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Temporal Stability for Unpredictable Annual Climatic Variability for Hevea Genotype Selection

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ABSTRACT

The objective of this study was to assess genotype-year interaction and determine temporal stable genotypes across six years of rubber yield evaluation. Stability analyses were performed by Eberhart and Russell method for rubber yield. Twenty-five genotypes were analyzed in a randomized complete block design with three replications. The best genotype in one year was not same always in the other year. The genotype components were partitioned into linear (genotypes within year) and nonlinear (pooled deviations) components. Significant mean square for linear components was predictable. This indicated that the performance of genotypes across the years for rubber yield could be predicted. Among the analyzed genotypes the IAC 40 also was considered highly productive and vigorous, with suitable adaptation.

Key words: *Hevea brasiliensis*, genotype x year interaction, rubber tree, rubber yield, stability parameters

INTRODUCTION

In rubber tree [*Hevea brasiliensis* (Willd. ex Adr. de Juss.) Muell.-Arg.], productivity performance consists mainly of girth growth and yield stability. Breeders search for genotypes that show a stable, high yield across the years and locations. Results of genotype evaluation trials can be used to select and recommend a clone adapted to a particular agro-climatic condition if the agro-climatic differences can be defined according to environmental factors (Gonçalves et al., 2003). No extensive studies have been performed on the temporal stability for rubber yield. In general, a large amount of variation across years in rubber yield is observed.

Genotype-year interaction in perennial crops such as rubber tree represents the differential response of genotypes to changing annual conditions. In the presence of genotype x year interaction, the relative performance of genotypes varies with the year.

Several methods have been proposed to analyze genotype x environment interactions of phenotypic stability (Lin et al., 1986; Becker and Leon, 1988; Piepho, 1998;Truberg and Huehn, 2000). The most widely used method to analyze phenotypic stability is to calculate a regression of the yield of a given genotype in different environments on the mean of all tested genotypes (Okuyama, et al., 2005). The coefficients of regression ($\hat{\beta}_i$) can be used to describe the general response to

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environmental conditions (Stringfield and Salter, 1934) while the mean squares deviations $(\hat{S}_{d_i}^2)$ from linear regression actually measure yield stability (Yates and Cochran, 1938; Eberhart and Russell, 1966). Kalil Filho (1982) was the first one to employ the methodology of Eberhart and Russell (1966) to study the temporal yield variation in 25 clones. Later, Gonçalves et al. (1992), using the same stability method, examined the magnitude of the genotype x environment interaction for rubber yield of 19 *Hevea* mother trees at the same age.

The objectives of this study were to assess genotype-year interaction and to determine temporally stable genotypes across six annual yield evaluations for future recommendations.

MATERIAL AND METHODS

One Malaysian (RRIM), eight Amazonic (IAN, Fx, RO) and 16 local (IAC) genotypes (clones) were used in this study (Table 1). The Amazonic clones comprised Fx 985 and Fx 3899 developed by Ford Motor Company by crossing primary selections of South American leaf blight (SALB) resistant materials with high yielding oriental clones to produce Fx clones. The IAN clones (IAN 3156, IAN 3703, IAN 4493, IAN 6323 and IAN 6721) were developed by the breeding program of the former Instituto Agronômico do Norte, nowadays Embrapa Western Amazon. One-yearold rootstock seedlings raised in nurseries were used to budgraft the genotypes (clone materials). Budgrafting was made in December 1988. The successful budgrafts were uprooted and planted in the polyethylene bags. After the first flush of leaves developed, the plants were established in the field.

The experiment was conductive at the Votuporanga Experimental Station located in Northwestern region of São Paulo State (Brazil) at 20°25'S latitude, 49°50'W longitude and 450m elevation. Annual mean temperatures varied from 22.3°C to 24.1°C. Annual rainfall ranged from 1,480 to 1,600mm. Winter drought varied from three to four months, with an average rain deficit of 180mm. The experiment was laid out in a randomized complete block design with three replicates, using eight trees per plot and 7.0 m x 3.0 m spacing. Missing plants were replaced with spares during the first two years after planting to

maintain plantation density, but they were not scored.

