

Genetic differentiation between sympatric sister species of eucalypts.

Rachael M. Fowler (School of Biosciences, The University of Melbourne) Harvey K. Orel (School of Biosciences, The University of Melbourne) David Cantrill (Royal Botanic Gardens Victoria) Daniel J. Murphy (Royal Botanic Gardens Victoria) Patrick Fahey (Research Centre for Ecosystem Resilience, The Royal Botanic Garden Sydney) Mike J. Bayly (School of Biosciences, The University of Melbourne) Todd G.B. McLay (School of Biosciences, The University of Melbourne; Botanic Gardens Victoria; National Biodiversity DNA Library, CSIRO).

Extensive hybridisation among related species is well-documented in the eucalypts, which raises questions about how speciation occurs and high species diversity is maintained. We explored genetic diversity and differentiation between two sister species of eudesmid eucalypts, Eucalyptus tetrodonta and Eucalyptus megasepala. The two species have overlapping distributions in northern Queensland, where E. megasepala is restricted to the Cape Yorke Peninsula, including areas where both species can occur within metres of one another; E. tetrodonta is widespread from northern Western Australia, through the Northern Territory and into Queensland. Despite their overlapping geographic ranges and apparently limited physical and environmental barriers to gene flow, significant genetic differentiation and population structure were observed with limited evidence of gene flow between the two species. Outlier loci associated with divergent selection were identified, suggesting genetic adaptations underpinning ecological differences. Additionally, flowering time analysis revealed temporal isolation, potentially contributing to reproductive barriers. Over its entire range, E. tetrodonta genetic diversity follows an isolation by distance pattern with a modestly supported genetic split occurring near the Gulf of Carpentaria. These findings provide valuable insights into the intricate mechanisms driving speciation in Eucalyptus species and have broader implications for understanding biodiversity in temperate tree genera.

Todd McLay: todd.mclay@csiro.au