Bioinformatics
doi.10.1093/bioinformatics/xxxxxx
Advance Access Publication Date: Day Month Year
Manuscript Category



## **Subject Section**

# mulTree: using phylogenetic uncertainty in comparative methods

## Thomas Guillerme 1,\*, Natalie Cooper 2 and Kevin Healy 3,

- <sup>1</sup> Silwood Park Campus, Department of Life Sciences, Imperial College London, Buckhurst Road, Ascot SL5 7PY, United Kingdom and
- <sup>2</sup> Department of Life Sciences, Natural History Museum, Cromwell Road, London, SW7 5BD, United Kingdom and
- <sup>3</sup> Department of Zoology, School of Natural Sciences, University of Dublin, Trinity College, Dublin, Ireland, Country.

Associate Editor: XXXXXXX

Received on XXXXX; revised on XXXXX; accepted on XXXXX

#### **Abstract**

Motivation:

Results:

Contact: t.guillerme@imperial.ac.uk

Supplementary information: Supplementary data are available at *Bioinformatics* online.

- 1 Introduction
- 2 Approach
- 3 Methods
- 4 Discussion
- 5 Conclusion
- **Acknowledgements**

### **Funding**

This work has been supported by the... Text Text Text Text.

<sup>\*</sup>To whom correspondence should be addressed.