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Research Article

Age-age correlation for early selection of rubber tree genotypes in São Paulo State, Brazil

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Abstract

In a field trial involving 68 rubber tree (*Hevea* spp.) clones calculation of genotypic correlation coefficients revealed significant age-age correlation from age 1 to 6 (immature period) for girth A and for age 7 to 12 (mature period) for girth B and for age 7 to 12 (production of latex) for yield. Rank correlation coefficients between all immature ages of girth (girth A), all ages of mature girth (girth B) and all annual rubber production (yield) were significant for the three traits, with the coefficients decreasing with increasing age. Selection of the sets of best 30, 15, 10 and 5 clones from the available 68 clones at a given age was generally accompanied by a descending order of percentage success. It was suggested: (a) to have the best 30 clones of age 6, select the set of best 36 clones at age 2, (b) to have the best 15 clones of age 6, select the set of best 3 clones of age 6, select the set of best 3 clones at age 5. More than 80% of the targeted clones on girth A or girth B basis and more than 76.7% clones on yield basis were found to get selected at steps (a) through (d). For achieving early multiplication of the most productive clone for deployment, multiplication should be started with the best 36 (*i.e.* 60%) clones selected at age 2.

Key words: Hevea brasiliensis, early clonal multiplication, genetic correlation, rank correlation, evaluation of clones. Received: June 15, 2004; Accepted: March 23, 2005.

Introduction

The major source of natural rubber is the rubber tree, *Hevea brasiliensis* (Willd. Adr. ex Juss.) Muell.-Arg. (family Euphorbiaceae), which is perennial and has na economically useful lifespan of '30-35 years. Adult trees shed their leaves almost completely for 3-4 weeks each year in a process known as wintering, subsequent to which they undergo "flushing" and produce new flowers, leaves and shoots. Flowering in *Hevea* normally occurs once a year after leaf fall and is directly affected by climatic factors, *Hevea* species having traditionally been cultivated in humid equatorial regions subject to monsoons (Rao and Vijayakumar, 1992) and characterized by adequate rainfall distribution and small yearly fluctulations in temperature and humidity (Rao *et al.*, 1993). In some countries to achieve self-

Send correpondence to Paulo de Souza Gonçalves. Instituto Agronômico de Campinas, Programa Seringueira, Caixa Postal 28, 13001-970 Campinas, SP, Brazil. E-mail: paulog@iac.sp.gov.br. sufficiency, attempts have been made to extend *Hevea* cultivation to marginal areas with varied climatic constraints, some areas being as far north as 29° in India and China and south to 23° in the Brazilian state of São Paulo (Pushparajah, 1983; Ortolani *et al.*, 1998).

Rubber trees are grown for their latex from which is collected by tapping, coagulated and processed to form rubber, hence it is the dry rubber yield that determines the level of output. Girth is considered to be a measure of vigor and is also an economically important character because it determines the age at which a *Hevea* clone can be commercially exploited by tapping and is, therefore, important in shortening the uneconomic immature period of the clone. However, there are several high yielding clones whose girthing is depressed by tapping (Abraham, 1992) because this procedure is controlled wounding and as such stresses the tree and retards growth, especially if carried out when the trees are budding.

In tree breeding, early selection has been shown to have three main advantages: increased overall selection intensity or a reduce in the amount of field-testing a shortened generation interval; and the fact that genetic information from early testing can be used to enhance selection efficiency maturity (Wu, 1998). Wu (1998) has developed a theoretical basis to study the effects of increased overall selection intesity and reduced field-testing but the most frequently cited advantage of early selection in tree breeding is shortened generation interval because this causes a reduction in the length of the breeding cycle (Nanson, 1970; Lambeth, 1980; McKeand, 1988; Adams et al., 1989; Adams and Atiken, 1992; Matheson et al., 1994). Shortening the breeding cycle of a tree through early selection can produce more genetic gains per unit time (year) if there is strong genetic correlation between early and mature traits.

Rubber is a forest tree crop and takes more than 25 years to complete a breeding cycle because of which genetic improvement of *Hevea* is normally a slow process, even for a single selection and breeding cycle. In *Hevea* breding there is usually a lapse of 3 years from pollination to field planting and since rubber trees generally require several years (often 6 or more) to reach tapping girth there is a wait of 7-9 years until tapping is started and then a 10-15 years period of tapping and clone evaluation.

