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# **Methods in Computational Biology**

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## **Message from the Guest Editors**

The focus of this special issue is methods in computational biology which can extract and test theory using large datasets. A primary goal of this issue is the communication of computational biology methods with enough detail to permit reproduction of results. Topics of interest span the gamut of fundamental and applied biology and can analyses of medical. environmental bioprocess systems. Modeling approaches of interest include, but are not limited to, flux balance analysis, elementary flux mode analysis, agent-based modeling as well as other dynamic modeling approaches. Papers combining experimental and computational studies are highly encouraged as are manuscripts that propose standards for model writing, storage and distribution. This issue aims to integrate highly interdisciplinary researchers such as biologists, computer scientists, engineers and mathematicians who focus on advances in biological systems analysis.

The issue is coordinated with the Metabolic Pathway Analysis 2017 conference held in Bozeman MT and select IMAG MultiScale Modeling (MSM) working groups.







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### **Editor-in-Chief**



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## **Aims and Scope**

Processes (ISSN 2227-9717) is an open access journal that provides an advanced forum for studies concerning process related research in chemistry, biology, and allied engineering fields. Our goals are to publish high impact articles of broad interest to the process systems community and to serve as a forum for major developments in process/systems research. The journal publishes regular research papers, communications, letters, short notes, and reviews. There are no restrictions on the length of published articles or on the use of color illustrations. All submitted manuscripts undergo rigorous peer review prior to publication.

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