



Genotype-environment interaction and phenotypic stability for girth growth and rubber yield of *Hevea* clones in São Paulo State, Brazil

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Abstract

The best-yielding, best vigour and most stable *Hevea* clones are identified by growing clones in different environments. However, research on the stability in *Hevea brasiliensis* (Willd. Adr. ex Juss.) Muell.-Arg. is scarce. The objectives of this work were to assess genotype-environment interaction and determine stable genotypes. Stability analysis were performed on results for girth growth and rubber yield of seven clones from five comparative trials conducted over 10 years (girth growth) and four years (rubber yield) in São Paulo State, Brazil. Stability was estimated using the Eberhart and Russell (1966) method. Year by location and location variability were the dominant sources of interactions. The stability analysis identified GT 1 and IAN 873 as the most stable clones for girth growth and rubber yield respectively since their regression coefficients were almost the unity ($\beta = 1$) and they had one of the lowest deviations from regressions (\overline{S}_{di}^2). Their coefficient of determination (R^2) was as high as 89.5% and 89.8% confirming their stability. In contrast, clones such as PB 235, PR 261, and RRIM 701 for girth growth and clones such as GT 1 for rubber yield with regression coefficients greater than one were regarded as sensitive to environment changes.

Key words: *Hevea brasiliensis*, rubber tree, coefficient of determination, stability parameters.

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Introduction

Genotype-environment interaction in crops such as the rubber tree [*Hevea brasiliensis* (Willd. Adr. ex Juss.) Muell.-Arg.] is the differential response of genotypes to changing environmental conditions. Such interactions complicate testing and selection in tree breeding programs and result in reduced overall genetic gain. The literature on genotype-environment interaction in *Hevea* is not extensive. Its effects have been recognized in Malaysia (Tan, 1995), Sri Lanka (Jayasekera, 1983; Jayasekera and Karunasekera, 1994 and Jayasekera *et al.*, 1977, 1984), Nigéria (Onokopise *et al.*, 1986) Indonésia (Daslin *et al.*, 1986), India (Menattoor *et al.*, 1991) and more recently in Brazil (Gonçalves *et al.*; 1990, 1991, 1998 abc, 1999 and, Costa *et al.*, 2000).

In *Hevea*, productivity performance is represented mainly by girth growth and yield stability. Breeders search for genotypes that show stability vigor and high yield over the years and locations. In general, a genotype is considered stable when its performance across environments does not deviate from the average performance of a group of standard genotypes.

Several methods have been proposed to analyze genotype x environment interactions and phenotypic stability (Lin *et al.*, 1986; Becker and Leon, 1988; Piepho, 1998; Truberg and Huhn, 2000). These methods can be divided into two major groups, univariate and multivariate stability statistics (Lin *et al.*, 1986). Joint regression is the most popular among the univariate methods because of its simplicity of calculation and application (Becker and Leon, 1988). Joint regression provides a conceptual model for genotypic stability (Becker and Leon, 1988; Romagosa and Fox, 1993). The regression of the yield or girth of an individual genotype on environment mean yields is determined. The

genotype x environment interaction from analysis of variance is portioned into heterogeneity of regression coefficients (β_i) and the sum of deviations ($\Sigma \bar{S}^2 d_i$) from regressions. Finlay and Wilkinson (1963) defined a genotype with regression coefficient equal to zero ($\beta_i = 0$) as stable, while Eberhart and Russell (1966) defined a genotype with $\beta_i = 1$ to be stable. Most biometricians consider $\bar{S}^2 d_i$ as stability parameter rather than β_i (Eberhart and Russell, 1966; Becker and Leon, 1988). According to the joint regression model, a stable genotype is one with a high mean yield, $\beta_i = 1$ and $\bar{S}^2 d_i = 0$ (Eberhart and Russell, 1966).

In Brazil, Kalil Filho (1982) was the first researcher to use Eberhart and Russell (1966) as a stability method. He studied temporal yielding, where the environments were represented by different months and years through an environmental index obtained by the mean performance of all the genotypes in each month and year. Later, Gonçalves *et al.* (1992, 1999), using the same stability method, examined the magnitude of the genotype x environment interaction on girth growth of *Hevea* at different ages, searching for clone stability and genetic gains.