Determining trends in age-age correlation is relatively simple for growth and yield traits, requiring only patience and repeated assessment. While there have been many reported age-age correlations for growth in conifers (*e.g.* Lambeth, 1980; McKeand, 1988; Riemenschneider, 1988; King and Burdon, 1991; Matheson *et al.*, 1994) there are few for *Hevea* species (Gonçalves *et al*, 1988.)

The objective of the research described in the present paper was to use age-age correlations and selection efficiency estimates to study the potential of early selection of *Hevea* for forward breeding using data from a series of assessments of a *Hevea* spp. clone trial carried out in the northwest of the Brazilian state of São Paulo.

Material and Methods

In this study we used 55 Brazilian, 10 Malaysian and three Indonesia genotypes (clones) from a small-scale clone trial belonging to the Campinas Agronomic Institute (Instituto Agronômico de Campinas - IAC), Campinas, São Paulo state, Brazil) breeding program. The Brazilian clones were developed by the IAC breeding program while the Malaysian genotypes consisted of seven Rubber Research Institute of Malaysia (RRIM) clones and three clones belonging to the Prang Besar (PB) private rubber plantation. The Indonesian genotypes (clones) came from Proefstation voor Rubber (PR) in Sumatra. All the clones were budded onto established rootstock (Tjir 1 x Tjir 16) at the nursery. One-year-old rootstock seedlings raised in nurseries were used to budgraft clonal materials in December 1990, successful budgrafts being transplanted into polyethylene bags and the plantlets established in the field after the first flush of leaves developed.

The clones were grown at the Votuporanga Experimental Station (20°25' S, 49°59' W, elevation 450 m) in the nortwest of the Brazilian state of São Paulo. Mean monthly temperatures varied from 20 to 25°C and the average annual rainfall ranged from 1.087 mm to 1.537 mm and the winter drought varied from four to six dry months with an average water deficiency of 180 mm. The trial was laid out in a randomized complete block design with three replicates using six trees replications per plot and 7 x 3 m spacing. Missing plants were replaced with spares during the first two years after planting to maintain plantation density but were not scored. One row of the commercial clone RRIM 600, acquired from a commercial nursery, was planted around the plot. Annual fertilization consisted of 400 g of 10-10-10 NPK formula per plant (Bataglia and Gonçalves, 2002).

Twelve months after planting the diameter of the stems of the plantlets were measured 0.5 cm above ground level using a slide caliper and this measurement converted to girth assuming that the stem was cylindrical. Immature girth (girth A) was measured at 2, 3, 4, 5 and 6 years and mature grith (girth B) at 8, 9, 10, 11 and 12 years. At the end of the sixth year all the trees with a girth of 45 cm or more were tapped for annual latex production 1.2 m above the highest point of the bud union using a half-spiral four-day tapping system (seven tappings per month) for 11 months per year. Yield was recorded on the days when normal tapping, which starts around 7:30 AM, was possible, the latex being collect in individual plastic cups for each tree and coagulated by adding 2% (v/v) acetic acid solution, with stirring, to the cup once the latex flow had stopped. The coagulated rubber in each cup was made into a "biscuit" which was dried for about 30 days by hanging it on a wire tied to the tree from which it had been extracted. After dry, each rubber biscuit was weighed and the dry rubber content for each tree recorded. Attempts were made to record 12 girths measurements and six annual yields.

A combined three-factor cross classification analysis of variance (ANOVAR) was performed on the data using the statistical model:

$$Y_{ijk} = \mu + g_i + a_j + b_k + (ga)_{ij} + (gb)_{ik} + (ab)_{jk} + (gab)_{ijk},$$

where, $Y_{ijk} =$ observed value of the *i*th genotype in the year in the *k*th replication; $\mu =$ general mean; $g_i =$ fixed effect of the *i*th genotype (*i* = 1, 2,g, g = 68); $a_j =$ effects of the *j*th year (*j* = 1, 2,s, s = 6); $b_k =$ effect of the *k*th randomized block (k = 1, 2, b, b = 3); (ga)_{*ij*} = interaction between *i*th genotype with the *j*th year; (gb)_{*ik*} = interaction between *i*th genotype with the *k*th replication; (gab)_{*ijk*} = interaction between genotype, year and replication. It the assumed that the interactions (ab)_{*ij*} and (gb)_{*ik*} were not significant, these being included in the residual together with (gab)*ijk*. Data from more than three traits were used for computation of genotypic age-age correlation using the Selegen software (Resende and Oliveira, 1997). Values of genotypic correlation were computed among all traits and age combinations. Genotypic correlations were worked out according to Kempthorne (1966) as follows:

