

Sandalwood spike disease: a brief synthesis

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Abstract

Sandalwood spike disease (SSD) is likely to be the most destructive of known diseases to infect Indian sandalwood, *Santalum album* L., which is valued for its wood and scented essential oil derived from the heartwood. Even though the history of research on SSD extends for well over a century, most of the earlier work was primarily descriptive in macromorphological aspects. In 1969, a breakthrough was achieved by three independent research groups that confirmed the causal agent of SSD to be a phytoplasma. Despite this discovery, the identity of the causal agent was only confirmed by a single group using RFLP analysis and nested PCR of 16S rDNA. The phytoplasma that was identified showed high sequence similarity to aster yellows subgroup 16SrI-B. Tissue culture to derive disease-free clones and transgenic strategies to derive disease-resistant trees may be the only realistic solutions for ridding the limited *S. album* forests of SSD. This brief synthesis aims to stimulate greater interest in SSD, especially since sandalwood production has expanded to the Chinese mainland and to Australia.

Key words: Indian sandalwood, leaf curl, phytoplasma, *Santalum album*, spike, yellow disease.

Abbreviations: ELISA, enzyme-linked immunosorbent assay; RFLP, restriction fragment length polymorphism; SSD, sandalwood spike disease.

Introduction and history

Indian sandalwood (*Santalum album* L.), a highly valuable hardwood species of the Santalaceae, is popular for its fragrant wood and essential oil, and has importance in traditional and pharmaceutical medicine, as well as in the cosmetic and perfumery industries. Therefore, sustainable cultivation methods are required for the large-scale propagation and production of disease-free *S. album* plants. This can be successfully achieved using biotechnological approaches such as tissue culture (Teixeira da Silva et al. 2016). The survival and growth of seedlings, or of tissue-cultured plantlets at the *ex vitro* stage, even those with an established root system, can be a difficult task due the hemiparasitic nature of *S. album*, which requires one or more specific hosts (Teixeira da Silva et al. 2016). There is evidence that *S. album* is susceptible to various insect pests (Sundararaj, Muthukrishnan 2011) and to sandalwood spike disease (SSD) (Arun Kumar, Joshi 2012; Kumar 2014). SSD of *S. album* was first reported in Coorg state, South India in 1899 (McCarthy 1900; Barber 1903) with over a million trees having been removed as a result of SSD from Mysore and Coorg states during 1903-1916. SSD, which is a critical disease of *S. album*, has been a topic of fascination for at least 100 years (McCarthy 1900; Barber 1903; Hole 1917; Ayyar 1918; Hodgson 1918; Hole 1918; Lushington 1918; Jivana Rao 1920; Jivanna Rao 1921; Coleman 1923;

Hart, Rengaswamy 1926; Sreenivasaya 1930a; 1930b; Sreenivasaya, Rangaswami 1934; Sastri 1936).

Given the expanding nature of *Santalum* spp. and Indian sandalwood plantations throughout the Indian subcontinent and the Pacific (Teixeira da Silva et al. 2016), greater research into the causal agents, methods of transmission, molecular mechanisms underlying infectivity and methods to control or eradicate SSD would be beneficial. Research into SSD has stagnated in recent years, restricted to a very limited number of laboratories, primarily in India. This synthesis thus serves to add an impulse of interest among phytopathologists who might find this disease to be worthy of research.

Discovery of the pathogen

Research to investigate the causal organism for SSD spanned over more than half a century, from 1900 to 1968, but without any conclusive findings (Barber 1903; Brandis 1903; Hole 1917; Ayyar 1918; Fisher 1918; Hodgson 1918; Hole 1918; Lushington 1918; Rama Rao 1918; Jivana Rao 1920; Jivanna Rao 1921; Coleman 1923; Hart and Rengaswamy 1926; Sreenivasaya 1930a; Sreenivasaya 1930b; Sreenivasaya, Rangaswami 1934; Sastri 1936; Ramaiah et al. 1961; Parthasarathi, Rao 1962; Parthasarathi et al. 1962; Ramaiah et al. 1962; Parthasarathi et al. 1963; Parthasarathi et al. 1966), even though various news articles

