Identification of *Ficus religiosa* by DNA SCAR Marker

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**ABSTRACT**

*Ficus religiosa* Linn. is a popularly used bark drug in Indian system of medicine, an ingredient of *Panchatvalaka* (Five bark drugs). Because of morphological similarity adulteration is common with bark where quality therapeutics is always uncertain. DNA markers are reliable in this regard as the genetic composition is unique for each species and is not affected by age, habitat, environmental factors and physiological conditions. Hence DNA SCAR marker development of *F. religiosa* was carried out. **Methods:** Fresh bark samples of *F. religiosa* along with samples of *Panchatvalaka* tree were collected from various parts of India. DNA was isolated and purified using column. The bands that were found only in *F. religiosa* species and not in others were selected for further processing. The unique band thus obtained was further purified from the gel and used for cloning into T vector. The white colonies obtained were inoculated in LB media with antibiotic and plasmid isolation was done. The plasmids with retarded mobility were selected for confirmation with PCR using M13 primers. Positive clone was further purified and sequenced. The sequence obtained was used for designing SCAR primers. **Results:** For identification of *F. religiosa* by DNA marker DNA marker has been developed for *F. religiosa*. **Conclusion:** SCAR marker developed can be used to identify and differentiate the plant wherever it is used in preparation of medicines.

**KEYWORDS** *Ficus religiosa*, DNA SCAR marker, CTAB method, RAPD.

**PICTORIAL ABSTRACT**

![Graph](Image)

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Ficus religiosa* Linn. is a large glabrous tree, without aerial roots from the branches, with sessile figs in axillary pairs, found all over India as an avenue tree[7]. It is held sacred by Hindus and Buddhists. It belongs to family *Moraceae* and the genus *Ficus*, which is a large genus of trees or shrubs and about 65 species are reported from India[6]. Bark of *F. religiosa* is popularly used in Indian system of medicine, for the treatment of many disease conditions. It is said to be astringent, antiseptic, laxative, haemostatic, vaginal disinfectant and used in diabetes, diarrhoea, leucorrhoea, menorrhagia, nervous diseases and also in skin problems[9]. Bark contains beta-sitosterol, 1-D-glucosid, Vit K, n-otacosanol, methyl oleanolate, lanosterol, stigmasterol, lupen-3-one[9]. Paste of the powdered bark is used as absorbent in inflammatory swellings. In Ceylon the juice of the bark is used as a mouth wash for toothache and for strengthening the gums. Water in which the freshly burnt ashes of the bark have been steeped is said to cure, obstinate cases of hiccup and stop vomiting gums. Water in which the freshly burnt ashes of the bark have been steeped is said to cure, obstinate cases of hiccup and stop vomiting gums. Water in which the freshly burnt ashes of the bark have been steeped is said to cure, obstinate cases of hiccup and stop vomiting gums. Water in which the freshly burnt ashes of the bark have been steeped is said to cure, obstinate cases of hiccup and stop vomiting gums.

Identification and quality assurance of botanical material is an essential requirement in herbal medicine to ensure reproducible quality of herbal medicine, which contributes to its safety and efficacy[8]. Most of pharmacopoeia’s, quality control guidelines suggest macroscopic, microscopic evaluation, chemical profiling for herbal drug standardization. Correct chemotype of plant can provide clinical efficacy. DNA markers are reliable in this regard as the genetic composition is unique for each species and is not affected by age, habitat, environmental factors and physiological conditions[11]. Various types of DNA based molecular techniques are utilized to evaluate DNA polymorphism such as, DNA based molecular techniques, hybridization based methods, PCR based methods and sequencing based methods[12]. Random Amplified Polymorphic DNA (RAPD) markers are DNA fragments from polymerase chain reaction amplification of random segments of genomic DNA with single primer of arbitrary nucleotide sequence[13]. The limitation of RAPD is that it is not possible to distinguish whether a DNA segment is amplified from a locus that is heterozygous or homozygous. To overcome these problems SCAR(sequence characterized amplified region) marker is developed by cloning and sequencing RAPD marker(unique band) and further designing specific primers for the sequence obtained. These SCAR primers are used to identify and differentiate the strain of interest from others[14].