One row of the commercial clone RRIM 600, acquired from a commercial nursery, was planted around the plot. Annual fertilizations consisted of 400 g of 10-10-10 NPK formula per plant, according to Bataglia and Gonçalves (2003). At the end of the sixth year, all the trees that reached a girth of 45.0 cm or more were opened for tapping at a height of 1.20 m above the highest point of the bud union, for annual rubber yield. Attempts were made to record six annual rubber yields. The latex extracted from the panel followed a half-spiral four-daily tapping system (seven tapping/month) for 11 months/year. Yield was recorded on the days when normal tapping, which started around 7:30 AM, was possible. After tapping, latex was allowed to collect in plastic cups provided for each useable tree. Upon stoppage of the latex flow, rubber was coagulated in the cup itself by adding 2% of acetic acid solution and stirring it well. The coagulated rubber in each cup was then made into a "biscuit," which was dried, hung by a wire tied in each tree for about 30 days, then weighed and the dry rubber content for each tree was recorded.

All the analyses were performed using "Genes" computer program, Windows version, 2001 (Cruz, 2001). The method of Eberhart and Russell (1966) was used to characterize genotypic stability in different years. For the temporal stability study, various environments were represented by the six different annual rubber yields through an environmental index, i.e., mean performance of all the selections in each year. The following linear model was used:

 $Y_{ij} = m + \beta_i I_j + \delta_{ij} + \overline{E}_{ij},$

where Y_{ij} is the mean of the genotype *i*th at the year *j*; *m* is general mean of all *i* genotypes over all years; β_i is the regression coefficient of the *i*th genotype on the annual index which measures the response of this genotype to varying years; I_j is the annual index which is defined as the mean deviation of all genotypes at a given year from the overall mean ; δ_{ij} is the deviation from regression of the *i*th genotype in the *j*th year and \overline{E}_{ij} is the mean of experimental error.

Two stability parameters were calculated: (a) the regression coefficient, $\hat{\beta}_i$, which is the regression

of the performance of each genotype under different years, means across all the genotypes; (b) mean square deviation $(\hat{S}_{d_i}^2)$ from linear regression for each genotype. They were estimated according to Cruz and Carneiro (2003) as follows: a) Computation of regression coefficient $(\hat{\beta}_i)$ for each genotype

$$\hat{\beta}_i = \sum_j Y_{ij} I_j / \sum_j I_j^2$$

where $\sum_{j} Y_{ij} I_{j}$ is the sum of products of environmental index (I_{j}) with the corresponding mean (m) of the genotype in each year. These values were obtained by the following manner:

 $[m] [I_j] = [\sum_j Y_{ij}I_j] = [S]$, where: [m] is the matrix of means; $[I_j]$ is the vector for environmental index and [S] is the vector for sum of products, i.e. $\sum_j Y_{ij}I_j$; $\sum_j I_j^2$ is the sum of environmental index squares for each genotype; I_j is the environmental index defined as the deviation of the mean of all genotypes at a year from the overall mean:

$$I_{j} = \frac{\sum_{i} Y_{ij}}{t} - \frac{\sum_{i} \sum_{j} Y_{ij}}{ta}, \text{ where: } \sum_{i} Y_{ij} \text{ is the total of all}$$

the genotypes at j^{th} year; $\sum_{i} \sum_{j} Y_{ij}$ is the

grand total; t = number of genotypes and a = number of years.

b) Computation of mean square deviations $(\overline{S}_{d_i}^2)$ from linear regression for each genotype

$$\overline{S}_{d_i}^2 = \frac{\sum_j \delta_{ij}^2}{(s-2)} - \frac{S_e^2}{r} \text{ where } \sum_j \delta_{ij}^2 \text{ is the variance}$$

due to deviations from regression:

$$\sum_{j} \delta_{ij}^{2} = \left[\sum_{j} Y_{ij}^{2} - \frac{Y_{i.}^{2}}{t} \right] - \frac{\left(\sum_{j} Y_{ij} I_{j} \right)^{2}}{\sum_{j} I_{j}^{2}}, \text{ where:}$$

$$\sum_{j} Y_{ij}^{2} - \frac{Y_{i.}^{2}}{t} \text{ is the variance due to dependent}$$
variable; $\left(\sum_{j} Y_{ij} I_{j} \right)^{2} / \left(\sum_{j} I_{i}^{2} \right)$ is the variance due to regression and S_{e}^{2} is the mean square for pooled error.