The objectives of this study are (1) to examine the magnitude of genotype x environment interaction in ten-year girth growth and four-year rubber yield in *Hevea* clones, (2) to characterize these clones for stability.

Material and Methods

Plant material

One Brazilian, three Indonesian and three Malaysian genotypes (clones) were used in this study (Table 1). The Brazilian clone IAN 873 was developed by the breeding program of the former Instituto Agronômico do Norte, nowadays, Embrapa Western Amazon. The Indonesian clones comprised GT 1, PR 261 and PR 255 from Gondang Tapen and Proefstation voor Rubber stations. The Malaysia cultivars consisted of RRIM 701, RRIM 600 and PB 235 from Rubber Research Institute of Malaysian and Prang Besar private rubber plantation. Clones PB 235, GT 1, RRIM 600 and IAN 873 are recommended production clones, whereas RRIM 701, PR 255 and PR 261 are used as parents in local rubber breeding programs. The rubber clones were budded onto established rootstocks (Tjir 1 x Tjir 16) in the nursery. One-year-old rootstock seedlings raised in nurseries were used to budgraft clonal materials. Budgrafting was made on December, 1989. The successful budgrafts were uprooted and planted in polyethylene bags. After the first flush of leaves developed, the plants were established in the field.

Experimental locations

The clones were grown in a total of five contrasting test environments for ten years in the plateau region of São Paulo State (Table 2). These locations represent the most

Table 1 - Origin and individual genealogy of seven *Hevea* genotypes (clones) tested at different locations in São Paulo State, Brazil.

Clones ¹	Parentage	Origin
GT 1	Primary clone	Indonésia
PR 261	Tjir 1 x PR 107	Indonésia
PR 255	Tjir 1 x PR 107	Indonésia
IAN 873	PB 86 x FA 1717	Brazil
RRIM 701	44/553 x RRIM 501	Malaysia
PB 235	PB 5/51 x PB S 78	Malaysia
RRIM 600	Tjir 1 x PB 69	Malaysia

¹GT – Godang Tapen; RRIM – Rubber Research Institute of Malaysia; IAN – Instituto Agronômico do Norte; PR – Proefstation voor Rubber; PB – Prang Besar; Tjir – Tjirandji; FA – Ford Acre.

important continental climate non-traditional rubber production area. The experimental design at each test location was randomized complete blocks with three replications. Thirty plants were used in each plot in all locations except Matão. At Matão, a larger plot size of forty plants was used. Plots consisted of five rows of six or eight plants each. Trees were spaced 8.00 m between and 2.50 m within rows, resulting in approximately 500 trees ha.

Measurements

The girth growth and rubber yield of each tree were measured. In the first year diameter was recorded because plants were too small to measure the girth. Measurements were converted into girth assuming that plants were measured 0.50 m above ground level by a slide caliper. The other girth measurements were recorded 1.20 m from the highest point of the bud union.

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 latex production. Attempts were to record 10 girth measurements and four annual yields. The latex extracted from the panel followed a half-spiral four-daily tapping system (seven tapping/month) for 11 months/years. Yield was recorded on the days when normal tapping, which starts around 7:30 AM, was possible. After tapping, latex was allowed to collect in plastic cups provided for each recording tree. Once the latex was flow stopped, rubber was coagulated in the cup itself by adding 2% acetic acid solution and stirring it well. The coagulated rubber in each cup was then made into a “biscuit” which was dried ranged by wire tied in each tree for about 30 days, was weighed and the dry rubber content for each tree was recorded.

Statistical analysis

A combined three-factor analysis of variance was performed on data collected for all locations and years using the statistical model

Table 2 - Details of experimental locations and planting dates from five different test locations where seven *Hevea* clones were evaluated in São Paulo State, Brazil.

