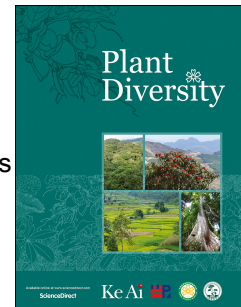


# Accepted Manuscript

Using a mega-phylogeny of seed plants to test for non-random patterns of areal-types across the Chinese tree of life

Rong Li



PII: S2468-2659(16)30080-4

DOI: [10.1016/j.pld.2016.08.003](https://doi.org/10.1016/j.pld.2016.08.003)

Reference: PLD 28

To appear in: *Plant Diversity*

Received Date: 16 August 2016

Accepted Date: 16 August 2016

Please cite this article as: Li, R., Using a mega-phylogeny of seed plants to test for non-random patterns of areal-types across the Chinese tree of life, *Plant Diversity* (2016), doi: 10.1016/j.pld.2016.08.003.

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

**Using a mega-phylogeny of seed plants to test for non-random  
patterns of areal-types across the Chinese tree of life**

Rong Li

*Key Laboratory for Plant Diversity and Biogeography of East Asia, Kunming Institute  
of Botany, Chinese Academy of Sciences, Kunming, 650201, China*

E-mail address: [lirong@mail.kib.ac.cn](mailto:lirong@mail.kib.ac.cn)

Download English Version:

<https://daneshyari.com/en/article/5767089>

Download Persian Version:

<https://daneshyari.com/article/5767089>

[Daneshyari.com](https://daneshyari.com)