Table 1 - Parentals and annual means for 25 genotypes (clones) of rubber yield in grams/tree/tapping evaluated over six years in Votuporanga Experimental Station, Northwestern region of São Paulo State, Brazil.

| Genotypes | notypes Parentals | | 1 st year | | 2 nd year | | 3 nd year | | 4 th year | | 5 th year | | 6 th year | |
|-----------|--------------------|------------|----------------------|-------------|----------------------|-------------|----------------------|-------------|----------------------|-------------|----------------------|-------------|----------------------|--|
| | | Mean | Rank | Mean | Rank | Mean | Rank | Mean | Rank | Mean | Rank | Mean | Rank | |
| Fx 985 | F 315 x AVROS 183 | 17.87 | 7 | 42.61 | 9 | 35.27 | 17 | 34.84 | 23 | 41.09 | 14 | 46.63 | 14 | |
| Fx 3899 | F 4542 x AVROS 363 | 18.63 | 6 | 42.91 | 8 | 51.75 | 8 | 48.32 | 13 | 33.04 | 23 | 35.25 | 22 | |
| IAC 40 | RRIM 608 x AVROS | 25.61 | 2 | 69.97 | 1 | 72.69 | 2 | 76.09 | 2 | 75.27 | 1 | 80.80 | 2 | |
| IAC 56 | RRIM 608 x F 3810 | 15.33 | 9 | 50.69 | 6 | 70.93 | 4 | 78.32 | 1 | 54.85 | 3 | 56.28 | 7 | |
| IAC 300 | RRIM 605 x AVROS | 22.32 | 3 | 62.42 | 2 | 53.56 | 7 | 66.36 | 4 | 49.93 | 8 | 61.73 | 4 | |
| IAC 301 | RRIM 501 X AVROS | 20.52 | 4 | 57.12 | 3 | 77.22 | 1 | 68.30 | 3 | 51.63 | 5 | 56.56 | 5 | |
| IAC 302 | RRIM 501 x AVROS | 19.13 | 5 | 37.99 | 11 | 41.08 | 13 | 47.55 | 14 | 49.26 | 9 | 53.64 | 9 | |
| IAC 303 | RRIM 511 x AVROS | 15.05 | 10 | 41.37 | 10 | 48.27 | 11 | 61.27 | 7 | 63.33 | 2 | 86.26 | 1 | |
| IAC 306 | AVROS 49 x RRIM | 14.80 | 12 | 33.27 | 17 | 34.55 | 18 | 36.22 | 22 | 26.77 | 25 | 26.08 | 25 | |
| IAC 307 | AVROS 1328 x PR | 13.77 | 13 | 36.97 | 13 | 47.07 | 12 | 53.28 | 10 | 50.35 | 7 | 54.33 | 8 | |
| IAC 308 | AVROS 49 x PR 107 | 12.19 | 17 | 35.65 | 14 | 38.27 | 15 | 53.13 | 11 | 50.62 | 6 | 48.75 | 12 | |
| IAC 309 | RRIM 626 x Fx 25 | 11.64 | 18 | 33.28 | 16 | 31.96 | 21 | 40.85 | 17 | 39.73 | 15 | 48.83 | 11 | |
| IAC 310 | AVROS 1328 x PB 86 | 12.50 | 16 | 31.39 | 19 | 33.08 | 20 | 36.87 | 20 | 36.29 | 19 | 39.00 | 20 | |
| IAC 311 | AVROS 509 x Fx 25 | 8.99 | 24 | 27.81 | 21 | 28.03 | 24 | 40.38 | 18 | 38.86 | 17 | 46.15 | 15 | |
| IAC 312 | RRIM 600 x Fx 25 | 10.64 | 19 | 29.48 | 20 | 23.62 | 25 | 32.47 | 25 | 36.29 | 20 | 41.53 | 18 | |
| IAC 313 | RRIM 626 x FX 25 | 10.53 | 20 | 34.17 | 15 | 48.56 | 10 | 45.78 | 15 | 46.80 | 10 | 50.78 | 10 | |
| IAC 314 | AVROS 1328 x Fx 25 | 9.48 | 23 | 25.63 | 23 | 30.08 | 22 | 34.70 | 24 | 30.55 | 24 | 28.63 | 24 | |
| IAC 316 | AVROS 1328 x RRIM | 13.50 | 14 | 25.78 | 22 | 28.43 | 23 | 36.24 | 21 | 37.34 | 18 | 44.22 | 16 | |
| IAN 3156 | IAN 4493 x PB 86 | 38.67 | 1 | 54.52 | 5 | 56.63 | 5 | 65.93 | 5 | 44.63 | 12 | 31.94 | 23 | |
| IAN 3703 | Fx 4371 x PB 86 | 9.51 | 22 | 37.70 | 12 | 51.31 | 9 | 57.21 | 9 | 39.28 | 16 | 36.79 | 21 | |
| IAN 4493 | Fx 4421 x Tjir 1 | 13.35 | 15 | 31.77 | 18 | 35.72 | 16 | 43.31 | 16 | 34.00 | 22 | 41.25 | 19 | |
| IAN 6323 | Tjir 1 x Fx 3810 | 9.97 | 21 | 25.15 | 24 | 34.25 | 19 | 39.50 | 19 | 34.28 | 21 | 48.67 | 13 | |
| IAN 6721 | Fx 43-655 x PB 86 | 6.93 | 25 | 23.83 | 25 | 39.25 | 14 | 48.38 | 12 | 42.38 | 13 | 41.74 | 17 | |
| RO 45 | Primary clone | 14.92 | 11 | 56.09 | 4 | 71.62 | 3 | 65.00 | 6 | 44.78 | 11 | 56.55 | 6 | |
| RRIM 600 | Tjir 1 x PB 86 | 17.26 | 8 | 48.08 | 7 | 56.25 | 6 | 58.43 | 8 | 51.90 | 4 | 68.47 | 3 | |
| Overall | | 15.32±6.63 | 3 | 39.83±12.61 | | 45.58±15.38 | | 50.75±13.70 | | 44.13±10.76 | | 49.23±14.48 | 3 | |

