# Green Journals

International Approved Journals

## Journal Information

| | |
|---|---|
| **Journal Title** | BMC Systems Biology |
| **Publisher** | BioMed Central |
| **ISSN** | 1752-0509 |
| **E-ISSN** | |
| **Database** | SCOPUS |
| **Domain** | Life Sciences, Physical Sciences |

## Classification History

| Year | A+ | A | B | C |
|---|---|---|---|---|
| **2026** | | | | |
| **2025** | | | | |
| **2024** | | | | |
| **2023** | | | | |
| **2021/2022** | | | ✓ | |
| **2020** | | | ✓ | |
| **2019** | | ✓ | ✓ | |
| **2018** | | ✓ | | |

**Classification:** The table above shows all journal classifications recorded for each classification year in the Green Journals database. A journal may have more than one classification in the same year.

## Recent Articles

**Correction to: A quantitative systems pharmacology (QSP) model for Pneumocystis treatment in mice**
2019-8-12 Vol. 13 Issue 1
Authors: Guan-Sheng Liu, Richard Ballweg, Alan Ashbaugh et al.
DOI: 10.1186/s12918-019-0708-9

**Identification of Hürthle cell cancers: solving a clinical challenge with genomic sequencing and a trio of machine learning algorithms**
2019-4 Vol. 13 Issue S2
Authors: Yangyang Hao, Quan-Yang Duh, Richard T. Kloos et al.
DOI: 10.1186/s12918-019-0693-z

**FCMDAP: using miRNA family and cluster information to improve the prediction accuracy of disease related miRNAs**
2019-4 Vol. 13 Issue S2
Authors: Xiaoying Li, Yaping Lin, Changlong Gu et al.
DOI: 10.1186/s12918-019-0696-9

**Fusing gene expressions and transitive protein-protein interactions for inference of gene regulatory networks**
2019-4 Vol. 13 Issue S2
Authors: Wenting Liu, Jagath C. Rajapakse
DOI: 10.1186/s12918-019-0695-x

**GNE: a deep learning framework for gene network inference by aggregating biological information**
2019-4 Vol. 13 Issue S2
Authors: Kishan KC, Rui Li, Feng Cui et al.
DOI: 10.1186/s12918-019-0694-y

**Predicting disease-related phenotypes using an integrated phenotype similarity measurement based on HPO**
2019-4 Vol. 13 Issue S2
Authors: Hansheng Xue, Jiajie Peng, Xuequn Shang
DOI: 10.1186/s12918-019-0697-8

**Ultrafast clustering of single-cell flow cytometry data using FlowGrid**
2019-4 Vol. 13 Issue S2
Authors: Xiaoxin Ye, Joshua W. K. Ho
DOI: 10.1186/s12918-019-0690-2

**Anti-TNF- αtreatment-related pathways and biomarkers revealed by transcriptome analysis in Chinese psoriasis patients**
2019-4 Vol. 13 Issue S2
Authors: Lunfei Liu, Wenting Liu, Yuxin Zheng et al.
DOI: 10.1186/s12918-019-0698-7

**Boolean network modeling of β-cell apoptosis and insulin resistance in type 2 diabetes mellitus**
2019-4 Vol. 13 Issue S2
Authors: Pritha Dutta, Lichun Ma, Yusuf Ali et al.
DOI: 10.1186/s12918-019-0692-0

**Network-based characterization of drug-protein interaction signatures with a space-efficient approach**
2019-4 Vol. 13 Issue S2
Authors: Yasuo Tabei, Masaaki Kotera, Ryusuke Sawada et al.
DOI: 10.1186/s12918-019-0691-1