

PhyloJIVE (and others): Integrating biodiversity data with the Tree of Life

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Centre for Australian National Biodiversity Research
National Science Foundation

Collaborators

Funding: DEWR, CANBR and ALA

- * CANBR

- * Garry Jolley-Rogers

- * ALA

- * Temi Varghese

- * Rebecca Pirzl

- * Nic dos Remedios

- * iDigBio

- * Andréa Matsunaga

- * others

- * Andrew Thornhill-ATH

- * Darren Crayn-ATH

- * John Pickering –
DiscoverLife

- * Dan Rosauer-ANU

Our Data

- * Biological collection informatics community
 - * Individual datasets
 - * iDigBio, GBIF, EOL, ALA, Map of Life
 - * Dryad, Morphbank, Morphobank, Traitbase, IDLife
- * Evolution and phylogenetics community
 - * Individual datasets
 - * Open Tree of Life
 - * Treebase

Our Data

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 - * Open Tree of Life

Data is not integrated.

Why Integrate Data?

The image features a blue header with a wavy, layered design. The top part is a solid blue bar containing the text 'Why Integrate Data?'. Below this bar, the background transitions into a white space with several overlapping, semi-transparent blue wavy lines that create a sense of depth and movement.

Integrated Data?

Currently done for individual studies

- * Currently done for individual studies
 - * Manual data generation and curation
- * Goal: Community resource
 - * Tools to link data

Why Integrate Data?

- * To understand our data
- * Study biodiversity at all levels, not just species
- * Facilitate comparative biology research
 - * Visualize patterns
 - * Develop and test hypotheses
- * Conservation planning
- * Evidence-based decision making

- 
- * “the ability to search multiple databases using the nodes of a phylogenetic tree may be the single most important contribution of systematics to conservation and sustainable use of biodiversity”

Cracraft (2002)

Outline

- * PhyloJIVE
- * Numerous screencasts
- * Other integrating and visualization tools

PhyloJIVE

Phylogeny Javascript Information Visualiser and Explorer

A

B

- External Links
- View Map
- Expand/Collapse
- Rotate
- Set Root

C

D *Acacia camptoclada*

Growth form: erect shrub

Inflorescence shape: globular

Inflorescence_arrangement: racemose

plant_height: 0.6000

Phyllode_length__median: 12.0000

E Legend

Growth form

- erect shrub
- prostrate shrub
- tree
- multiple

Inflorescence shape

- globular
- obloid to ellipsoid

F Taxa

ALA API EOL APNI [Acacia prainii](#)

ALA API EOL APNI [Acacia camptoclada](#)

ALA API EOL APNI [Acacia anthochaera](#)

ALA API EOL APNI [Acacia suaveolens](#)

ALA API EOL APNI [Acacia iteaphylla](#)

G ATLAS of LIVING AUSTRALIA
sharing biodiversity knowledge

Species Locations Collections Mapping & analysis

***Acacia murrayana* F.Muell. ex Benth.**

Colony Wattle

Overview Gallery Names Classification Records Literature Sequences

H

***Acacia murrayana* F.Muell. ex Benth. [apni \(as xml\)](#)**

URI <http://biodiversity.org.au/apni.name/12669>

LSID <urn:lsid:biodiversity.org.au:apni.name:12669>

source http://www.anbg.gov.au/cgi-bin/apni?taxon_id=12669

modified 2013-01-08

Title *Acacia murrayana* F.Muell. ex Benth.

