

Subject: CRAN submission P2C2M 0.7.6
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To: CRAN <cran@r-project.org>

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The following package was uploaded to CRAN:

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Package Information:

Package: P2C2M

Version: 0.7.6

Title: Posterior Predictive Checks of Coalescent Models

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Maintainer: Michael Gruenstaeudl <mi.gruenstaeudl@gmail.com>

Depends: R (>= 3.0.0)

Suggests: genealogicalSorting (>= 0.92), phybase (>= 1.3.1), Rmpi (>= 0.6-5), xtermStyle (>= 2.2-4)

Description: Conducts posterior predictive checks of coalescent models using gene and species trees generated by 'BEAST' or '*BEAST'. The functionality of P2C2M can be extended via two third-party R packages that are available from the author websites: 'genealogicalSorting' and 'phybase'. To use these optional packages, the installation of the Python libraries 'NumPy' (>= 1.9.0) and 'DendroPy' (= 3.12.0) is required.

License: GPL (>= 2)

Imports: ape (>= 3.1-4), ggplot2 (>= 1.0.0), rPython (>= 0.0-5), stringr (>= 0.6.2)

The maintainer confirms that he or she
has read and agrees to the CRAN policies.

Submitter's comment: Update to version 0.7.6

Checking performed via "R

CMD check --as-cran P2C2M_0.7.6.tar.gz"; no errors,
warnings or relevant notes encountered