were published in Nature® during the same period (Editorial News, 1933a, 1933b, 1933c, 1940, 1941). Barber (1903) observed the following, with relation to SSD: “Results of my studies has been to trace the disease to the roots, but the exact cause of this diseased condition has not been determined”, noting further that “It is difficult to see how effected trees could touch each other” referring to the steady progress of SSD from one plant to another as a communicable disease, irrespective of weather or soil conditions. Microscopic observations of *S. album* samples infected with SSD revealed a high accumulation of starch relative to healthy shoots, leading Barber to conclude that diseased plants die due to undigested nutrients (Barber 1903). He also observed that haustoria, which are specialized organs that assist in the hemiparasitic association of *S. album* with a host, were dead in the infected trees. In contrast, other researchers were of the opinion that SSD was not caused by infection, but was instead induced by an unbalanced circulation of sap (Hole 1917; Ayyar 1918) or a virus or viral-like microscopic organism (Coleman 1923; Dijkstra 1968). Dijkstra and Ie (1969) observed the presence of mycoplasma in the phloem sap of infected *S. album* trees, with Hull et al. (1969) and Verma et al. (1969) independently confirming the causal agent to be a mycoplasma. Mycoplasmas are now referred as phytoplasmas, i.e., prokaryotes lacking cell walls, infecting a wide range of plant taxa (Marcone 2015). Plant diseases caused by, or associated with, phytoplasmas occur in many agricultural and horticultural plants species of importance (Bertaccini et al. 2014; Arunkumar et al. 2016; Gurr et al. 2016). There are several reviews on various aspects of phytoplasma research (Marcone 2011; Kube et al.

2012; Marcone 2012; Bertaccini et al. 2014; Bertaccini 2015; Marcone 2015; Gurr et al. 2016; Trivedi et al. 2016).

Symptoms

SSD is characterised by chlorosis, a reduction in leaf size, and shortened internodes, causing leaves to become crowded on twigs with a “bushy” appearance and as stems stand out stiffly, they acquire a spike-like appearance (Barber 1903; Hole 1917; Fig. 1A, 1B). Spiked plants do not bear flowers or fruits and trees generally die within 1 to 2 years after the appearance of symptoms at a young age (Swaminathan et al. 1998; Balasundaran, Muralidharan 2004; Upul Subasinghe, personal communication). Due to SSD, *S. album* populations in the forests of Kerala and Karnataka in India became considerably depleted (Swaminathan et al. 1998; Kumar 2014; Arunkumar et al. 2016). Binu et al. (2014) also indicated that *S. album* production in India has been decreasing at an annual rate of 20% since 1995 for various reasons, including SSD, illicit felling and poor regeneration ability. Phytoplasmas are unculturable bacteria of the class *Mollicutes*; *Candidatus* (*Ca.*) that are associated with several symptoms, including “witches’ broom (clustering of branches) of developing tissues; phyllody (retrograde metamorphosis of the floral organs to the condition of leaves); virescence (green coloration of non-green flower parts); bolting (growth of elongated stalks); formation of buncy fibrous secondary roots; reddening of leaves and stems; generalized yellowing, decline and stunting of plants; and phloem necrosis” in many important food, vegetable, and fruit crops, ornamental plants, and timber and shade



Fig. 1. Control sandalwood (*Santalum album* L.) plant (A) vs plant infected with sandal spike disease (B). Both photos from Arun Kumar, A.N., Joshi, G., 2012. Incidence of sandal spike symptoms in a one-year-old plantation in Karnataka. *Curr. Sci.* 103: 613–614. With kind permission from Dr. Arun A. N. Kumar.

trees (Lee et al. 2000; Hogenhout et al. 2008; Bertaccini, Duduk 2009).

Detection

At the time when phytoplasmas were discovered, the diagnosis of plant pathogens was difficult since detection was based on the “observation of symptoms, insect or dodder/graft transmission to host plant and electron microscopy of ultra-thin sections of the phloem tissue” (Bertaccini, Duduk 2009; Delić 2012). Serological diagnostic techniques for the detection of phytoplasmas began to emerge in the 1980’s with direct and indirect enzyme-linked immunosorbent assay (ELISA)-based methods (Martin et al. 2000; Delić 2012). Methods based on ELISA and electron microscopy were reported by Balasundaran and Muralidharan (2004) and Thomas and Balasundaran (2001). ELISA could detect SSD six months before the expression of external symptoms (Balasundaran, Muralidharan 2004) but serological methods are insufficiently sensitive to detect various phytoplasmas (Delić 2012). PCR-based methods were used by Khan et al. (2004; 2006; 2008) and Murli et al. (2013). Khan et al. (2008) revealed, following restriction fragment length polymorphism (RFLP) analysis, and using nested PCR of 16S rDNA (Khan et al. 2004), that two SSD strains showed high sequence similarity to the aster yellows subgroup 16SrI-B. PCR-based detection of phytoplasmas has been reviewed by Delić (2012). PCR-based methods revealed that *Ca. Phytoplasma cynodontis* belongs to 16SrI, which is one of the major groups of phytoplasmas associated with different plant species reported to be infected with phytoplasmas worldwide (Lee et al. 1998; Hogenhout et al. 2008; Bertaccini, Duduk 2009; Laimer et al. 2009; Marcone 2011; Rao et al. 2011; Delić 2012; Kube et al. 2012; Gurr et al. 2016). An understanding of phytoplasma ecology and epidemiology has increased as a result of the use of these PCR-based methods (Martin et al. 2000). Phytoplasmas can be detected by real-time PCR, nested-PCR and microarrays, but most of these need 16S rRNA gene sequences (Hogenhout et al. 2008). About 40% of open reading frames in phytoplasmas are dissimilar to GenBank sequences, and about 200 genes are conserved in the OY and AY-WB genomes (Bai et al. 2006). These distinctive genes can be useful targets for the detection of phytoplasmas (Hogenhout et al. 2008). Analyses of 16S rRNA gene sequence analyses of about 200 phytoplasmas as well as members of *Ca. Phytoplasma asteris* (16SrI group phytoplasmas) showed that about 30 clades could form in 80 monocotyledonous and dicotyledonous plant species (Hogenhout et al. 2008).

