

PILOCARPINE CONTENT AND MOLECULAR DIVERSITY IN JABORANDI

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ABSTRACT: Pilocarpine is an imidazol alkaloid exclusively found in *Pilocarpus* genus and *P. microphyllus* accumulates its highest content in the leaves. There is no report in the literature on the variability of the pilocarpine content in this genus. A population of 20 genotypes of *P. microphyllus* from the state of Maranhão, Brazil, was analyzed for Random Amplification of Polymorphic DNA (RAPD) markers and pilocarpine content. Although it was not possible to establish any correlation between these features, the absence or presence of some markers could indicate in some genotypes a possible association with the content of the alkaloid.

Key words: RAPD, *Pilocarpus jaborandi*, molecular diversity, molecular markers

CONTEÚDO DE PILOCARPINA E DIVERSIDADE MOLECULAR EM JABORANDI

RESUMO: Pilocarpina é um alcalóide imidazólico encontrado exclusivamente em plantas do gênero *Pilocarpus*, sendo que as folhas de *P. microphyllus* acumulam o maior conteúdo deste alcalóide. Não há na literatura nenhum relato sobre a variabilidade do conteúdo de pilocarpina nesse gênero. Uma população de 20 plantas de *P. microphyllus* do estado do Maranhão, Brasil, foi analisada por marcadores Aplicação de DNA polimórfico randomica (RAPD) e quanto ao conteúdo de pilocarpina. Apesar de não ter sido possível estabelecer uma associação entre as variáveis estudadas, a ausência ou a presença de alguns loci marcadores em certos genótipos puderam ser associados ao teor do alcalóide.

Palavras-chave: RAPD, *Pilocarpus jaborandi*, diversidade molecular, marcadores moleculares

INTRODUCTION

Pilocarpine is an imidazol alkaloid found in plants of the genus *Pilocarpus*. Plants of this genus are designated by the name jaborandi but only *P. microphyllus*, which accumulates the highest pilocarpine content, is considered the true jaborandi (Pinheiro, 1997; Vieira, 1999). Jaborandi grows as a shrub and it is found in the understory of the pre-Amazonian rain forest and occurs more intensively in the state of Maranhão (Vieira, 1999; Pinheiro, 2002).

Pilocarpine has important pharmaceutical properties. It is used to reduce the intraocular pressure in the treatment of glaucoma (Migdal, 2000), as a stimulant of salivation and perspiration, and recently has been prescribed for the treatment of xerostomia, which is the reduction of saliva production (Davies et al., 2001).

In spite of the importance of the plant and the pharmacological activity of pilocarpine, only a few reports have been published on the content of this alka-

loid in *Pilocarpus* (Andrade-Neto et al., 1996; Avancini et al., 2003). A preliminary study was carried out on the genetic diversity in jaborandi accessions using Random Amplification of Polymorphic DNA (RAPD) (Moura et al., 2003). Curiously, the highest genetic variability was observed within populations and not among populations. Pilocarpine was not determined in these plants.

The above-mentioned results prompted to the analysis of a population of *P. microphyllus* growing in a greenhouse which was obtained from seeds from a private farm in the Maranhão State, Brazil. Jaborandi is domesticated and is now cultivated as a crop, to our knowledge not having been submitted to any breeding program. Although having been explored in Brazil for three decades, only in 1989/90 a program started to domesticate the plant. Domestication was accomplished from the knowledge accumulated by peasants, when all leaves involved in pilocarpine extraction were harvested from jaborandi plants in the forest. This

know-how was mainly based on the observation that some collection areas contained plants with more pilocarpine than others and such plants presented larger leaflets. However, this might be related to soil fertility since the leaves containing higher contents were obtained from areas of pre-Amazonian rain forest while those with lower contents were either from 'cerrado' (Brazilian savanna) or areas along the coast (Pinheiro, 2002). Avancini et al. (2003) observed that nutrient-deprived jaborandi seedlings produced less pilocarpine in the leaves.

