

Arbuscular Mycorrhizal Symbiosis in *Trema orientalis*: Effect of a Naturally-Occurring Symbiosis Receptor Kinase Mutant Allele

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Abstract : The *Trema* genus represents a group of fast-growing tropical tree species within the Cannabaceae. Interestingly, five species nested in this lineage -known as *Parasponia*- can establish rhizobium nitrogen-fixing root nodules, similar to those found in legumes. *Parasponia* and legumes use a conserved genetic network to control root nodule formation, among which are genes also essential for mycorrhizal symbiosis (the so-called common symbiotic pathway). However, *Trema* species lost several genes that function exclusively in nodulation, suggesting a loss-of the nodulation trait in *Trema*. Strikingly, in a *Trema orientalis* population found in Malaysian Borneo we identified a truncated SYMBIOSIS RECEPTOR KINASE (SYMRK) mutant allele lacking a large portion of the c-terminal kinase domain. In legumes this gene is essential for nodulation and mycorrhization. This raises the question whether *Trema orientalis* can still be mycorrhized. To answer this question, we established quantitative mycorrhization assay for *Parasponia andersonii* and *Trema orientalis*. Plants were grown in closed pots on half strength Hoagland medium containing 20 μ M potassium phosphate in sterilized sand and inoculated with 125 spores of *Rhizopagus irregularis* (Agronutrition-DAOM197198). Mycorrhization efficiency was determined by analyzing the frequency of mycorrhiza (%F), the intensity of the mycorrhizal colonization (%M) and the arbuscule abundance (%A) in the root system. *Trema orientalis* RG33 can be mycorrhized, though with lower efficiency compared to *Parasponia andersonii*. From this we conclude that a functional SYMRK kinase domain is not essential for *Trema orientalis* mycorrhization. In ongoing experiments, we aim to investigate the role of SYMRK in *Parasponia andersonii* mycorrhization and nodulation. For this two *Parasponia andersonii* symrk CRISPR-Cas9 mutant alleles were created. One mimicking the TorSYMRKRG33 allele by deletion of exon 13-15, and a full *Parasponia andersonii* SYMRK knockout.

Keywords : endomycorrhization, *Parasponia andersonii*, symbiosis receptor kinase (SYMRK), *Trema orientalis*

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