$$r_{g(xy)} = \frac{COVg_{(xy)}}{\sqrt{\sigma_{gx}^2 \cdot \sigma_{gy}^2}}$$

where $COVg_{(xy)}$ = the genotypic covariance for traits x and y; σ_{gx}^2 = genotypic variance for trait x; σ_{gy}^2 = genotypic variance for trait y.

For each trait, Spearman's rank correlation coefficient (r_s) was computed between age 6 and younger ages. Rank 1 was assigned to the lowest value and rank 68 to the highest value within each age for a given trait. The significance of these correlations was tested using the Table XXV of Fisher (1941), for genotypic correlations for 5 and 1% probability. The genotypic correlations were tested at the degrees of freedom of the error. Percent success in correctly selecting the best 5, 10, 15 or 30 clones at age 2 and above was calculated using the ranks of clones method. Scatter plots between ranks at age 6 and at younger ages were plotted for each trait. Scatter plots of ranks for girth between age 2 and 3, age 3 and 4, and age 4 and 5 were also produced, showing the identification numbers of respective clones against the plotted points.

Results

All effects including the genotype versus years interaction were highly significant (p < 0.01) statistically in the overall analysis of variance for girth A, girth B and yield (Table 1).

Significant positive genotypic correlation coefficients (r_g) were observed for all the combinations of traits and ages (Table 2). The r_g values were higher for traits of the same age group and as the difference increased r_g usually decreased. The r_g values within a trait varied from 0.3916 to 0.9916 for girth A, 0.5116 to 0.9517 for girth B

and 0.3916 to 0.9612 for yield. The r_g values between girth A and girth B were high, although both girth values showed low r_g values in relation to yield.

Similarly, significant of rank correlation coefficient (r_s) values were observed between age 6 and younger ages for girth A, girth B and yield (Table 3). The values of r_s with reference to age 6 ranged from 0.5498 (for age 1) to 0.9420 (for age 5) for girth A, from 0.5114 (for age 1) to 0.9471 (for age 5) for girth B, and from 0.3451 (for age 1) to 0.8311 (for age 5) for yield. Since r_s was significant for all age combinations, it shows that there were no drastic changes in the ranks of the different clones between age 1 and 6.

Table 4 shows that selecting the best 30 clones at age 2 would lead to the best 69.93% of clones on a girth A basis (or 76.59% on a girth B or yield basis) being correctly selected at age 6 and the proportion increased to between 83.25% and 89.91% if selection was made at age 3 and 89.91% to 96.57% at age 4. The success in correctly selecting the set of the best 15 clones at age 6 by selection at age 3 was 66.70% on a girth A basis and 80.04% on a girth B or yield basis, the success being much lower when the trees were selected at less than 3 years of age. The success in correct selection of the set of the best 5 clones at age 4 was 80% on a girth A or girth B basis and only 60% on a yield basis.

Figure 1 shows the scatter plot of ranks of clone means for girth A by plotting the ranks at age 6 along X-axis and ranks at younger ages along the Y-axis. With an increase in age along the Y-axis, the plotted points tend to converge along the imaginary diagonal drawn from the origin. If the ranks on the X and Y-axes belong to the same age, the plotted points would be axiomatically be on the diagonal. As the difference in age in the two axes increased, the values of X and Y fall more and more out of step, resulting in increased scattering of the points and lower r_s values.

Of the best 34 girth A clones (ranking from 35 to 68) at age 6, 24 were also among the best 34 clones at age 2 while 27 clones were among the best 36 clones at this age. By selecting the best 34 girth A clones at age two we could select 76.7% of the best 34 clones at age 6 and this success rate could be increased to 90% if 38 age 2 clones were used. Similarly, by selecting the best 15 girth A clones at age 3 we

Table 1 - Analysis of variance for immature girth ages (girth A), girth mature ages (girth B) and rubber production (yield) for 68 *Hevea* genotypes (clones) grown in Votuporanga, São Paulo State, Brazil.