The plants growing in the greenhouse of this study presented different height, leaf shape, branch pattern and leaf color. Therefore, the aim was to investigate the content of pilocarpine in leaves of 20 jaborandi genotypes selected on the basis of their phenotypes, and try to establish a possible association with the genetic diversity revealed by molecular markers.

MATERIAL AND METHODS

Plant material

Seeds of *Pilocarpus microphyllus* Stapf. ex Holm. were germinated in vermiculite. A voucher specimen was deposited in the Herbarium of the State University of Campinas (UNICAMP) Campinas, SP State, Brazil. When the seedlings reached approximately 5 cm (five months old) they were transferred to 0.5 L plastic pots containing a mixture of sand and soil (1:1, v/v) and each pot received complete nutrient solution (50 mL) twice a week (Hoagland & Arnon, 1950). The 20 genotypes of this study were selected according to their height, branching pattern, leaflet size, and the color of young leaves. Two lateral leaflets positioned in the middle of the third composite leaf (from the apex) were harvested and one was used for extraction and the other for pilocarpine and anthocyanin determinations. The length of the selected leaf and leaflet were measured. Leaflets for chemical determinations were freeze-dried before extraction.

Extraction and analysis procedures

Pilocarpine was extracted and analysed by High Performance Liquid Chromatography (HPLC) using a standard method (Avancini et al., 2003). The absorbance of anthocyanins was determined in methanol extracts containing 1% HCl prepared from the same amount of dried material (Rabino & Mancinelli, 1986).

DNA isolation and amplification

Total genomic DNA was extracted from 0.3-0.4 g fresh leaves ground in liquid nitrogen using 0.1 M Tris-HCl, 1.25 M NaCl, 0.02 M EDTA, 2% mixed alkyltrimethylammonium bromide, and 1% β -mercaptoethanol. After 90 min incubation at 65°C with

slow stirring, the solution was treated twice with an equal volume of chloroform/isoamylalcohol (24:1) and the supernatant treated with RNase. DNAs was precipitated with 0.8 vol. of isopropanol and after washing with 70% ethanol, it was vacuum dried and dissolved in 200 μ L of TE buffer. DNA was used in Polymerase Chain Reaction (PCR) reactions carried out in a 25 μ L reaction mixture containing 1X reaction buffer, 1.5 mM MgCl₂, 10 ng template DNA, 1.0 μ M primer, 100 μ M of each dNTP and 1 unit of Taq polymerase. The following conditions were used: 1min at 94°C, 1min at 35°C and 1.5 min at 72°C, repeated 42 times. Amplified fragments were separated on 1.5% agarose gel and visualized with ethidium bromide. About 100 primers (Operon Technology, US) were tested for variation and polymorphism in 20 genotypes of the jaborandi tree.

Data Analysis

Only clearly amplified polymorphic fragments were analyzed. The RAPD fragments were scored as present (1) or absent (0) and these data used to calculate an index of genetic similarity between pairs of genotypes using the DICE coefficient of genetic similarity: $SD=(2a/n-d)$, where a is the number of positive matches, n is the sample size and d is the number of negative matches. A hierarchical classification was obtained from these coefficients using the UPGMA method (unweighted pair-group method using arithmetical averages) of aggregation and bootstrap values. All statistical analysis were performed using NTSYS-pc (Numerical Taxonomy and Multivariate Analysis System for personal computers), version 2.1 software (Rohlf, 2000).

RESULTS AND DISCUSSION

The content of pilocarpine in the 20 genotypes was variable, and it was possible to separate them in four groups (Table 1). Group 1 was formed by genotypes containing less pilocarpine, with pilocarpine below 50 μ g g⁻¹ f.w. A second group was formed by genotypes 5, 8, 9, 11, 16, 18 and 20 with pilocarpine contents within the range 60 to 90 μ g g⁻¹ f.w. pilocarpine. Group 3, by genotypes 6, 14, 15 and 19 with pilocarpine within the range 100-200 μ g g⁻¹ f.w. and the group 4 with three genotypes containing more than 200 μ g g⁻¹ f.w. pilocarpine in the leaves.