Aim of present work is to develop SCAR primers for *Ficus religiosa* that is used in preparation of *Panchatvalaka* along with *F. bengholensis*, *F. glomerata*, *F. lacor* and *Thespesia populnea* bark powders. This would help in quality control stage of the formulations prepared from this important medicinal plant.
1. Plant material
Fresh bark samples of *F. religiosa* tree were collected from various parts of India. Plant samples were authenticated and Bark samples were frozen at -70°C till further use.

2. Preparation of bark powder
Bark samples were crushed into small pieces using autoclaved pestle and mortar and powdered thoroughly using dry ice intermittently. Individual pestle and mortar were used for each sample and collected the powder in a fresh container till further use.

3. DNA isolation
DNA isolation was attempted with different methods including urea method, DNAzol method, CTAB method and modified CTAB method. CTAB method DNA isolation was found better for RAPD PCR, though the yields were better with other methods. The crushed powder was treated with 70% ethanol before DNA extraction. Water wash to the powder was given to remove soluble pigments and compounds that may interfere in the experiment[15].

4. CTAB method DNA extractions
Ethanol and water washed bark powder was homogenized with CTAB buffer thoroughly. Beta Mercapto ethanol was added and incubated for 90 minutes at 60°C in a water bath. This was centrifuged at 10,000 rpm for 15 minutes and supernatant was decanted into fresh tubes. Equal volume of 100% chloroform was added and mixed well for 10 minutes by inverting tube. This was, centrifuged at 10,000 rpm for five minutes. Separated aqueous layer was transferred in to fresh tube and added equal volumes of iso-propyl alcohol. Centrifuged at 10,000 rpm for 15 minutes. Discarded supernatant and washed DNA pellets with 0.5 ml of 70% ethanol and centrifuged at 10,000 rpm for 5 minutes. Discarded ethanol and air dried DNA pellet. DNA was suspended in 1X TE buffer. The DNA was further purified using silica membrane based column and quantitated on agarose gel[16].

5. RAPD PCR
Templates of all the 5 species were used for RAPD. PCR was set up using master mix consisting of 100µM each of dNTPs, 100mM KCl, 1.5mM MgCl2, 0.1% each of tween 20 and Nonidet P40, 15p moles of random primer and1u of taq polymerase in 40 µl volume. 100ng of template was used. PCR Cycle conditions were ,initial denaturation at 94°C for 5 minutes, denaturation for 30 seconds, annealing for 1 minute at 45°C, extension for 90 seconds at 72°C and final extension for another 7 minutes.

After amplification, the samples were run in 2% agarose gel using DNA marker with 0.1, 0.2, 0.3, 0.6, 1.0, 1.5, 2, 2.5, 3 and 3.5 Kb fragments. Unique band of ~900bp was observed for *F. religiosa* which was not seen in other species. This fragment was gel purified, amplified and used for cloning into T vector. Clones were confirmed by amplification using vector primers and was sequenced. Sequence obtained was analysed and primers were designed to give ~550bp product for this region. Randomly selected ficus samples were used for checking the efficacy of the primers designed[17].

Samples were used from all the five *Panchavalkala* plant species for RAPD (Figure 1). The bands that were found only in religiosa species and not in others were selected for further processing. The unique band thus obtained was further purified from the gel and used for cloning into T vector. The white colonies obtained were inoculated in LB media with antibiotic and plasmid isolation was done. The plasmids with retarded mobility were selected for confirmation with PCR using M13 primers. Positive clone was further purified and sequenced. The sequence obtained was used for designing SCAR primers. Highlighted sequence and primer designed is shown in Figure 2 and Table 1.

The primer selectively amplifies only *Ficus religiosa* samples (Figure 3). Thus SCAR marker for *F. religiosa* is successfully developed and can be used to identify and differentiate the same wherever it is used.