Vectors and disease transmission

Coleman (1917) first reported that SSD can be transmitted through grafting, and later added that haustorial

connection of infected plants to a healthy host can be a possible means of SSD transmission (Coleman 1923). Grafting-based disease transmission was studied by histological observations (Dijkstra, van der Want 1970), in which cross sections of SSD-infected plants had phloem, the cambial area showed the presence of necrosed cells, while spongy parenchyma and palisade tissues remained unaffected. Insect transmission of SSD was suspected by Coleman (1917; 1923). Much later it was confirmed that SSD can be transmitted by insect vectors (Rangaswami, Sreenivasaya 1934). Insect vectors such as *Moonia albimaculata* (Dover, Appanna 1933; Sen-Sarma 1981; Sen-Sarma 1982), leafhopper *Jassus indicus* [Rangaswami, Griffith 1941; *J. indicus* is now known as *Coelidia indica* Walker (Marcone 2015)] and *Nephotettix virescens* Distant (Shivaramakrishnan, Sen-Sarma 1978; Sen-Sarma 1981; 1982). The transmission of phytoplasmas via insect vectors has been extensively reviewed (Bai et al. 2006; Weintraub, Beanland 2006; Hogenhout et al. 2008; Marcone 2012). According to Weintraub and Beanland (2006), phytoplasmas are restricted to the phloem, therefore only insects that feed on phloem sap are responsible for the transmission of phytoplasmas. The physiology, biochemistry and molecular biology of the mechanism of the insect-phytoplasma-plant interaction has been described in detail by Weintraub and Beanland (2006), Kube et al. (2012), Marcone (2012) and Trivedi et al. (2016). Recently, Konnerth et al. (2016) reviewed the immunodominant membrane proteins found on the outer membrane of phytoplasmas, which can help to understand the phytoplasma-host interaction and its biology, virulence and mode of action.

Control of SSD

Absolute characterization of the phytoplasma responsible for SSD has still not been achieved and research is still required to reveal its definitive source and the best form of disease management (Bertaccini 2015; Food and Agriculture Organization of the United Nations 2015). Nayar (1980) reviewed various methods such as the use of heat therapy and chemical treatment using arsenic, benlate, or tetracyclines to control SSD, the latter having the most positive effect on SSD control. However, the application of tetracyclines to a large forest area is impractical. Thus, the most effective methods to eliminate SSD from sandalwood tissue are likely to be the use of tissue culture to create disease-free superior genotypes (Balasundaran, Muralidharan 2004), or to employ genetic transformation to introduce sequences that, once overexpressed, will reduce the phytoplasma titre, or the development of insect-resistant plants. Phytoplasma-induced disease can be controlled by proper management of insect vectors (Weintraub, Beanland 2006; Gurr et al. 2016), removal of infected plants from the field (Gurr et al. 2016), and of the application of plant growth-promoting microbes that not

only enhance nutrient levels but also provide protection against microbial and insect vectors (Trivedi et al. 2016).

Conclusions

The mass production of *Santalum* spp., including Indian sandalwood, is now taking place not only on the Indian sub-continent, but also on mainland China and in Australia, for the production of high quality heartwood for use in the wood and perfume industries. The greatest limitation to field establishment lies with the selection of a suitable host as a result of the plant's hemiparasitic nature. SSD does not appear to pose a significant threat (yet) to Chinese and Australian cultures, but expanded research into understanding the causal agent and a molecular method to better control of the phytopathogen would be desirable.

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