There are few reports in the literature to have this data compared. The amounts observed in the 20 genotypes are in agreement with previous values reported for jaborandi plants grown in greenhouse (Avancini et al., 2003). Plants in the field seem to accumulate more pilocarpine, with a mean values between

Table 1 - Pilocarpine, anthocyanin and leaf/leaflet length of the 20 tree genotypes of *P. microphyllus* grouped according the alkaloid content.

Tree genotype	Group	Pilocarpine $\mu\text{g g}^{-1}$ dry weight	Anthocyanin (Abs \times 1000)	length	
				Leaf	Leaflet
4	1	16.3	38	7.1	1.8
17	1	19.8	16	5	1.1
12	1	24.1	1	7	2.5
3	1	24.1	77	8.5	2.5
10	1	29.6	262	6.5	1.6
13	1	39.4	52.0	11.1	2.5
11	2	63.8	66	10.3	1
8	2	67.5	113	9.5	2.5
5	2	71.8	222	11.5	3
16	2	74	36	8.5	1.5
18	2	84.5	87	7.1	2
20	2	85	51	9.1	2
9	2	86.4	172	11.2	3.2
14	3	129	55	12.4	2.5
15	3	130.2	41	11.8	3.1
19	3	145.1	52.0	10	2.6
6	3	166.1	220	10.5	2.5
2	4	215.7	620	7.8	2.3
1	4	232.5	148	10.3	1.5
7	4	235.9	589	8	2

400-500 $\mu\text{g g}^{-1}$ (Sousa et al., 1991; Pinheiro, 1997; 2002), indicating that the content of pilocarpine might respond to environmental conditions. Indeed, it has been shown that secondary metabolites may have their concentration in plants strongly affected by biotic and abiotic factors (Waller & Novack, 1978; Dixon & Paiva, 1995). Avancini et al. (2003) observed that mineral, salt and oxygen stresses affect the content of pilocarpine in jaborandi leaves. In the evolution process mutations usually cause alterations in the biosynthetic routes of secondary metabolites modulating the final content of the compounds (Haslam, 1986). Such variations are very well illustrated in caffeine-containing plants where the content of this alkaloid may vary even in cultivars of the same species (Mazzaferri et al., 1994).

Out of the 100 tested primers, 33 were selected for further studies on the basis of amplification patterns (Figure 1). These 33 profiles were analyzed in detail and it was observed that 18 primers (OPAC-1, OPAC-8, OPAC-10, OPAC-11, OPAC12, OPAC-15, OPAC-17, OPY-20, OPAB-3, OPI-16, OPE-12, OPAL-9, OPX-1, OPJ-1, OPZ-9, OPX-18, OPX-10 and OPX-7) revealed genetic polymorphisms. When this variation was compared with the results of total pilo-

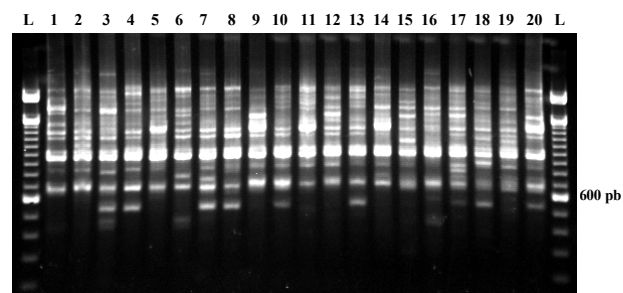


Figure 1 - Random Amplification of Polymorphic DNA (RAPD) amplification of twenty jaborandi genotypes using the primer OPX-18 from Operon Technology. Lateral lanes are 100 pb ladder. The 600 pb fragment is indicated.

carpine of the 20 jaborandi genotypes, it was observed that 9 of them (OPAC-1, OPAC-8, OPAC-10, OPAC-11, OPAC12, OPAC-17, OPY-20, OPAB-3, OPX-18) could be used for selection of RAPD molecular markers to differentiate the 20 genotypes regarding their amounts of pilocarpine.