Sources of variation	d.f.	Girth A (cm)	Girth B (cm)	Yield (g)
Replications	2	6.1755	27.3758	164.6510
Genotypes	67	138.7458**	206.8648**	2002.3679**
Years	5	12642.7641**	4747.0498**	12670.7851**
Genotypes x years	335	12.0233**	8.6361ns	29.9479**
Residual	815	3.8935	8.8062	65.70636
Mean		21.82	56.98	40.71
Coef. of variation (%)		9.02	5.20	19.91

**significant for p < 0.01.

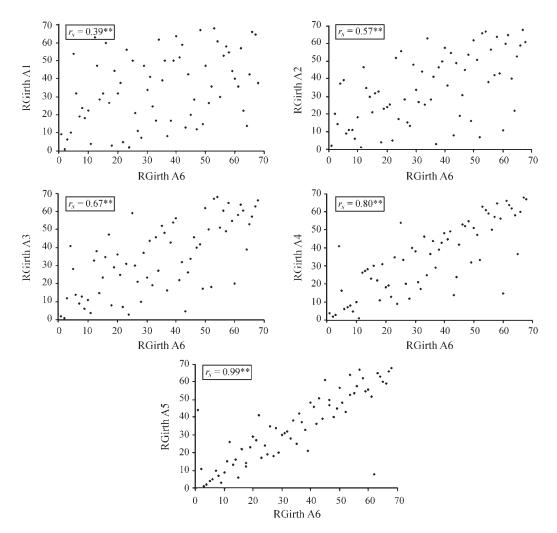


Figure 1 - Scatter plot ranks recorded at age six ranks at age 1 (i), 2 (ii), 3 (iii), 4 (iviv) and 5 (v) for mean girth A values of 68 clones. The rank of the girth A values at 1 year, 2 years, etc are indicated as Rgirth A1, Rgirth A2.

could select 80% of the set of the best 15 clones at age 6 and this success rate could be increased to 93.3% if 20 age 2 cloens were used. Selecting the best 5 girth A clones at age 4 meant that 60% of the set of the best 5 clones could be selected at age 6 and this could be increased to 100% if 8 clones were selected at age 4 (Figure 1). The scatter plots of girth B and yield (not shown) also showed a similar trend. Selection of the set of the 3 most productive clones at age 6 could be made with accuracy at age 5, although the ranks of the three clones changed among themselves during the sixth year.

Of the best 20 clones at age 3, the number of clones that simultaneously existed in the set of the best 36 clones at age 2 was 20 on a girth A basis [Figure 2(i)] and 19 each on a girth B or yield basis. The best 8 clones at age 4 invariably appeared in the set of the best 20 clones at age 3 for girth A (Figure 2 (*ii*)), girth B or yield. Each of the best 5 clones at ages 5 and 6 were among the best 8 clones at age 4 for girth A (Figure 2 (*ii*)), or girth B, while only 4 clones out of 5 top-ranking clones at age 6 could do so in respect of yield.

Discussion

The presence of significant clone versus site interaction indicates that genotypes perform differently between years for girth A, girth B and yield.

Existence of a strong, positive genotypic correlation (Table 2) as well as rank correlation (Table 3) between girth A, girth B and yield suggests that selection of clones for superiority in respect of one of these traits would inevitably lead to selection of superior clones in respect to the other two traits as well. However, any decision on the selection of clones should preferably be taken on the basis of the girth A trait because vigor, represented by girth volume, is more closely related with girth A than with girth B or yield only. We also suggest that for early evaluations girth A should be used in preference to secondary characters such as height, number of laticiferous vessel rings, etc.

It was not possible to predict with any fair degree of success as to which clone would occupy a particular rank at

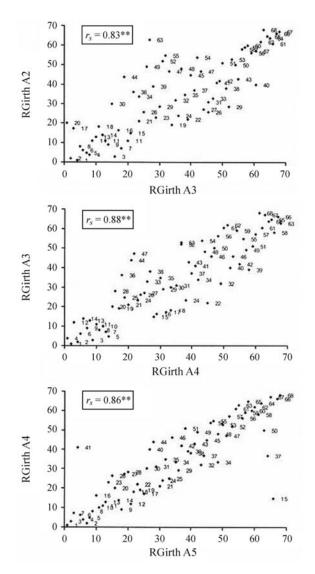


Figure 2 - Scatter plot of ranks for Girth A recorded between (i) age 2 and 3, (ii) age 3, (ii) age 3 and 4 (*iii*) age 4 and 5. The clone identification numbers are printed above the corresponding points in the plot.

panel opening age (Figure 1). At younger ages, a realistic attempt can be made to determine only the set of the best n clones (n > 1) at panel opening age, without allocating the n ranks of panel opening age to individual clones. Therefore, exchange of ranks between clones within the selected set of clones was not viewed as a waste of selection effort during this study.