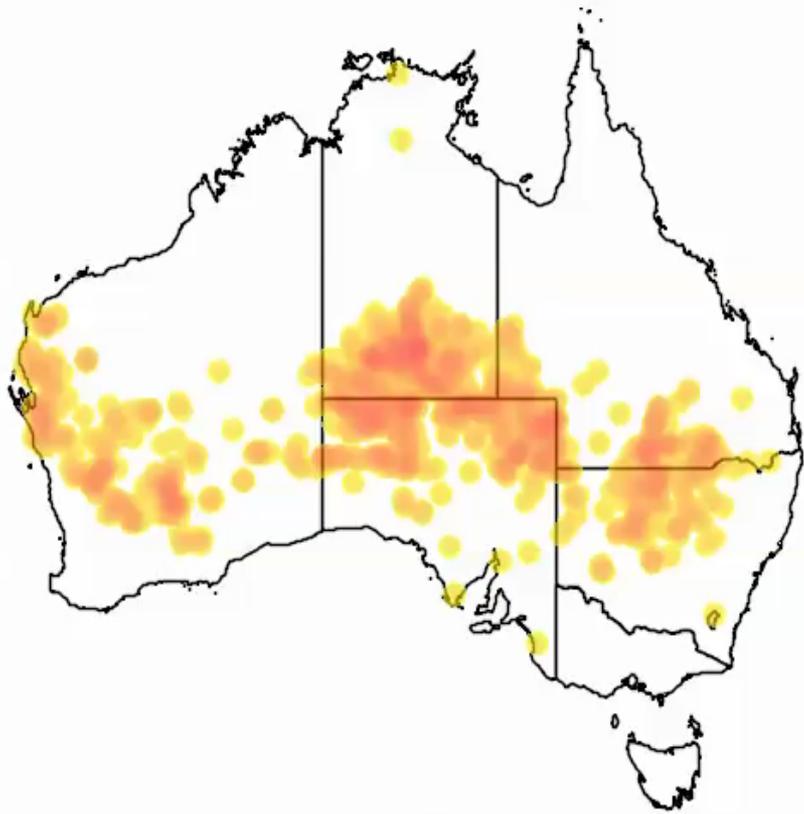
I Australian Plant Image Index

[Acacia triquetra](#)

Taken at. ANBG

Photographer. Fagg, M.

dig 22097





Create a new tree

Tree for Case Study 1: Acacia

Admin actions: [Tree list](#) [Edit tree](#)

