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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

EvoSeq-ML: Advancing Data-Centric Machine Learning with Evolutionary-Informed Protein Sequence Representation and Generation
2026-11-3
Authors: Mehrsa Mardikoraem, Nathaniel Pascual, Joelle N Eaves et al.
DOI: 10.1093/protein/gzag033

A tethered yeast surface display system facilitates enriching epitope-specific binding interactions
2026 Vol. 39
Authors: Rojhae A Panton, Jakob Lind, Amanda Richardson et al.
DOI: 10.1093/protein/gzag019

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2026 Vol. 39
Authors: Melissa A Walsh, Paul L Blanchard, Xihong Zhang et al.
DOI: 10.1093/protein/gzag025

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2026 Vol. 39
Authors: Logan Mavar, Mikhail Pavlenok, Avraneel Paul et al.
DOI: 10.1093/protein/gzag030

Rapid selection of high-affinity DARPins using click display
2026 Vol. 39
Authors: Joshua D Bryant, Yu Zeng, Monisha Yerram et al.
DOI: 10.1093/protein/gzag028

Towards directed evolution of rare earth element binding peptides using genetically encoded sensitizing amino acids
2026 Vol. 39
Authors: Riley M Bednar, Jason G Marmorstein, Kathleen J Stebe et al.
DOI: 10.1093/protein/gzag026

Prioritizing stability-enhancing mutations using the ESM protein language model in conjunction with physics-based MM/GBSA predictions
2026 Vol. 39
Authors: Emily R Rhodes, Guido Scarabelli, Jonathan Jou et al.
DOI: 10.1093/protein/gzag016

Free energy–guided Q302P mutation in GH1 β -glucosidase from Aspergillus fumigatus improves enzyme function via catalytic tunnel perturbation
2026 Vol. 39
Authors: Shalini Das, Subba Reddy Dodda, Musaddique Hossain et al.
DOI: 10.1093/protein/gzag015

SARS-CoV-2 Orf3a protein interaction mapping using unnatural amino acid incorporation
2026 Vol. 39
Authors: Eryn Lundrigan, Noreen Ahmed, John Paul Pezacki
DOI: 10.1093/protein/gzag024

Mapping functional dynamics hotspots for protein engineering with NMR peak intensity analysis
2026 Vol. 39
Authors: Adam M Damry, Serena E Hunt, Sandrine Legault et al.
DOI: 10.1093/protein/gzag014