RESULTS AND DISCUSSION

The average rubber yields, their ranks and differences among averages for the 25 genotypes tested across six years are presented in Table 1. Except for IAC 40 and IAC 301 genotypes, yield ranks of the genotypes across the years changed. Similarly, a study (Gonçalves et al., 1992) undertaken on rubber yield stability across five years also revealed very high fluctuations in one growing trial of the Experimental Station of Pindorama, of São Paulo State. This showed the difficulties encountered by the breeders in selecting new clones for release. These difficulties arose mainly from the masking effects of variable annual yield. Thus, would be important to study in depth the yield and vigor levels, adaptation patterns and stability of genotypes across several years (Romagosa and Fox, 1993). Pham and Kang (1988) indicated that genotype x environment interactions minimized the usefulness of genotypes by confounding their yield performance. The highest rubber yield of 86.26 dry rubber grams per tree/tapping was obtained from IAC 303 in the sixth year, while the lowest was from IAN 6721 in the first year. Across the years, however, IAC 40 and IAC 301 surpassed all other genotypes with an average rubber yield of 66.74 g and 55.22 g, respectively. Their average rubber yield exceeded that of IAC 314 by about 150%, indicating their high yield and good adaptability to the Heveaproducing years in Votuporanga Experimental Station. Genotype IAC 40 was particularly outstanding, ranking first during two of six years. The sixth year was the most productive with a mean yield of 80.80 g of rubber, nearly 55 g higher than that for the first year.

The partitioning of variance components revealed that unpredictable environment (years) was an important source of variation (Table 2). When the genotype x year interaction was due to variation in predictable environment factors, *Hevea* breeders would have the alternatives of either developing specific genotypes for different environments (locations, soil types, management systems, etc.) or broadly adapted genotypes that could perform well under annual variable conditions. However, when genotype x environment interaction resulted from variation in unpredictable environmental factors, such as year-to-year variation in rainfall distribution, as was the case of this study, the breeder would need to develop stable genotypes that could perform reasonably well under a range of conditions. Such breeding strategies could assist the rubber producer in risk avoidance. Ceccarelli (1994) and Piepho (1998) indicated that producers perceived the yield stability as the most important economic aim to minimize crop failure, especially in marginal environments.