Options Character Legend
Map Search About

Actions

Tree Options

Align Names

Ladderize

Animate

Branch Length

Length Multiplier x0.1 x1 x2 x5

Load characters via

Identify Life

Add to Map ▾ Tools ▾ Import ▾ Export ▾ Help ▾

☑ dunii

Map options

dunii

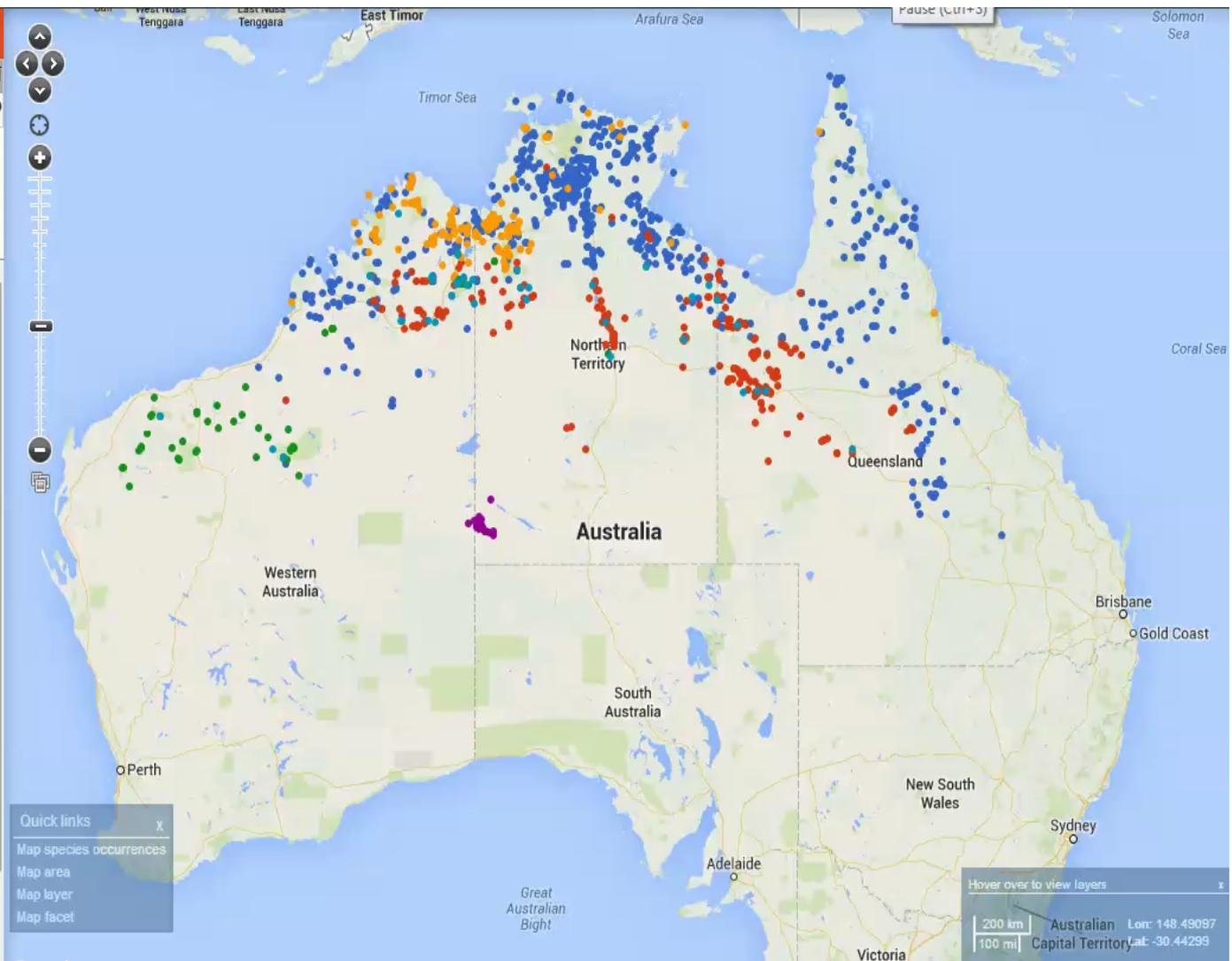
Records selected: 0

Highlight occurrences on the scatterplot that are in an area
 Australia

Selected species

Temperature - annual mean (Bio01)

Precipitation - annual (Bio12)



Home → Phylojive

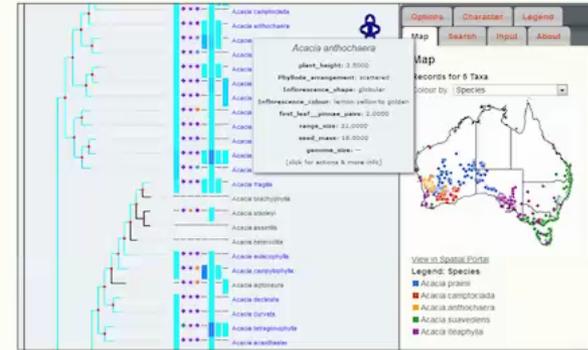
Phylojive

PhyloJive ([Phy](#)logeny [J](#)avascript [I](#)nformation [V](#)isualiser and [E](#)xplorer) is a web based application that places biodiversity information aggregated from many sources onto compact phylogenetic trees.

The project is the brainchild of [Garry Jolley-Rogers](#) and [Joe Miller](#) and was developed by Temi Varghese and [Garry Jolley-Rogers](#) as part of the [Taxonomy Research & Information Network \(TRIN\)](#) – see the [original project page](#), [original code repository](#) and [ALA code repository](#). The ALA has contributed to the PhyloJive codebase to integrate a number of web services: occurrence data, maps and character data from Identify Life. This work has been undertaken with help and advice from [Joe Miller](#).

The [getting started](#) page outlines the steps for creating a new phylogenetic tree and contains demo data sets that can be used to get up and running.

