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Journal Information

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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

DFRL-Mol: a dual-stage framework of reinforcement learning for multi-scenario molecule optimization
2026-9-1 Vol. 27 Issue 5
Authors: Zhenyi Wu, Pengcheng Zhao, Jianing Li et al.
DOI: 10.1093/bib/bbag493

TransSurv: compatibility-aware transfer learning for target-cohort survival prediction
2026-9-1 Vol. 27 Issue 5
Authors: Zheng Qu, Hongzhe Zhang, Katherine L Nathanson et al.
DOI: 10.1093/bib/bbag532

Supervised and covariate-adjusted dimension reduction for prediction and visualization using microbiome data
2026-9-1 Vol. 27 Issue 5
Authors: Crystal Zang, Lu Tang, Rebecca A Deek
DOI: 10.1093/bib/bbag590

MMP2Mol: a matched molecular pairs-based framework for ligand-based de novo drug design
2026-9-1 Vol. 27 Issue 5
Authors: Dekun Chen, Hui Li, Jianwang Liu et al.
DOI: 10.1093/bib/bbag548

MHCXGraph: a graph-based approach to detecting T-cell receptor cross-reactivity
2026-9-1 Vol. 27 Issue 5
Authors: Carlos D M S Simões, Rocío L B R Maidana, Samuel C de Assis et al.
DOI: 10.1093/bib/bbag555

PMPIHGLL: predicting metabolite–protein interactions using dual hypergraph convolutional networks and large language models
2026-9-1 Vol. 27 Issue 5
Authors: Lei Chen, Zhitong Jin, Ying Shao et al.
DOI: 10.1093/bib/bbag489

spMosaic: multi-sample integration for scalable spatial domain discovery based on spatial transcriptomics
2026-9-1 Vol. 27 Issue 5
Authors: Shiwei Fu, Han Li, Wei Vivian Li
DOI: 10.1093/bib/bbag527

Dynamic perturbation analysis: a modular rewiring framework for decoding cancer stage progression and identifying therapeutic vulnerabilities
2026-9-1 Vol. 27 Issue 5
Authors: Bin Yang, Mi Li, Xingyi Liu et al.
DOI: 10.1093/bib/bbag572

Correction to: scAnno: a deconvolution strategy-based automatic cell type annotation tool for single-cell RNA-sequencing data sets
2026-9-1 Vol. 27 Issue 5
DOI: 10.1093/bib/bbag462

Correction to: Master of Metals2: a graph neural network based architecture for the prediction of zinc binding sites in protein structures
2026-9-1 Vol. 27 Issue 5
DOI: 10.1093/bib/bbag617