Project 3.5 – Project Executive Summary.

Phylogenetic position of Puccinia psidii within the Pucciniales

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Project Executive Summary.

The phylogenetic position of *Puccinia psidii* (guava rust, myrtle rust or Eucalyptus rust) within the Pucciniales, in particular its relationship with Pucciniaceae, was investigated. The study adopted a multi-step approach, which firstly included an extensive number of taxa covering 12 of the 13 currently known rust families and constructed a phylogeny based on sequences from the nuclear large subunit ribosomal gene (LSU). This phylogeny was then used to identify the phylogenetic affinities of P. psidii within the rust fungi. The nuclear small subunit ribosomal gene (SSU) was then obtained for these closely related taxa and, together with representatives from the Pucciniaceae and representative families, a combined two-locus phylogeny was derived. Finally partial coding regions of the β -tubulin 1 gene (β -tub1) were sequenced for a smaller sample of taxa to corroborate or contradict the LSU/SSU phylogenies. The results obtained clearly show that *P. psidii* does not belong in the Pucciniaceae. Its closer relatives include Dasyspora, Sphearophragmium, Diorchidium, as well as *Uredo baruensis* and *P. popowiae*. Since none of these genera are suitable systematic home for P. psidii, it is recommended that a new genus be erected to indicate that P. psidii is unambiguously phylogenetically distinct from representatives of *Puccinia s.s.*

The Final Report for this project is not available as manuscripts describing this work are in preparation. For further information, please contact Lead Researcher Dr Edward C.Y. Liew Edward.Liew@rbgsyd.nsw.gov.au

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