The stability analysis conducted for six-year rubber yield in Votuporanga is presented in Table 2 it revealed that the genotypes (clones) differed significantly. The genotypes x year interaction component were further partitioned into linear (genotypes within years effects) and non-linear (pooled deviations) components. Mean squares for both these components were tested against pooled error mean squares. The linear component was highly significant, indicating that the unpredictable components were shared in the genotype x year interaction. Preponderance of linear genotype x year interaction was of great practical importance, implying that there wose differences among linear regression coefficients for each genotype.

The stability parameters for all genotypes are given in Table 3. Eberhart and Russell (1966) emphasized the need for considering both linear $(\hat{\beta}_i)$ and non-linear (\hat{S}_{di}) components of genotype-environment interactions in judging the stability of a genotype. A highly stable genotype was defined as one with ($\hat{S}_{di} = 0$) and an unstable as one with $\hat{S}_{di} > 1$. In addition, a widely adapted genotype was defined as one with $\hat{\beta}_i = 1$; specifically adapted to favorable environments as one with $\hat{\beta}_i > 1$, and specifically adapted to unfavorable environments as one with $\hat{\beta}_i < 1$. In this study, values for the regression coefficient $(\hat{\beta}_i)$ ranged from 0.3521 (IAN 3156) to 1.5952 (IAC 303) for rubber yield. The regression coefficients of genotypes IAC 302, IAC 307, IAC 310, IAC 313, IAN 4493 and RRIM 600 were non-significantly different from unity ($\hat{\beta}_i = 1.0$) and had a small deviation from regression (\hat{S}_{di}) and thus possessed fair stability.

On the other hand, Eberhart and Russell (1966) stated that genotypes with wide adaptability were those with high mean yield, regression coefficient

equal to unity $(\hat{\beta}_i=1)$ and deviation from regression as small as possible $(\hat{S}_{di}=0)$. Accordingly, IAC 302, IAC 307 and IAC 313 were the most widely adapted genotypes for rubber production, because their regression coefficients were almost equal to the unity $(\hat{\beta}_i=1)$ and they had the lowest deviations from regression and high mean yields.

Table 2 - Mean squares (MS) of the analysis of variance for temporal stability parameter for 25 *Hevea* genotypes (clones) evaluated over six years rubber yield in Votuporanga, Experimental Station in Northwestern region of São Paulo State, Brazil.

| Source of variation | | DF | | MS | |
|-------------------------|-----|-----|-----|--------------|---------|
| Genotypes | 24 | | | 2002.3679** | |
| Years | 5 | | | 12670.7850** | |
| Genotypes x Years | 120 | | | 177.1590** | |
| Genotypes within Years | 125 | | | 676.9040** | |
| Year (linear) | | 1 | | 63353.9654** | |
| Genotype x Year (linear | r) | 24 | | 284.4732** | |
| Pooled deviation | | 100 | | 144.3172 | |
| Fx 38 | 399 | | 4 | 210.4008* | |
| Fx 9 | 985 | | 4 | 118.5464 | |
| IAC | 300 | | 4 | 142.0634 | |
| IAC | 301 | | 4 | 311.1483** | |
| IAC | 302 | | 4 | 60.3465 | |
| IAC | 303 | | 4 | 545.1644** | |
| IAC | 306 | | 4 | 79.1920 | |
| IAC | 307 | | 4 | 31.5173 | |
| IAC | 308 | | 4 | 87.9197 | |
| IAC | 309 | | 4 | 76.9825 | |
| IAC 3 | 310 | | 4 | 10.8315 | |
| IAC 3 | 311 | | 4 | 102.6332 | |
| IAC 3 | 312 | | 4 | 117.9125 | |
| IAC 3 | 313 | | 4 | 50.9312 | |
| IAC 3 | 314 | | 4 | 14.6253 | |
| IAC 3 | 316 | | 4 | 95.3139 | |
| IAC | 40 | | 4 | 52.5829 | |
| IAC | 56 | | 4 | 238.2053 ** | |
| IAN 31 | 156 | | 4 | 515.1784** | |
| IAN 37 | 703 | | 4 | 209.8998** | |
| IAN 44 | 493 | | 4 | 89.2256 | |
| IAN 63 | 323 | | 4 | 102.7960 | |
| IAN 67 | 721 | | 4 | 95.1471 | |
| RO | 45 | | 4 | 290.5618** | |
| RRIM (| 500 | | 4 | 57.7847 | |
| Pooled error | | | 288 | | 65.1700 |

