

Special Issue

Open Science Phyloinformatics: Resources, Methods, and Analyses

Message from the Guest Editors

Phylogenetic research, today, is increasingly complex and data-intensive, placing increased demands on effective data sharing, analytical reproducibility, and knowledge synthesis. Large trees, vast datasets, and long multi-step analysis pipelines demand that biologists follow best practices in open access, sharing, and data integration. This Special Issue aims to assemble articles on phyloinformatic resources, tools, and research reports that exemplify best practices supporting open science reproducibility and/or demonstrate inference of broad-scale patterns through the integration of large datasets in phylogenetics.

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Life (ISSN 2075-1729) is an international, peer-reviewed open access journal that publishes scientific studies related to fundamental themes in life sciences. Some papers are published individually, while others are submitted for inclusion in special issues with guest editors. You are invited to contribute a research article, essay, or a review to be considered for publication.

Editor-in-Chief

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