Remarks	Locations				
	Mococa	Ribeirão Preto	Votuporanga	Indiana	Matão
Year of planting	1989	1989	1989	1989	1989
Planting material	Budded stumps	Budded stumps	Budded stumps	Budded stumps	Budded Stumps
Spacing (m)	7.00 x 3.00	7.00 x 3.00	7.00 x 3.00	7.00 x 3.00	8.00 x 2.50
Planting density	5 rows x 6 trees 30 trees/plot	5 rows x 6 trees 30 trees/plot	5 rows x 8 trees 40 trees/plot	5 rows x 6 trees 30 trees/plot	5 rows x 8 trees 40 trees/plot
Design of experiment	RBD ¹	RBD	RBD	RBD	RBD
Numbers of replications	3	3	3	3	4
Total area (ha)	1.3	1.3	1.76	1.3	2.2
Elevation (m) (mean sea level)	665	467	450	621	551
Latitude (S)	21°28'	22°11'	20°25'	32°21'	21°25'
Longitude (EE)	47°01'	47°48'	49°50'	51°30'	48°25'
Temperature (annual mean)	24.5 °C	29.10 °C	22.3 °C	28.10 °C	21.0 °C
Annual rainfall (mm) (mean annual)	1500	1534	1480	1257	1430
Soil type ²	Eustrtox	Kandiudox	Paleudalf	Paleudalf	Paleudox
Terrain	Flat to undulating	Flat to undulating	Falt	Flat	Flat
Year of planting	1989	1989	1989	1989	1989

¹Randomized block design.

$$Y_{ijkl} = \mu + g_i + p_i + t_k + (gp)_{ij} + (gt)_{jk} + (tp)_{ik} + (gpt)_{ijk} + e_{ijkl}$$

where Y_{ijkl} is the i^{th} observation on the l^{th} clone in j^{th} location in the k^{th} year. The first four terms are the mean and the main effects of genotypes (clones), locations and years. The next three terms are the first order interactions, then the second order interaction and finally the micro-environmental deviation within locations and years. It is usually assumed that genotypes (clones) and locations are fixed effects and years random effects, so that the model is a mixed effects model.

All analysis were performed using the GENES computer program, Windows version, 2001 (Cruz, 2001). The method of Eberhart and Russell (1966) was used in this study to characterize genotypic stability. The following linear regression model was used:

$$Y_{ij} = \mu + \beta_i I_j + \delta_{ij} + \bar{\epsilon}_{ij}$$

where Y_{ij} is the mean of the clone i^{th} at the location j ; μ is the general mean of clone i ; β_i is the regression coefficient of the i^{th} clone at the location index which measures the response of this clone to varying location; I_j is the environmental index which is defined as the mean deviation of all clones at a given location from the overall mean; δ_{ij} is the deviation from regression of the i^{th} clone in the j^{th} location; $\bar{\epsilon}_{ij}$ is the mean of experimental error.

Two stability parameters were calculated: (a) the regression coefficient, which is the regression of the performance of each clone under different locations on the

environment, means over all the genotypes. This is estimated according to Sing and Chaudhary (1979) as follows:

$$\beta_i = \frac{\sum_j Y_{ij} I_j}{\sum_j I_j^2}$$

where $\sum_j Y_{ij} I_j$ is the sum of products and $\sum_j I_j^2$ is the sum of squares

(b) Mean square deviations (\bar{S}_d^2) from linear regression,

$$= \frac{\sum_j \delta_{ij}^2}{(s-2)} - \frac{\bar{S}_d^2}{r}$$

where $\sum_j \delta_{ij}^2 = \left[\sum_j Y_{ij}^2 - \frac{Y_i^2}{t} \right] - \frac{\left(\sum_j Y_{ij} I_j \right)^2}{\sum_j I_j^2}$ and $S_e^2 =$ the

estimate of pooled error.

The contribution of locations in generating interactions was studied using a method developed by Burdon (1977). It considers growth in different locations as separate traits and calculates the genetic correlations between pairs of locations.

Results and Discussion

The results of the combined analysis of girth growth and rubber yield across locations and years are given in Ta-

ble 3. Effects of years, locations, genotypes and their interactions except for genotype year (G x Y) for rubber yield ($p < 0.05$) were highly significant ($p < 0.01$). Out of the total variance, a larger portion of variation was caused by the locations effect and their interaction with years (S x Y), followed by the years effect. The former accounted for about 17.0% while 14.5% could be attributed to the latter. The highly significant differences for S and Y ($p < 0.01$) indicated the fluctuations of genotypes in their responses to the different environments.