The molecular markers OPAC-1₈₀₀, OPAC-8₈₀₀ and OPAC-11₁₀₀₀ were present in genotype 1 and absent in all other genotypes, therefore they might be considered as RAPD markers for these genotypes. Molecular marker OPAC-8₂₀₀₀ presented polymorphism

in all 19 tree genotypes but was not found in tree specimens of genotype 6, which produce significant amount of pilocarpine and hence the absence of this particular DNA fragment could be treated as marker for this genotype. Molecular marker OPAC-10₁₁₀₀ had a strong amplification in genotype 15 but was also observed in genotype 2 and 19, and these three genotypes have significant levels of pilocarpine. Similarly, OPAC-12₁₁₀₀ had amplification in genotype 2, 15 and 19. OPAC-17₄₀₀ showed very good amplification in genotype 14 as well as in tree genotypes 2 and 7. OPX-18₁₂₀₀ was amplified only in tree genotype 7 and absent in all other 19 genotypes and hence could be selected as molecular marker for this highest pilocarpine producing genotype. Another fragment OPY-20₇₂₅ was detected in genotype 2 and 19. It was also noticed that OPAB-03₅₅₀ was found only in the lowest pilocarpine producing genotype (4) and therefore it might be treated as molecular marker for this low pilocarpine producing tree.

Before collection for the analysis of pilocarpine and anthocyanin contents, the leaves had their length measured and a leaflet at the center of the composite was also measured. Only the third expanded leaf from the apex was used. One may argue that there might be an opposite relationship between leaf length and tree genotypes with low pilocarpine as seen for genotypes 7 and 2 with small leaves and high pilocarpine contents. However, group 3 showed the highest mean leaf length. Furthermore, genotype 1 with 232.5 $\mu\text{g g}^{-1}$ of pilocarpine had a long leaf. At the other extreme, short leaves were those with the lowest contents. The correlation coefficient calculated for these characteristics were: pilocarpine \times anthocyanin = 0.643, pilocarpine \times leaf length = 0.305, pilocarpine \times leaflet length = 0.103, anthocyanin \times leaf length = -0.113, anthocyanin \times leaflet length = 0.06, leaf length \times leaflet length = 0.568. Although a good relationship was observed between leaf and leaflet lengths, leaf length correlated better with pilocarpine. One may also argue that young leaves usually present more anthocyanin which decreases with aging. However, leaf length, that could also be an indication of leaf age, did not correlate well with anthocyanin. Furthermore, for all genotypes, leaves of the same age were used. Therefore, it seems that the anthocyanin content might not be correlated only to a developmental stage but also to genetic variability.

From the presence/absence (binary data) of the polymorphic *loci* the genetic similarity was calculated for the genotypes resulting in mean, minimum and maximum values of 0.66, 0.51 and 0.77, respectively. The most similar genotypes were the genotype-pairs J7/J10, J3/J7, J13/J17, J2/J19 and J10/J13 while the

most different were J1/J6, J4/J15, J4/J18 and J8/J9. From the genetic similarities among the genotypes and using the hierarchic classification (UPMGA aggregation method) a dendrogram was obtained (Figure 2) from which it was possible to define four main heterotic groups (A, B, C and D), represented by the genotypes J1 and J20 (A), J2, J3, J5, J7, J8, J10, J11, J12, J13, J14, J16, J17, J18, J19 (B), J9 and J15 (C) and J4 and J6 (D).

The results of the genetic diversity study using RAPD markers revealed four groups genetically distinct, suggesting a genetic structure. However, it was not possible to establish a relationship between the genetic diversity revealed by RAPD markers and pilocarpine and anthocyanin contents.