Create a new tree



Trees with character data

- quantitative characters
- demo
- ARFK1
- joe test July 26
- Boletes
- Damien
- Joe's acacia
- kimberley land snails3
- opentree
- Asteraceae coll bias
- treetest
- Tests

"Bare" Trees

- Mollusca-tree of life
- amphibia - frogs, salamanders and caecilians
- Damien
- Maluridae
- Bird super tree
- kimberley land snails3
- 4000 lizards
- test3
- Characteristics 1-8
- dfd
- Tests
- opentree



iDigBio Implementation

← → ↻ phylojive.acis.ufl.edu/PhyloJive/index.jsp

iDigBio & ALA PhyloJIVE
Integrated Digitized Biocollections

About iDigBio | Portal | Technical Information | Education

iDigBio Home | iDigBio Portal | **PhyloJIVE Home** | OpenTree Studies | Sample Trees | Tutorial | Research Tools | Feedback

PhyloJIVE
integrating phylogenies
... with online biodiversity data sources

for developers, [wiki instructions + known problems](#), [download PhyloJIVE](#)

[View on GitHub](#)

tar.gz .zip

The growing catalogue of online biodiversity data sources has reached a scale where data integration and mashups can be truly useful. Information about species is being aggregated at national and international scales. [PhyloJIVE](#) ([Phylogeny Javascript Information Visualiser and Explorer](#)) is one such mashup placing biodiversity web-services into an evolutionary context.

