Plastid phylogenomics reveals evolutionary relationships in the mycoheterotrophic orchid genus *Dipodium* and provides insights into plastid gene degeneration

















Heterotrophic plants





-Epipogium roseum-



-Gastrodia umbrosa-

& Low 201 Gray B. Gray; Photo:



-Rhizanthella gadneri-

(Myco)-heterotrophy





Orchidaceae – Life cycle – nutrition strategies





Evolution of mycoheterotrophy



Reduction in leaf size

Loss of photosynthetic activity

Loss of chlorophyll

Degradation of the plastid genome



-Gastrodia umbrosa-



-Rhizanthella gadneri-



Delannoy et al. 201

hoto: D.L. Jon

-Epipogium roseum-

Plastid genome (plastome) of mycoheterotrophs

Rampant Gene Loss in the Underground Orchid Rhizanthella gardneri Highlights Evolutionary Constraints on Plastid Genomes

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Associate editor: Charles Delwiche

plant plastomes:

152,858 kb







Plastid genes - degradation and loss



Functional gene



Functional loss: pseudogenised gene



Physical loss: non-detectable gene

Plastid genes – gene groups





Plastid genes - degradation and loss





Dipodium R.Br.





-Dipodium pandanum-

-Dipodium ensifolium-





-Dipodium elegantulum-

Dipodium R.Br.



Common name: Hyacinth orchids

Subfamily: Epidendroideae Tribe: Cymbidieae

• 39 species

Divided in two section:

- Section Dipodium
- Section Leopardanthus (Blume) O. Kuntze



https://commons.wikimedia.org/wiki/File:BlankMap-World.svg

Section Leopardanthus in Australia

- 'Leafy', autotrophic species
- Adventitious roots, epiphytes and terrestrials



Section Dipodium in Australia

- 'Leafy', autotrophic species
- Non-adventitious roots, terrestrials



Section Dipodium in Australia

- 'Leafless', mycoheterotrophic species
- Non-adventitious roots, terrestrials



Previous studies



CORRIGENDUM

https://doi.org/10.1071/BT22075_CO



Australian Journal of Botany

Corrigendum to: Retention of an apparently functional plastome in an apparently mycoheterotrophic orchid, Dipodium roseum D.L.Jones & M.A.Clem. (Orchidaceae)

Todd G. B. McLay, Michael J. Bayly, Michael R. Whitehead and Rachael M. Fowler



ORIGINAL RESEARCH published: 21 February 2020 doi: 10.3389/fpls.2020.00022



Plastome Evolution and Phylogeny of Orchidaceae, With 24 New Sequences

Young-Kee Kim¹, Sangjin Jo¹, Se-Hwan Cheon¹, Min-Jung Joo¹, Ja-Ram Hong¹, Myounghai Kwak² and Ki-Joong Kim^{1*}

¹ Division of Life Sciences, Korea University, Seoul, South Korea, ² Department of Plant Resources, National Institute of Biological Resources, Incheon, South Korea

Phylogenomic resolution: Orchidaceae

Maximum likelihood phylogenetic tree: Based on 68 plastid loci and 148 taxa



• SH-aLRT/ufboot < 80/95











Divergence-time estimations: Orchidaceae I Epidendroideae I Cymbidieae

Bayesian analyses performed with BEAST 2: based on 30 plastid loci and 134 taxa; Best-fit model for the data set: Optimised relaxed clock/ birth-death (Model comparison by AICM; Fabozzi *et al.* 2014)



Orchid chronogram



Divergence-time estimations: Orchidaceae I Epidendroideae I Cymbidieae

Scotese, C.R., 2014

Bayesian analyses performed with BEAST 2: based on 30 plastid loci and 134 taxa; Best-fit model for the data set: Optimised relaxed clock/ birth-death (Model comparison by AICM; Fabozzi *et al.* 2014)



Early Oligocene



Bayesian analyses performed with BEAST 2: based on 30 plastid loci and 134 taxa; Best-fit model for the data set: Optimised relaxed clock/ birth-death (Model comparison by AICM; Fabozzi *et al.* 2014)



Middle to late Miocene



Australia has arrived at today's geographical position

Bayesian analyses performed with BEAST 2: based on 30 plastid loci and 134 taxa; Best-fit model for the data set: Optimised relaxed clock/ birth-death (Model comparison by AICM; Fabozzi *et al.* 2014)



Late Miocene



95% highest posterior density (HDP) values



Australia's climatic conditions became increasingly arid

Bayesian analyses performed with BEAST 2: based on 30 plastid loci and 134 taxa; Best-fit model for the data set: Optimised relaxed clock/ birth-death (Model comparison by AICM; Fabozzi *et al.* 2014)



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climatic oscillations

eopardanthus.

Dipodium

Dipodium plastome: general features

Plastome assembly

• 24 Dipodium plastomes were assembled using Geneious Prime and a de novo & reference-guided assembly approach (reference plastomes D. roseum (MN200386) & M. coccinea (KP205432)).



Dipodium plastome: general features

Plastome assembly

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Dipodium plastome: structural hotspots – *ndh* genes pseudogenisation and loss

Plastome assembly

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L**L**.,

tmV-UAC* ndhCΨ ndhKΨ ndhJΨ

5

ndh gene degradation: Dipodium plastomes

Maximum likelihood phylogenetic tree: Based on 68 concatenated plastid loci (69,335 bp) and 145 taxa; Best-fit model: GTR+I+I+F+R4 (IQ-TREE)



ndh gene degradation: Dipodium plastomes

Maximum likelihood phylogenetic tree: Based on 68 concatenated plastid loci (69,335 bp) and 145 taxa; Best-fit model: GTR+I+I+F+R4 (IQ-TREE)



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Plastid genes - degradation and loss





Plastid genes - degradation and loss







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