* and ** significant for P<0.05 and P<0.01 respectively

| | Genotypes | \hat{m}_{i} | $\hat{\beta}_{i}$ | $\hat{S}_{d_i}^2$ | ${}^{4}R{}^{2}_{i}\%$ |
|-----|-----------|---------------|-------------------|-------------------|-----------------------|
| 1 | IAC 40 | 66.7372 a | 1.5485** | -4.1957 | 96.6544 |
| 2 | IAC 301 | 55.2256 a-b | 1.3158* | 81.9927** | 77.9023 |
| 3 | IAC 56 | 54.3983 a-c | 1.5661** | 57.6784** | 86.7083 |
| 4 | IAC 300 | 52.7217 a-d | 1.1437 | 25.6311 | 85.3666 |
| 5 | IAC 303 | 52.5911 a-d | 1.5952** | 160.3314** | 74.6954 |
| 6 | R O 45 | 51.4944 a-e | 1.3888** | 75.1372 ** | 80.7889 |
| 7 | RRIM 600 | 50.0683 a-f | 1.3119 | -2.4617 | 94.9672 |
| 8 | IAN 3156 | 48.7189 a-g | 0.3521** | 150.0028** | 13.2301 |
| 9 | IAC 307 | 42.6317 b-h | 1.1681 | -11.2175 | 96.4821 |
| 10 | IAC 302 | 41.4411 b-h | 0.8952 | -1.6078 | 89.3772 |
| 11 | IAC 308 | 39.7683 b-g | 1.1092 | 7.5832 | 89.8646 |
| 12 | IAC 313 | 39.4367 b-h | 1.1419 | -4.7462 | 94.1934 |
| 13 | IAN 3703 | 38.6322 b-h | 1.1446 | 41.3518* | 81.4344 |
| 14 | Fx 3899 | 38.3161 b-h | 0.7265 | 48.4102* | 61.3825 |
| 15 | Fx 985 | 36.3861 b-h | 0.6478* | 17.7921 | 69.1596 |
| 16 | IAC 309 | 34.3839 c-h | 0.9109 | 3.9375 | 87.2256 |
| 17 | IAN 4493 | 33.2333 d-h | 0.8120 | -18.4234 | 97.6848 |
| 18 | IAN 6721 | 31.9506 e-h | 1.0056 | 9.9923 | 87.0699 |
| 19 | IAC 311 | 31.7033 e-h | 0.9378 | 12.4877 | 84.4452 |
| 2 0 | IAC 310 | 31.5206 e-h | 0.7356 | -18.1128 | 96.9370 |
| 21 | IAN 6323 | 31.4172 f-h | 0.9318 | 12.5420 | 84.2536 |
| 22 | IAC 316 | 30.9200 g-h | 0.7342 | 10.0479 | 78.1839 |
| 23 | IAC 312 | 29.0028 g-h | 0.7145 | 17.5808 | 73.2885 |
| 24 | IAC 306 | 28.6161 g-h | 0.4983** | 4.4740 | 66.5147 |
| 25 | IAC 214 | 26 5004 h | 0 6624* | 16 9 4 9 2 | 05 0171 |

Table 3 - Estimates of temporal stability and adaptability parameters for rubber yield in grams/tree/tapping for 25 genotypes evaluated over six years in Votuporanga Experimental Station, Northwestern region of São Paulo State, Brazil.