There was also tremendous change in girth growth ranks of the genotypes across locations (Table 4). Similarly, a recent study (Gonçalves *et al.*, 1999) undertaken on the girth growth stability at six years also revealed very high fluctuations in the growing environments of São Paulo State. This shows the difficulties encountered by breeders in selecting new genotypes (clones) for release; these difficulties arise mainly from the masking effects of variable environments. Pham and Kang (1988) indicated that genotype x environment interactions minimize the usefulness of genotypes by confounding their yield performance. Thus, it is important to study in depth the vigor levels, adaptation patterns and stability of genotypes in multi-location trials. Becker and Leon (1988) also indicated that assessment of stability across many locations and years could increase both repeatability and heritability of important traits.

The genotype means for girth growth and rubber yield, their ranks and differences among genotypes means for the seven genotypes tested across five (girth growth) and four (rubber yield) locations over the tenth and fourth years, respectively, are presented in Table 4. The highest girth growth of 62.67 cm was obtained from PB 235 clone at Votuporanga, while the lowest was from RRIM 701 at Mococa. Across the locations, however, PB 235 surpassed

Table 3 - Analysis of variance of girth growth (cm) and rubber yield for seven *Hevea* genotypes (clones) tested at five locations in São Paulo State, Brazil.

Sources of variation	Girth growth		Rubber yield	
	D.F.	M.S.	D.F.	M.S.
Replications (P/Y)	100	5.0317	32	19.7662
Genotypes (G)	6	219.3001**	6	2,408.9117**
Years (Y)	9	33,506.2674**	3	916.8731**
Locations (L)	4	4,220.9545**	3	14,991.8680**
G x Y	54	8.1058**	18	26.6794*
G x L	24	67.2242**	18	298.1749**
Y x L	36	111.7173**	9	1,089.6314**
G x Y x L	216	4.2320**	54	64.5435*
Residual (mean)	600	3.7666	192	18.7875
Média		32.1829		36.2198
CV(%)		6.0305		11.9671

* and ** significant for $p < 0.05$ and $p < 0.01$, respectively.

Table 4 - Mean of ten-years girth growth (cm) and four-years rubber yield (g/tt)¹ of seven *Hevea* clones grown at five locations São Paulo State, Brazil.

Genotypes	Indiana		Votuporanga		Matão		Ribeirão Preto		Mococa		Overall mean	
	Girth growth	Rubber yield	Girth growth	Rubber yield	Girth growth	Rubber yield	Girth growth	Rubber yield	Girth growth	Rubber yield	Girth growth	Rubber yield
GT 1	56.35a(1)	62.00a(1)	58.74a(5)	36.46b(6)	62.19a(2)	-	56.30a(2)	30.65c(7)	52.21ab(3)	29.19bc(5)	57.16	39.58
IAN 873	54.80a(3)	49.37bc(5)	51.09b(7)	37.33b(5)	59.26a(3)	-	54.70a(4)	31.45c(6)	50.56ab(5)	32.43abc(3)	54.08	37.65
PB 235	55.81a(2)	55.80ab(1)	62.67a(1)	67.97a(1)	62.41a(1)	-	55.66a(3)	40.10a(3)	52.55ab(2)	43.19a(1)	57.82	51.76
PR 255	52.17a(7)	56.33ab(2)	55.94ab(6)	43.26b(2)	58.09ab(4)	-	51.28a(7)	42.53a(1)	50.71ab(4)	39.78ab(2)	53.64	45.47
PR 261	54.11a(5)	39.97c(7)	58.96a(4)	31.90b(7)	57.51ab(6)	-	53.22a(5)	32.73bc(5)	49.69bc(6)	24.80c(6)	54.70	32.35
RRIM 600	54.53a(4)	55.50ab(4)	60.82a(2)	40.00b(4)	52.50b(7)	-	56.39a(1)	42.10a(2)	56.36bc(1)	32.37abc(4)	56.12	42.49
RRIM 701	53.36a(6)	43.10c(6)	60.53a(3)	42.70b(3)	57.56a(5)	-	53.06a(6)	38.47b(4)	44.28b(7)	24.54c(7)	53.76	37.20
Mean	54.45	51.72	58.39	42.80	58.50	-	54.37	36.86	50.91	32.33	55.33	40.93
CV (%)	2.41	14.16	14.16	25.45	5.31	-	3.27	12.87	6.63	20.27	3.08	15.47

Values in girth growth and rubber yield columns followed by the same letters are not significantly different at $p = 0.05$ according Tukey's test. Ranks are given in the parentheses
¹g/tt = grams per tree per tapping.

all other genotypes with a mean girth growth of 57.16 cm. Its mean girth growth exceeded that of PB 255 by about 5% indicating its high vigour and good adaptability to the *Hevea*-growing environment of São Paulo.