Gottardi *et al.*, (1995) reported that the ranking in girth A among 11 *Hevea* clones did not vary much after 3 years up to 7 years of age. The mean data for each clone showed good genotypic correlation for girth between age 7 and young ages. The coefficient increased with increasing age suggesting that early selection was possible. Gonçalves *et al.*, (1998) suggested that selection of fastest growing

	Girth A2	Girth A3	Girth A4	Girth A5	Girth A6	Girth B1	Girth B2	Girth B3	Girth B4	Girth B5	Girth B6	Yield 1	Yield 2	Yield 3	Yield 4	Yield 5	Yield 6
Girth A1	0.7516**	0.6320^{**}	0.5514^{**}	0.4912^{**}	0.3916^{**}	0.3817^{**}	0.3325**	0.3134^{**}	0.3136^{**}	0.3142**	0.3233**	-0.0848ns	-0.846ns	-0.0242ns	$-0.0931 \mathrm{ns}$	-0.0529ns	0.0122ns
Girth A2	0.79**	0.8312^{**}	0.7715^{**}	0.6412^{**}	0.5734**	0.5236^{**}	0.4532**	0.4741^{**}	0.4255**	0.4151**	0.3353**	-0.0842ns	-0.0643ns	-0.1215ns	-0.0917ns	0.0116ns	0.1722ns
Girth A3			0.8812**	0.7633**	0.6736**	0.5638**	0.4842^{**}	0.4846^{**}	0.4141^{**}	0.3156^{**}	0.3632**	-0.1731ns	-0.1722ns	-0.1827ns	-0.0423ns	-0.0420ns	0.0212ns
Girth A4				0.8626^{**}	0.8018^{**}	0.7116^{**}	0.6712^{**}	0.6514^{**}	0.5722**	0.4416^{**}	0.4817**	-0.2032ns	-0.1941ns	-0.2322*	-0.0912ns	0.0313ns	0.0414ns
Girth A5					0.9916^{**}	0.8533**	0.8542**	0.8145**	0.7646^{**}	0.6431^{**}	0.6521^{**}	0.2512*	-0.2319ns	-0.3026*	0.3038*	0.1617ns	0.0416n.s
Girth A6						0.8442**	0.8633^{**}	0.8138^{**}	0.8036^{**}	0.6117^{**}	0.6122^{**}	0.2632**	-0.2735**	-0.3527**	0.3429**	0.2244ns	0.0717 ns
Girth B1							0.8202^{**}	0.7855**	0.7636^{**}	0.5327**	0.5116^{**}	0.3417**	-0.3316^{*}	-0.3714**	0.3713*	0.1116ns	-0.0315ns
Girth B2								0.9255**	0.9016^{**}	0.7127^{**}	0.6918^{**}	-0.0532ns	-0.0622ns	-0.1016ns	0.3217ns	0.2514ns	0.0516ns
Girth B3									0.9012^{**}	0.6933**	0.6932**	-0.0342ns	-0.0449ns	-0.0646ns	0.2932ns	$0.2941 \mathrm{ns}$	0.1953ns
Girth B4										0.8332**	0.7741^{**}	-0.0232ns	-0.0433ns	-0.0712ns	-0.2718ns	0.2355ns	0.1114ns
Girth B5											0.9517^{**}	-0.1715ns	-0.1312ns	-0.0316ns	-0.1732ns	0.1642ns	0.0916ns
Girth B6												-0.2712ns	-0.1217ns	-0.1333ns	-0.0723ns	0.1222ns	0.0916ns
Yield 1													0.9612^{**}	0.8717^{**}	-0.7222**	0.6125**	0.3916^{*}
Yield 2														0.8914^{**}	0.7816^{**}	0.7132**	0.4412**
Yield 3															0.8817 * *	0.7016^{**}	0.5022**
Yield 4																0.8413^{**}	0.4821**
Yield 5																	0.6914^{**}