 ${}^{1}\hat{m}_{i}$ = mean of the genotypes (clones), tested according to Tukey's test (p < 0.05), ${}^{2}\hat{\beta}_{1}$ = mean regression coefficient, * and ** significant for p < 0.05 and p < 0.01 respectively (t-test); ${}^{3}\hat{S}_{d_{1}}^{2}$ = deviation from regression coefficient, * and ** significant for p < 0.05 and p < 0.01 (F-test); ${}^{4}R_{1}^{2}$ = coefficient of determination



Figure 1 - A representation of the genotype regression coefficients ($\hat{\beta}$) plotted against rubber yield means in grams/tree/tapping for 25 *Hevea* clones evaluated across six year in Votuporanga Experimental Station, Northwestern region of São Paulo State, Brazil.

Their coefficient of determination, R^2 , (Pinthus, 1973) were as high as 89.38, 96.48 and 94.19%, their importance. In contrast, confirming genotypes IAC 301, IAC 303, IAC 40, IAC 56 and RO 45 with regression coefficients greater than one, were regarded as adaptable to annual changes. Fig. 1 represents genotype regression coefficients plotted against the rubber yield clonal means. The genotype IAC 40 ($\hat{\beta} = 1.55$) had a regression coefficient greater than unity and was highly vigorous and high yielding, hence it could characterized as suitable for the specific adaptation in favorable and unfavorable years. In addition, even the second place, the IAC 301, being less previsible with significant deviation from regression, should be selected because it presented low rank in all the annual evaluations. The locally adapted cultivars had regression coefficients close to unity, had above average yields and could, therefore, be characterized as well adapted to temporal changes. These genotypes also had smallest deviations from regression and, hence, could be regarded as stable genotypes. The genotypes IAN 6721, IAC 311 and IAC 316 that had regression coefficients smaller than unity and below average rubber yields, indicated that they offered a greater resistance to temporal changes.

The regression coefficient measures the "relative" performance of the genotype. Clair and Kleinschmit (1986) emphasized that in forest tree breeding, this information was useful to distinguish genotypes for specific environments, but if all environments tested was in one planting zone, and each one represented the same proportion of area to be planted, then this information was irrelevant. Selection based on the overall mean was all that was necessary to assure the largest overall gains (Dias et al., 2003). Therefore the genotypes IAN 6721, IAC 313 and IAC 308 could be considered superior in future breeding programs in order to incorporate stability for rubber yield. According to Singh and Gupta (1988), it was possible that stable genotypes carried genes for the stability, useful in breeding programs because it facilitated economic production.

Finally, the following major findings could be summarized from this study. (1) Genotypes IAC 302, IAC 307 and IAC 313 were widely adaptable for rubber yield, and were thus, recommended for commercial release in Votuporanga. (2) The significant genotype-year interaction and the change in ranks of genotypes across the years suggested a breeding strategy of specifically adapted genotypes in homogenously grouped environments. (3) Whenever new genotypes are proposed for commercial release, information on genotype-year interaction and stability clearly indicating their specific and or general adaptations needs to be available to the user. (4) The genotype IAC 40 was high yielding and vigorous, hence it could be considered as having suitable adaptation for favorable and unfavorable years.

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RESUMO

A interação genótipo x ano em culturas perenes como a seringueira Hevea brasiliensis (Willd. ex Adr. de Juss.) Muell.-Arg. representa o diferencial de resposta dos genótipos às mudanças climáticas anuais. O objetivo deste trabalho foi estudar a interação genótipo-ano e determinar genótipos temporalmente estáveis por meio da avaliação de seis anos de produção de borracha. Análises de estabilidade foram realizadas pelo método de Eberhart and Russell para produção de borracha na Estação Experimental de Votuporanga, região Noroeste do Estado de São Paulo, Brasil. Vinte e cinco genótipos foram analisados num delineamento de blocos ao acaso com três repetições. Os resultados mostraram que o melhor genótipo em um ano nem sempre foi o melhor em outro. Os componentes genéticos foram repartidos em lineares (genótipos dentro do ano) e não lineares (desvios agrupados). Quadrados médios significativos para os componentes lineares foram previsíveis indicando que o desempenho dos genótipos através dos anos em relação ao rendimento de borracha é passível de previsão. Entre os clones estudados o IAC 40 também foi considerado altamente produtivo e vigoroso, com adaptabilidade adequada.

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