gain in the traits
SCREEN	r i i i i i i i i i i i i i i i i i i i				
S 17	105.23	121.55	17	116 19	130 36
S 15	22.96	30.28		110.15	130.30
S 13	-37.1	-42.68	15	37.59	38.26
S 11	-58.7	-60.49	13	29.79	30.56
Sm	5.70	6.11	.11	54.96	59.52
YIELD1				6 12	6 30
cycle	-5.29	-6.61	_	0.12	0.00
CfCo	-3.86	-4.12	2		
CfPb	-7.91	-8.78		7.08	7.53
CoPb	-3.31	-5.10		7.18	7.68
BEAN TYPE				40.84	52.82
flat	4.89	5.18			
peaberry	-22.59	-24.00			
EMPTY				4.90	4.99
empty	-28.48	-31.81		21.69	22.08

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Table 5. Percentage of predicted gains in Marilândia for the traits that compose the complexes SCREEN, YIELD1, BEAN TYPE, and EMPTY by the selection via complexes and direct selection in the traits

gains by the complexes. However, direct selection did not obtain adequate simultaneous gains for any of the traits in the other involved traits, which was the case with selection via complexes.

Based on the obtained results, the simplification and structuring of the original data of *Coffea canephora* were obtained in the two experiments by the factor analysis. In Marilândia 12 traits were represented by four complexes, and in Sooretama 10 traits were represented by three complexes. The factors preserved a maximum of information of the traits in the experiments, made quite clear by the estimates of the predicted gains obtained by the complexes, which were close to the prediction of the direct selection gains in the traits.

The analyses of the two experiments demonstrated that the use of the factor analysis technique was adequate and there was evidence that it could be applied in Coffea *canephora* improvement programs.

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Predição de ganhos genéticos em *Coffea canephora* com base em escores fatoriais

RESUMO - Avaliou-se a utilização da técnica de análise de fatores na seleção simultânea de caracteres e predição de ganhos genéticos em Coffea canephora var. conilon. Foram avaliados 14 caracteres em 40 genótipos ensaiados em dois locais. A técnica foi utilizada visando a estruturação e a simplificação dos dados, sem perda de informação e com interpretação biológica. Os dados analisados pertencem ao programa de melhoramento do INCAPER/ES. O delineamento experimental utilizado foi o de blocos casualizados com quatro repetições, cada parcela composta por duas plantas úteis. A técnica foi eficiente na simplificação dos dados. As estimativas de ganhos preditos nas características envolvidas nos fatores apresentaram magnitudes próximas ao ganho com seleção direta, atestando o potencial de uso da técnica em programas de melhoramento da espécie.

Palavras-chave: café, genética e melhoramento de plantas, análise multivariada.

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