In a previous study, Moura et al. (2003) analysed several accessions from Maranhão e Pará states and showed a significant variation among the collection sites. Only individuals from a single place were grouped together with a great variation for the other accessions, indicating that jaborandi presents more variability among plants from the same area. They also analysed 10 accessions from the Merck farm and a mean similarity value of 0.72 was obtained. Such similarity varied from 0.55 to 0.91 when considering all the accessions.

The genus *Pilocarpus* has a wide distribution in the Brazilian territory, ranging from the northern state of Pará to the southern state of Rio Grande do Sul (Joseph, 1967) and *P. microphyllus* has been cited

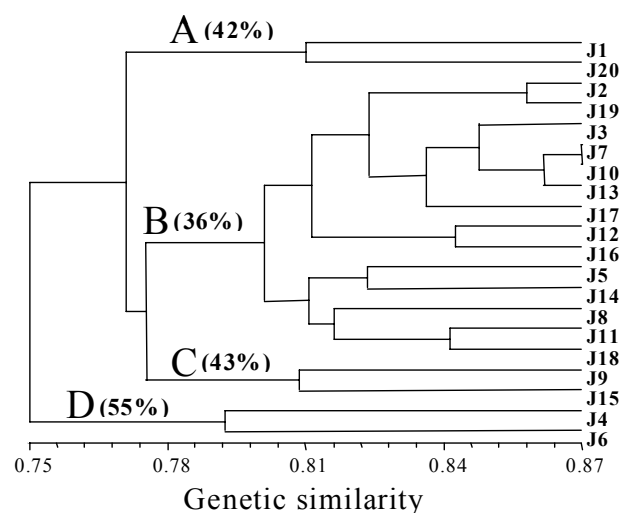


Figure 2 - Dendrogram unweighted pair-group method using arithmetical averages (UPGMA) obtained from similarity matrix based on DICE coefficient estimated among 20 jaborandi genotypes based on 170 Random Amplification of Polymorphic DNA (RAPD) markers. Values in parenthesis indicate the bootstrap values.

to be more restricted to the northern state of Maranhão (Vieira, 1999; Pinheiro, 2002). Unfortunately, there is no information available on the reproductive mechanism of this species. *P. microphyllus* is hermaphrodite and although several plants presenting hermaphroditism have cross pollination due to different maturation times of pollen and stigma, it is not sure that this really happens with this species and that this could be the reason for the observed variation.

The differences observed among the 20 genotypes here studied indicate intra-specific genetic variability. Since these plants were grown under the same environmental conditions in a greenhouse, the genetic diversity observed is probably related to the collection sites these genotypes came from. Although these plants came from the same place, information (Pinheiro, 2002) indicates that there was no breeding program for the selection of the most productive plants in terms of alkaloid content. Therefore, the plants growing at the original farm may have been collected from different localities and genetic variation was probably maintained.

Molecular markers are more stable and informative than isoenzymes. They are currently used to study the genetic diversity of several species (Virk et al., 1995) and they can be used more efficiently to examine the genetic diversity of plant germplasm (Colombo et al., 2000). Although it was not possible to establish a direct relationship between RAPD molecular markers and pilocarpine content, it was observed that the presence or even the absence of some markers might be used to select plants that tend to accumulate more pilocarpine. Therefore, a study on the cosegregation of these markers and pilocarpine content could provide more information on their genetic linkage.

ACKNOWLEDGEMENTS

S.S. Sandhu thanks CNPq (Brazil) and TWAS-UNESCO (Italy) for the financial support to visit IAC, Campinas, Brazil. I.N. Abreu thanks for the post-doctoral fellowship of FAPESP and P. Mazzafera thanks for the research fellowship of CNPq. This work was also partially granted by FAPESP and CNPq. To the Merek Company for providing the genetic material.

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Received June 20, 2005

Accepted July 21, 2006