The partitioning of variance components revealed that environment factors both predictable (locations) and unpredictable (year) were important sources of variation (Table 3). When genotype x environment is due to variation in predictable environment factors, *Hevea* breeders have the alternatives of either developing specific varieties for different environments (location, soil types, management systems, etc.) or broadly adapted varieties that can perform well under variable conditions. However, when genotype x environment interaction results from variation in unpredictable environmental factors, such as year to year variation in rainfall distribution, as in the case in this study, the breeder needs to develop stable genotypes that can perform reasonably well under a range of conditions. Such breeding strategies can assist the rubber producer in risk avoidance. Ceccarelli (1994) and Piepho (1998) indicated that producers perceive yield stability as the most important socio economic aim to minimize crop failure, especially in marginal environments.

The stability analysis conducted for ten-year girth growth and four-year rubber yield at five locations of the study is presented in Table 5, and revealed that the clones differ significantly for both characters. The clone x location interaction component was further partitioned into linear (locations and clones x locations) and non-linear (pooled deviations) components. Mean squares for both of these components were tested against pooled error mean square. The linear component was highly significant, indicating that the predictable-components shared with clone x location interactions. Preponderance of linear clone x location interaction is of great practical importance, implying that there are differences among linear regression coefficients for each clone.

Table 5 - Analysis of variance for stability parameter for seven *Hevea* genotypes (clones) of ten years girth growth and rubber yield tested at five and four different locations, respectively São Paulo State, Brazil.

Source of variation	Girth growth		Rubber yield	
	d.f.	M.S.	d.f.	M.S.
Clones	6	43.6412*	6	480.8276**
Locations (clone x location)	28	1,402.8983**	21	310.8017**
Locations (linear)	1	354.7185**	1	4416.9052**
Clones x locations (linear)	6	40.1737**	6	72.0071**
Pooled deviation	21	14.4541**	14	119.8492**
GT 1	3	5.6365	2	117.6850**
IAN 873	3	37.6206**	2	31.1334
PB 235	3	2.4694	2	462.0978**
PR 255	3	6.5224	2	30.6365
PR 261	3	1.5828	2	23.6860
RRIM 600	3	37.4819**	2	58.1025*
RRIM 701	3	9.8653	2	115.6029**
Pooled error	60	5.4635	48	14.2419

* and ** significant for $p < 0.05$ and $p < 0.01$, respectively.

The stability parameters for all clones are given in Table 6. Eberhart and Russell (1966) emphasized the need of considering both linear (β_i) and non-linear ($\bar{S}^2 d_i$) components of genotype environment interactions in judging the stability of a genotype. A wide adaptability genotype was defined as one with $\beta_i = 1.0$ and high stability as one with $\bar{S}^2 d_i = 0$. In this study values for the regression coefficient (β_i) ranged from 0.0723 (RRIM 600) to 1.8608 (RRIM 701) for girth growth. For rubber yield the range for β_i was 0.6875 (PR 261) to 1.7196 (GT 1). Plotting all clonal means x location means with the corresponding regression lines illustrates the differential reactions of clones to changing environments (Figure 1).

Table 6 - Estimates of stability and adaptability parameters of the traits girth growth (cm) and rubber yield (g/t)¹ for seven *Hevea* clones grown at five locations São Paulo State, Brazil.

Genotypes	² \bar{x}_i		³ $\hat{\beta}_i$		⁴ $\bar{S}^2 d_i$		⁵ R_i^2 (%)	
	Girth growth	Rubber yield	Girth growth	Rubber yield	Girth growth	Rubber yield	Girth growth	Rubber yield
GT 1	57.1573	39.5608	1.0874	1.7196**	0.0577	34.4810**	89.5169	88.7984
IAN 873	54.0813	37.6558	0.5314*	0.9299	10.7190**	5.6305	23.4013	89.7563
PB 235	57.8187	51.7633	1.3899	0.9260	-0.9980	149.2853**	96.9549	36.9281
PR 255	53.6367	45.4625	0.9262	0.8266	0.3530	5.4649	84.2595	87.5558
PR 261	54.6973	32.3525	1.1319	0.6875*	-1.2936	3.1480	97.0541	86.2946
RRIM 600	56.1661	42.4942	0.0723**	1.0665	10.6728	14.6202*	0.5651	86.0644
RRIM 701	53.7553	37.2025	1.8608**	0.8439	1.4673	33.7869**	93.4576	66.0275