Integration with [Atlas of Living Australia](#) and [iDigBio](#)

Joe Miller's has made [detailed](#) instructions on the use of the ALA implementation of phyloJIVE including videos, which are similar to the [instructions for the iDigBio instance of PhyloJIVE](#). Briefly, this instance offers the following additional features:

- linkage to specimens available on iDigBio
- visualization of iDigBio specimen on a world map
- external links to ALA Taxonomy, EOL, Discover Life, ALA maps
- access to a list of phylogenetic studies made available by OpenTree

Example trees on iDigBio+PhyloJive:

Pause (Ctrl+S)

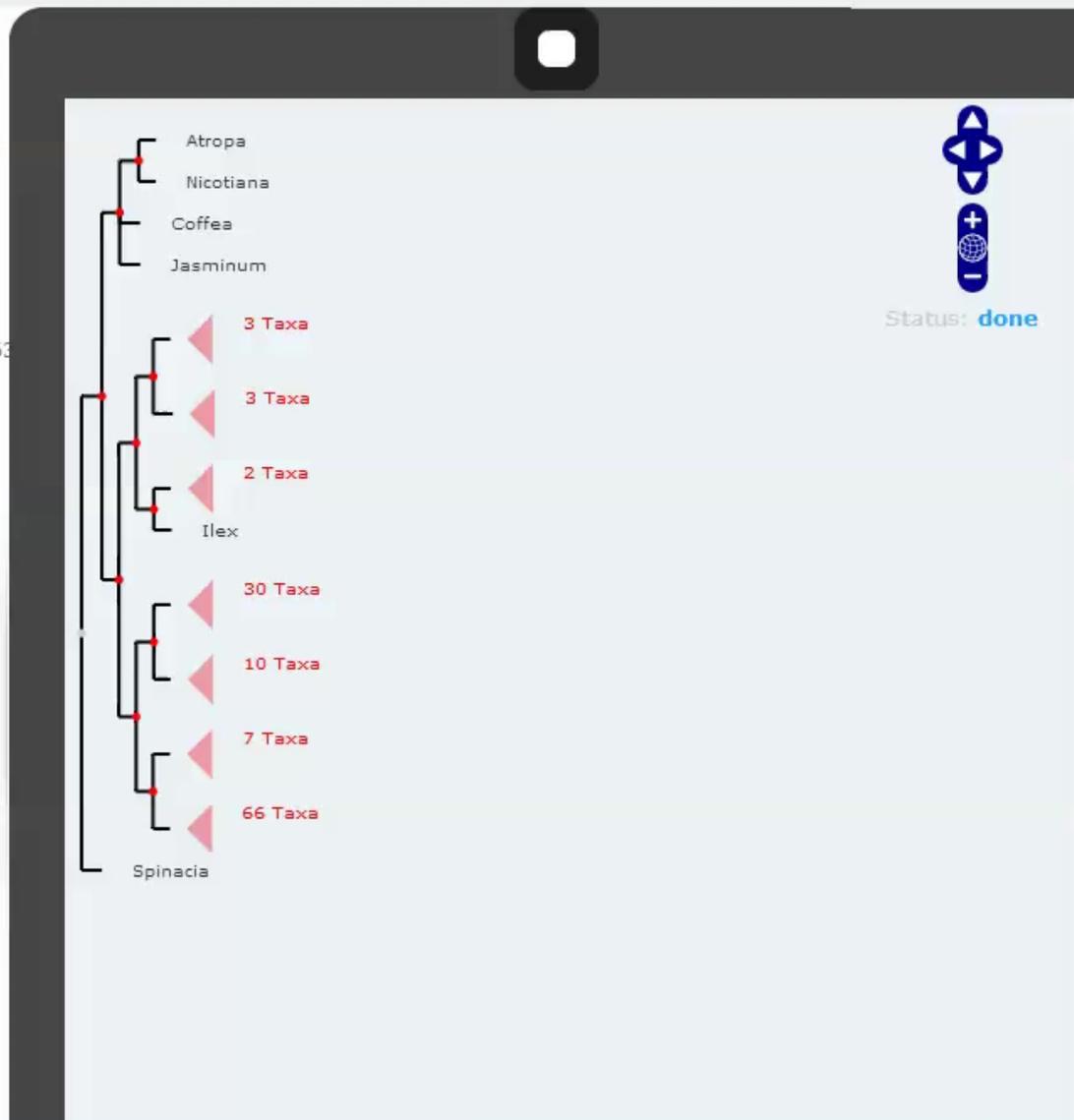
Andréa Matsunaga, iDigBio, UF

[OpenTree Study: pg_09](#)

Tank, David C., and Michael J. Donoghue. 2010. Phylogeny and Phylogenetic Nomenclature of the Campanulidae based on an Expanded Sample of Genes and Taxa. *Systematic Botany* 35, no. 2 (6): 425-441.
doi:10.1600/03636441079163

Select another OT study:

- Click the top button to get the navigation aid
- Click nodes to get maps and external services
- Try choosing characters (if available) to plot on the tree;
- Align-names feature; search; set-root; rotate, etc.



OneZoom, James Rosindell

<http://www.onezoom.org/>

Linking phylogenetic widget into you webpage



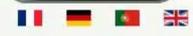
The image shows a screenshot of the OneZoom web application. At the top, there is a black header bar with the OneZoom logo on the left and four navigation icons (plus, minus, refresh, and zoom) on the right. Below the header, the main content area has a light blue background. In the center, a large, stylized phylogenetic tree is displayed. The tree is primarily green, with some branches in brown and red. The word "Tetrapods" is written in a large, black, serif font across the middle of the tree. At the bottom of the screenshot, there is a dark grey bar with the text "Click to see animation" in white.



Help save the Great Apes & their habitats



Gorilla
Africa



- Home
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- Events
- Working Groups
- Get Involved
- Ape Inspired
- The Apes
- About Us
- Shop