Table 2 - Genotypic (rg) correlation coefficients for/among all immature girth ages of (girth A), mature girth ages (girth B) and yield recorded at age 1, 2, 3, 4, 5, and 6 for 68 Hevea genotypes (clones) grown in

Table 3 - Values of Spearman's rank correlation coefficient (r_s) between age 6 and younger ages for immature girth age (girth (A), mature girth age (girth B) and rubber production (yield) for 25 *Hevea* genotypes (clones) grown in Votuporanga, São Paulo State, Brazil.

Trait		Age c	ombination	(years)	
	1 and 6	2 and 6	3 and 6	4 and 6	5 and 6
Girth A	0.5498	0.6773	0.7723	0.8005	0.9420
Girth B	0.5114	0.5685	0.8320	0.9046	0.9471
Yield	0.3451	0.5988	0.5185	0.6556	0.8311

All values in table are significant at p < 0.01.

Hevea clones could be done on a girth basis after 24 months.

Our study shows that in São Paulo it may be unwise to select Hevea clones at age 1, although the 2 to 4-year growth data can be used differentially in the selection process. Selection of the sets of the best 30, 15, 10 and 5 clones from the available 68 clones at a given age was generally accompanied by a descending order of percentage success (Table 4). Thus, for a given level of percentage success in selection, the set of clones to be selected at a relatively young age needs to be large, while selection of a small set needs a higher selection age. Therefore, it is suggested that: a) to have the best 30 (*i.e.* 44% of the available germplasm) clones at age 6 select the set of the best 36 (53%) clones at age 2; b) to have the best 15 (22%) clones at age 6 select the set of the best 20 (29.41%) clones at age 3; c) to have the best 5 (7.35%) clones at age 6 select the set of the best 8 (11.76%) clones at age 4; and d) to have the best 3 (4.4%)clones at age 6 just select the set of the best 3 (4.41%)clones at age 5. More than 80% of the clones targeted on a girth A or yield basis, and more than 76,7% clones targeted on a girth B basis, were found to be correctly identified during steps (a) to (d). Nevertheles, the 5 highest-ranking age 6 clones (viz. clones with identification numbers 68, 67, 66, 65 and 64) on a girth A (Figure 1 and 2) or girth B basis were found to be invariably present in the sets of clones selected at (a), (b) and (c).

One is often faced with the task of making early selection of clones on the basis of a field test to give interim recommendation about the choice of clone for further multi-location testing (large-scale trials). The age of early selection needs to be sufficently long to effect a reasonable level of selection accuracy. After making early selection, vegetative multiplication of the selected clones is started and it is continued for many years in order to build the budwood stock nursery which is large enough to supply buds to all rubber producers on a sustained basis. To shorten the time involved in this two-stage process of (i) selecting the most promissing clones, followed by (ii) multiplying their germplasm, an alternative approach involving concurrent short-listing and multiplication of clones is suggested. According to this approach, early selection of the best 60% of clones should be done at age 2 of the field trial