* and ** significant for $p < 0.05$ and $p < 0.01$, respectively; ¹g/t/t = grams per tree per tapping. ² \bar{x}_i = mean; ³ $\hat{\beta}_i$ = mean regression coefficient; ⁴ $\bar{S}^2 d_i$ = deviation from regression; ⁵ R_i^2 % = coefficient of determination.

The regression coefficient of clones GT 1, PB 235, PR 261 and PR 255 for girth growth was non-significantly different from the unity ($\beta_i = 1$) and had a small deviation from regression ($\bar{S}^2 d_i$) and thus possessed fair stability. Finlay and Wilkinson (1963) and Eberhart and Russell (1966) stated that genotypes with high mean yield, regression coefficient equal to the unity ($\beta = 1$) and deviation from regression as small as possible ($\bar{S}^2 d_i = 0$) are considered stable. Accordingly, GT 1 and IAN 873 were the most stable clones for girth growth and rubber yield, respectively, since their regression coefficients were almost equal to the unity and they had the lowest deviations from regression. Their coefficients of determination, R^2 (Pinthus, 1973), were as high as 89.5% and 89.8% confirming their stability. In contrast, the clones PB 235, PR 261 and RRIM 701 for girth growth and the clones GT 1 for rubber yield, with regression coefficients greater than one, were regarded as sensitive to environmental changes.

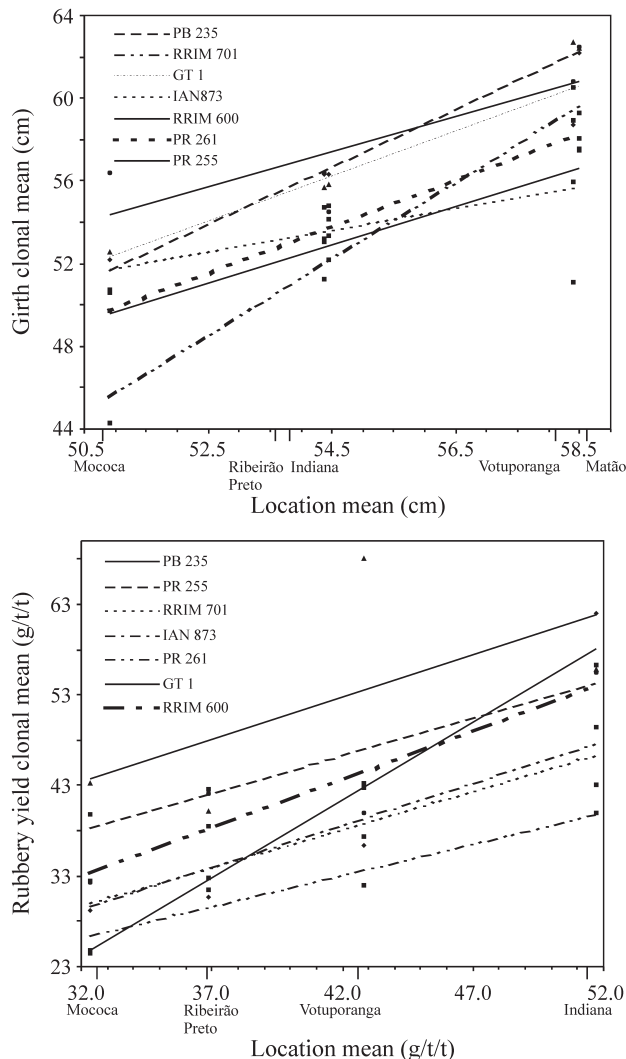


Figure 1 - Plot of girth growth and rubber yield clonal means for seven clones against site means with corresponding regression lines.