Mt. Kilimanjaro

Climb for Change

Sponsor Ian, Support the Ape Alliance



Gorilla

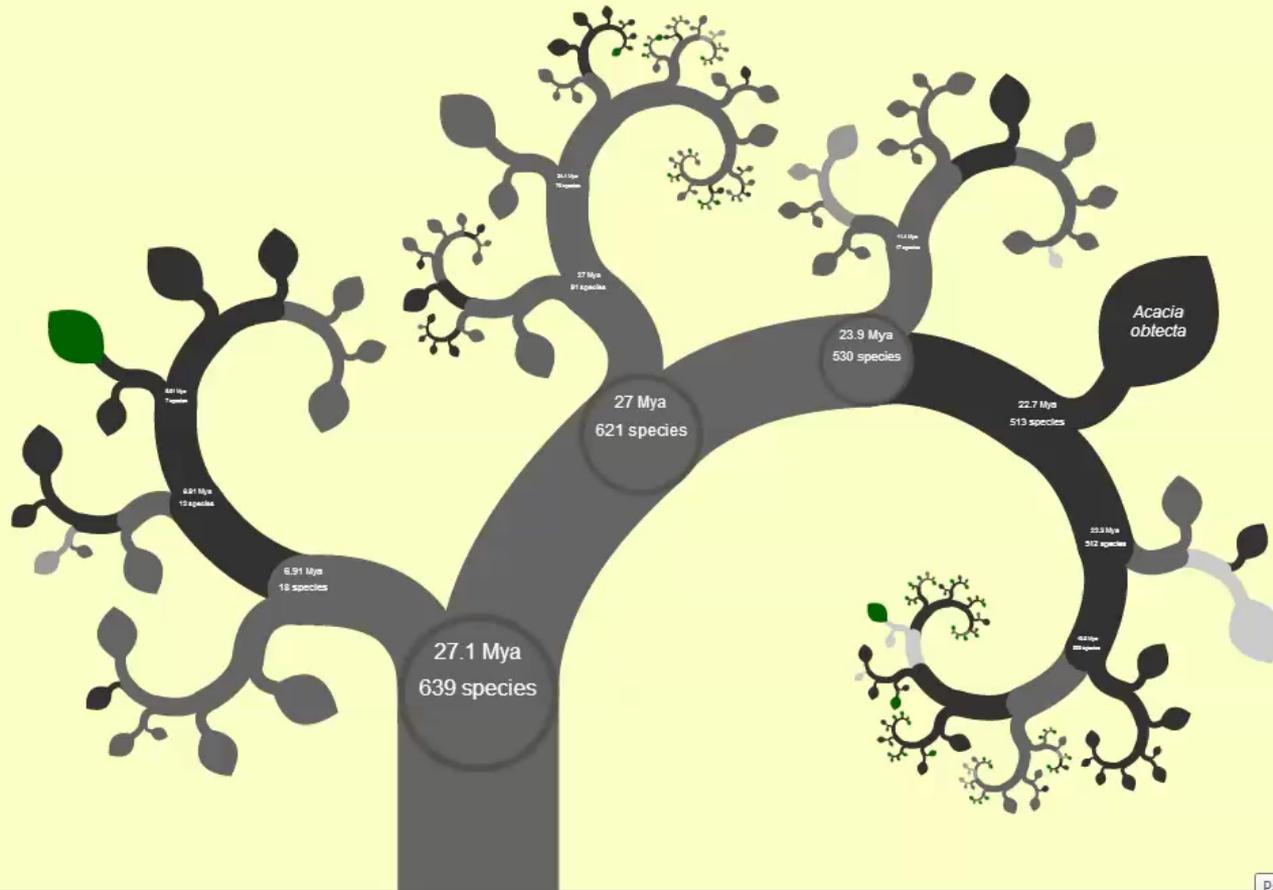
© Ian Redmond

[Back to Top](#)

OneZoom with PhyloJIVE



Inflorescence Arrangement ▾



Pause (Ctrl+S)

Growth animation Reverse Pause Play Stop Close Present day

Gavin Naylor, shakrays.org

shakrays.org

Chondrichthyes: Tree of Life

Phylogeny Atlas DNA Anatomy

Contact: Gavin Naylor

The image displays a phylogenetic tree of Chondrichthyes (sharks and rays). On the left, a vertical list of species names is provided, including *Callorhynchus callorhynchus*, *Rhinodimaera pacifica*, *Chimaera argiloba*, *Hydrolagus marmoratus*, and *Spharxiparus*. On the right, a circular tree diagram shows the relationships between major groups, with representative images of organisms from each group. The groups shown include *Rhinopristiiformes*, *Torpediniformes*, *Rajiformes*, *Carcharhiniformes*, *Lamniformes*, *Orectocheiliformes*, *Chimaeriformes*, *Heterodontiformes*, *Squaliformes*, *Pristiophoriformes*, *Echinorhiniformes*, and *Squatriniformes*. At the bottom, there are interactive buttons for 'Look', 'Delete', 'Expand', and 'Collapse'.

Future

- * All data integrated
 - * GoLife, Genealogy of Life
- * Other integration platforms
 - * Arbor
 - * ???



* “Physicists acknowledge the power of the fundamental forces, such as gravity, incorporating them into every calculation. To successfully conserve our dwindling global biodiversity which we rely upon economically and aesthetically for sustenance and pleasure, biology must integrate evolution into every assessment of biodiversity.”