<table>
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<th>Table 1: The primers designed</th>
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<td>ReFP</td>
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<td>ReRP</td>
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6. Sequence data

AGTCTCGGTAGCTCTCCGGCCGCATGTTCCCTGTGCTGAGCAGAGAGA
GTAATGGAGCGCGACGGGTATACACCCTCCCACGGAGTACCTTCCGGC
TTCCCTCTCCTGCTCTGCGACTCTGCTCTGACCGCTGAGGACCTGCT
AGCTGCTGCCCTTGGCCTGAGGAGAGCTGCTGCTGTAGGTCGG
CGTTCCGAGATTCCTGCGGAGATCCGCGCTGGTCTACGGGAGTCTAT
GTGCTGCGACGGACGGACGGAGAGGGGAGTATACGCGCACTCTCCGCA
GGTTTTTCTACCCCCCGGACGGATGCTGCTGCTGACCTCAGG
CTGCAACGCGATCTTAGGACGGGCGGTAGCTTGTGTCGGA
GTCCTGAGCGCGCTGTTGGAACCGTGGTCGTGTTGGATGCTTCCGA
AGCGCAGTGGCTGGACGCTGTTAGTTGACGCTGACCGCTATAGG
TTCTTGAGATGCTGCTGCCGGCAGGGAAGAAGCATATTTGTGACCTGAA
TCGCTGCTGCTGTGAGGTTCTTGAGCAGGTGGTTGGTGAATGACATTTGA
GAAAATGAGAGATTACCGACAGAAAATTTGAGAATCTGATGAAAGATTG
ATCCTGAGAGACTCATTGCTTCAAAATCCGCCGCTCCTCCTAGCAGAAAATCTG
GTGAAAAAGCTGCGGAGAGAAATTGGAGTACCCGGAAAGCAGTAAAT
TCCACCCAGATCTATGCGAAGCTGCAAGAGCAGGACTTTGGGT
GATGAGAAATGCTGCGAGCAATGAATATGAGCAGAGAAGGATCATG
CCGCCGGATATCTGACTGCGCGCGCTGAGTGCACCATATGGGAG
AGTCACCCAGGCGTGGATGCAG

Clarity and reliability of SCAR markers proves them best techniques in current herbal drug standardization. SCAR primes of *Ficus religiosa* can be used to authenticate the sample, preventing its admixture with other bark drugs.

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**Conflict of interest:** Authors declare no conflict of interest.

Quality control parameters with modern scientific techniques are of prime importance. The routine methods of herbal drug standardization address pharmacognostical, chemical, biological, biopharmaceutical and molecular approaches[18]. Isolation of purified DNA from plant bark is challenging because of secondary metabolites and other compounds, which interfere with isolation process and purity[19]. In this study DNA isolation was attempted with different methods and among these, CTAB method DNA isolation was found better for RAPD PCR, though the yields were better with other methods. The DNA obtained was further purified using silica membrane based column.

RAPD- a PCR based reaction introduced by Williams et al., which amplify segments of DNA those are essentially unknown to the researcher[20]. Still the fragment polymorphisms produced by RAPD-PCR amplifications are not always reproducible. This limitation can be overcome by converting RAPD’s into sequence characterized amplified region (SCAR)[21].

In this study, templates of all the 5 species were used for RAPD. After amplification, the samples were run in 2% agarose gel using DNA marker with 0.1, 0.2, 0.3, 0.6, 1.0, 1.5, 2, 2.5, 3 and 3.5 Kb fragments. Unique band of ~900bp was observed for *F. religiosa* which was not seen in other species and this fragment polymorphisms produced by RAPD-PCR amplifications were better with other methods. The DNA obtained was sequenced and primers were designed. The SCAR primers thus obtained were further validated using various ficus samples.

**Figure 3. Amplification using SCAR primers**

(Lane1: 500bp ladder, lane2-5 racemosa, lane6-9religiosa, 10-13 benghalensis, 14-16 lacor and 17-19 thespeia.)

**Contributors:** Dr. Suma Venkatesh Mallya collected plant samples and carried out DNA extraction, RAPD and SCAR marker at Aristogene Bangalore along with Dr.Vishwanatha. Dr.KN Sunil Kumar edited the manuscript giving suitable suggestions, to make it acceptable form. Suchitra Prabhu helped in authentication of plant samples.

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