Tr	Sa		Ct		Cs	Ре
	Years	Ν	%	Ν	%	%
(a)	(b)	(c)	$(d)^{1}$	(e)	$(f)^2$	$(g)^{3}$
Girth A	2	5	7.35	3	4.41	60.00
Girth A	3	5	7.35	4	5.88	80.00
Girth A	4	5	7.35	4	5.88	80.00
Girth A	5	5	7.35	4	5.88	80.00
Girth A	2	10	14.70	5	7.35	50.00
Girth A	3	10	14.70	8	11.76	80.00
Girth A	4	10	14.70	8	11.76	80.00
Girth A	5	10	14.70	10	14.70	80.00
Girth A	2	15	22.06	7	10.29	46.67
Girth A	3	15	22.06	10	14.70	66.70
Girth A	4	15	22.06	12	17.64	80.04
Girth A	5	15	22.06	12	17.64	80.04
Girth A	2	30	44.12	21	30.87	69.93
Girth A	3	30	44.12	25	36.75	83.25
Girth A	4	30	44.12	27	39.69	89.91
Girth A	5	30	44.12	28	41.16	93.24
Girth B	2	5	7.35	3	4.41	60.00
Girth B	3	5	7.35	3	4.41	60.00
Girth B	4	5	7.35	4	5.88	80.00
Girth B	5	5	7.35	4	5.88	80.00
Girth B	2	10	14.70	7	10.29	70.00
Girth B	3	10	14.70	6	8.82	60.00
Girth B	4	10	14.70	7	10.29	70.00
Girth B	5	10	14.70	9	13.23	90.00
Girth B	2	15	22.06	10	14.70	66.70
Girth B	3	15	22.06	12	17.64	80.04
Girth B	4	15	22.06	12	17.64	80.04
Girth B	5	15	22.06	13	19.11	86.71
Girth B	2	30	44.12	23	33.81	76.59
Girth B	3	30	44.12	25	36.75	83.25
Girth B	4	30	44.12	29	42.63	96.57
Girth B	5	30	44.12	30	44.10	100.00
Yield	2	5	7.35	3	4.41	60.00
Yield	3	5	7.35	2	2.94	40.00
Yield	4	5	7.35	3	4.41	60.00
Yield	5	5	7.35	4	5.88	80.00
Yield	2	10	14.70	5	7.35	50.00
Yield	3	10	14.70	8	11.76	80.00
Yield	4	10	14.70	8	11.76	80.00
Yield	5	10	14.70	10	14.70	100.00
Yield	2	15	22.06	7	10.29	66.70
Yield	3	15	22.06	12	17.64	80.04
Yield	4	15	22.06	12	17.64	80.04
Yield	5	15	22.06	14	20.58	96.38
Yield	2	30	44.12	23	33.81	76.59
Yield	3	30	44.12	27	39.69	89.91

 $^{1}(d) = (c) \ge 100/68$; $^{2}(f) = (e) \ge 100/68$ and $^{3}(g) = (e) \ge 100/(c)$.

30

30

Yield

Yield

4

5

Tr: Trait for selection of clones. Sa: Selection age. Ct: Clones targeted for selection at selection age. Cs: Clones successfully selected at selection age with the aim of selection for age. Per: Per cent success of selection at selection age. N: Number. %: % of 68.

44 12

44.12

27

28

39 69

41.16

89 91

93.33

Table 4 - Extent of success in correctly selecting a set of 5, 10, 15 or 30 best clones for age 6 (girth) and age 6 (yield) by making selection at age 2, 3, 4, 5 or 6 respectively.

and their vegetative multiplication should be immediately started in the nursery. Short-listing of the best 33% (of the original number of clones under test) clones should be done from within the above set of clones select at age 3 on the basis of (a) age 3 growth data in field trial and (b) grafting and survival performance of buds during the previous year in nursery. Propagation of the rejected clones should be stopped by uprooting and eliminating the rejected clones (roguing). This process of the simultaneous roguing of clones on the basis of current evaluation and the vegetative propagation of select clones can be continued in the subsequent years until the desired selection intensity and number of clones (ramets) are archieved, the best 5% of clones being shot-listed at age 5.

Early selection might result in some bias against clones which grow relatively slow in the initial stage but catch up with others at the later stages. However, Figure 1 and Table 2 suggest that such clones would be very few, and, in fact, such clones would lag behind others if planted in intimate mixture with other clones at high densities. Since the objective of a Hevea improvement program is to maximize gains per unit time it is undesirable to delay making interim recommendation about choice of clones and starting their vegetative multiplication on the basis of early evaluation just because sporadic clones would be disfavored. Final recommendation on the choice of clones for later deployment in plantations and for serving as parents for the next generation of Hevea improvement can be done at year 4 because in São Paulo Hevea normally does not flower before this age.

Conclusions

Early selection of *Hevea* clones for tapping panel opening at age of 6 years can be done effectively at age 4. Selection at age 2 is relatively less accurate but it results in early propagation of selected clones for further testing or deployment. To combine the advantages of these two early selection ages we suggest a differential selection strategy, *i.e.* the best 60% of clones of the original set should be selected at age 2,33% at age 3, 13% at age 4 and 5% at age 5. The sets of clones to be selected at age 3 and later should be contained in the sets selected in the perceding years. Multiplication of the selected clones should be started at the end of the second year of the field trial and continued in successive years. Selection at age 2 should be done on the basis of growth, preferably vigor (girth A) in the field trial.

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