Figure 2 shows a representation of the genotype regression coefficients plotted against the means. The genotype PB 235 ($\beta_i = 1.39$) had a regression coefficient for girth growth greater than the unity and was highly vigorous and high yielding, so it may be characterized as suitable for specific adaptation in favorable environments. The locally adapted cultivars had regression coefficients close to the unity, had above average yields and may, therefore, be characterized as well adapted to all environments. These genotypes also had the smallest deviations from regression and, hence, may be regarded as stable genotypes. The genotypes PR 261 and RRIM 701 that had regression coefficients of less than unity and below average rubber yields, indicating that they offer a greater resistance to environmental change and are specially adapted to poor environments.

The regression coefficient measures the “relative” performance of the clone. Clair and Kleinschmit, (1986) emphasizes that in forest tree breeding, this information is useful to distinguish clones for specific environments, but if all environments tested are in one planting zone, and each represents the same proportion of area to be planted, then this information is irrelevant. Selection on the overall mean is all that is necessary to assure the largest overall gains. Therefore clones GT 1 and IAN 873 could be considered superior in future breeding programs in order to incorporate stability for vigour and rubber yield. According to Singh and Gupta (1988) it is possible that stable genotypes carry genes for stability, useful in breeding programs because it facilitates economic production.

Table 7 shows the correlation matrix among locations. Girth growth of the trees at Mococa was significantly

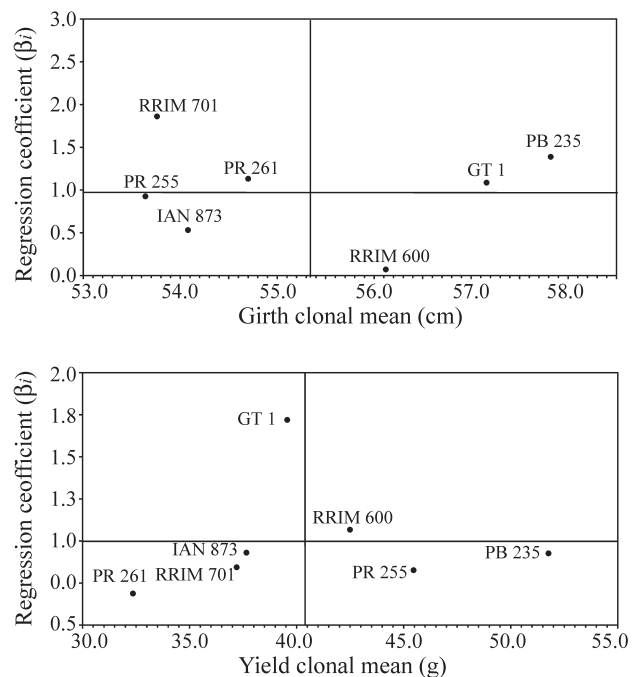


Figure 2 - Plot of deviation from regression coefficient against girth and yield clonal means in a stability study of seven *Hevea* clones.

Table 7 - Genetic correlation coefficients between locations for girth growth and rubber yield (between parenthesis) in São Paulo State, Brazil.

Locations	Ribeirão Preto	Indiana	Votuporanga	Matão
Mococa	0.623981n.s. (0.528938n.s.)	0.422400n.s. (0.600499n.s.)	0.104463n.s. (0.758129*)	0.783443* -
Ribeirão Preto		0.889579** (0.175431n.s.)	0.288359n.s. (0.506696n.s.)	0.888974** -
Indiana			0.219663n.s. (0.305830n.s.)	0.760962* -
Votuporanga				0.379341n.s. -

n.s.: no significant; * and **: significant for $p < 0.05$ and $p < 0.01$, respectively.

correlated with Matão. Therefore the girth growth of Mococa was not necessarily a good indication of vigour. Only at Ribeirão Preto did tree girth growth show very high and stable statistically significant correlations with the girth growth of Indiana and Matão. There was also correlation among tree girth growth between Indiana and Matão. On the other hand, rubber yield were not necessarily a good indicator of correlation among locations. Only Mococa showed statistically significant correlations with Votuporanga.

Finally the following major findings can be summarized from this study: (1) GT 1 and IAN 873 were the most stable clones for girth growth and rubber yield, respectively, and are thus recommended for commercial release in São Paulo. (2) The significant genotype-environment interactions and the change in rank of genotypes (clones) across environments suggest a breeding strategy of specifically adopted genotypes in homogeneously grouped environment. (3) Whenever new varieties are proposed for commercial release, information on genotype-environment interactions and stability, clearly indicating their specific and/or general adaptations, needs to be available to the user.

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