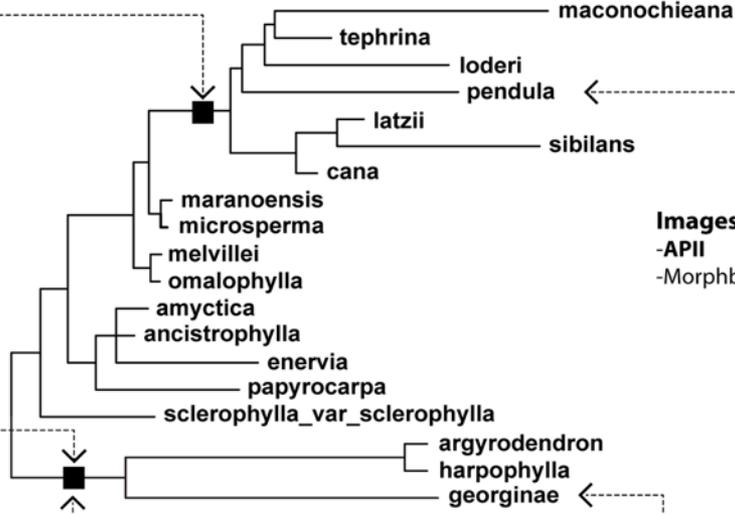
extra

| |
|---|
| Acacia latzii Maslin <i>sp. nov.</i> (A. 1007) (A. 1008) |
| LDI: http://www.identifylife.org/australia/nomenclature |
| LSID: http://dx.doi.org/10.31212/australia.nomenclature.1007 |
| Source: http://www.identifylife.org/australia/nomenclature |
| modified: 2012-10-24 |
| Title: Acacia latzii Maslin |
| Complex: Acacia latzii |
| Authority: Maslin |
| Contributor Authority: Maslin |
| Year: 2007 |
| Accepted: Acacia latzii Maslin <i>nom. nov.</i> http://www.identifylife.org/australia/nomenclature |
| Publication: Maslin, B. R. (2007) Acacia (Leguminosae-Mimosoideae): A contribution to the flora of central Australia. <i>Journal of the Adelaide Botanic Garden</i> 2(5): 313. fig. 1, 8. (image) |
| Base name: Acacia |
| Genus aut: Acacia |
| Specific ep: latzii |
| Cont. authority: Maslin |
| Type locality: "Shalwater Range, New Cavena Station, 25-08E(149-MEN)-6, 134-08E(126-MEN)-6, Northern Territory ... 21 April 1977. P.K. Lee 6999 (holo NY; see AD, CASL, K, NY) (1 tree seen with the AD and NY specimens)"; |
| Taxonomic events: Acacia latzii Maslin <i>nom. nov.</i> http://www.identifylife.org/australia/nomenclature |
| Acacia latzii Maslin <i>nom. nov.</i> http://www.identifylife.org/australia/nomenclature |
| Acacia latzii Maslin <i>nom. nov.</i> http://www.identifylife.org/australia/nomenclature |
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| Acacia latzii Maslin <i>nom. nov.</i> http://www.identifylife.org/australia/nomenclature |
| Acacia latzii Maslin <i>nom. nov.</i> http://www.identifylife.org/australia/nomenclature |

NOMENCATURE
 - APNI
 - IPNI

TAXONOMIC PAGES
 - EOL
 - ALA
 - GBIF
 - Wikipedia
 - DiscoverLife
 - Google

node support
 divergence times



ANCESTRAL CHARACTER STATES FOR CLADE
 - Foliage=phyllodes
 - inflorescence type=globular
 - Growth form= tree or shrub

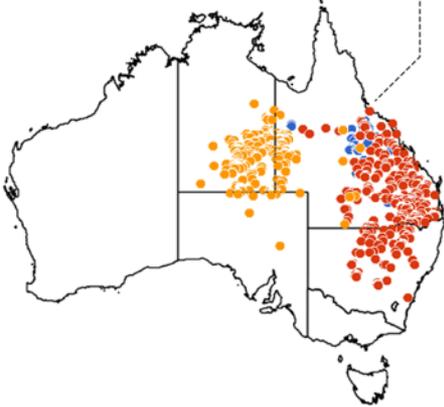
Images
 -APII
 -Morphbank



Australian Plant Image Index (APII)-
 Photo No. : a.19340

CHARACTER STATES FROM IDENTIFY LIFE
 - Foliage=phyllodes
 - inflorescence type-globular
 - Growth form= tree

TRAITS
 - user defined
 - ID keys Identify Life
 - webservices
 - text mining
 - floras



DISTRIBUTION MAPS (ALA)
 - environmental layers (climate and geography)
 - Google earth
 - analytical tools (endemism)

Generate a new tree



Create Phylogenetic Tree

Name *

Owner *

Is Public



Newick *

Characters

Input characters in [charJSON format](#). Alternatively, you can paste [CSV data](#) and then click:

Create Phylogenetic Tree

Name *

Owner *

Is Public



Tree (nexml or newick)



Search Treebase

Search

Search for a study in treebase

Treebase Tree ID

eg: Tr2026