

# Brian K Radak

## List of Publications by Year in descending order

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Version: 2024-02-01

25  
papers

2,521  
citations

567281

15  
h-index

552781

26  
g-index

28  
all docs

28  
docs citations

28  
times ranked

3081  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Precise Binding Free Energy Calculations for Multiple Molecules Using an Optimal Measurement Network of Pairwise Differences. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 650-663.                              | 5.3  | 7         |
| 2  | Fast Equilibration of Water between Buried Sites and the Bulk by Molecular Dynamics with Parallel Monte Carlo Water Moves on Graphical Processing Units. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 7366-7372. | 5.3  | 16        |
| 3  | Improved Alchemical Free Energy Calculations with Optimized Smoothstep Softcore Potentials. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 5512-5525.  | 5.3  | 35        |
| 4  | Accounting for the Central Role of Interfacial Water in Protein-Ligand Binding Free Energy Calculations. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 7883-7894.   | 5.3  | 24        |
| 5  | Scalable molecular dynamics on CPU and GPU architectures with NAMD. <i>Journal of Chemical Physics</i> , 2020, 153, 044130.   | 3.0  | 1,548     |
| 6  | Alchemical Binding Free Energy Calculations in AMBER20: Advances and Best Practices for Drug Discovery. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 5595-5623.  | 5.4  | 177       |
| 7  | Boosting Free-Energy Perturbation Calculations with GPU-Accelerated NAMD. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 5301-5307.  | 5.4  | 37        |
| 8  | Rigorous Free Energy Simulations in Virtual Screening. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 4153-4169.   | 5.4  | 114       |
| 9  | Finite-sample bias in free energy bridge estimators. <i>Journal of Chemical Physics</i> , 2019, 151, 034105.  | 3.0  | 1         |
| 10 | Enhanced configurational sampling with hybrid non-equilibrium molecular dynamics-Monte Carlo propagator. <i>Journal of Chemical Physics</i> , 2018, 148, 014101.  | 3.0  | 26        |
| 11 | A generalized linear response framework for expanded ensemble and replica exchange simulations. <i>Journal of Chemical Physics</i> , 2018, 149, 072315.   | 3.0  | 7         |
| 12 | A Multidimensional B-Spline Correction for Accurate Modeling Sugar Puckering in QM/MM Simulations. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 3975-3984.   | 5.3  | 12        |
| 13 | Constant-pH Molecular Dynamics Simulations for Large Biomolecular Systems. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 5933-5944.   | 5.3  | 139       |
| 14 | Efficiency in nonequilibrium molecular dynamics Monte Carlo simulations. <i>Journal of Chemical Physics</i> , 2016, 145, 134109.  | 3.0  | 15        |
| 15 | A Two-Metal-Ion-Mediated Conformational Switching Pathway for HDV Ribozyme Activation. <i>ACS Catalysis</i> , 2016, 6, 1853-1869.   | 11.2 | 24        |
| 16 | Assessment of metal-assisted nucleophile activation in the hepatitis delta virus ribozyme from molecular simulation and 3D-RISM. <i>Rna</i> , 2015, 21, 1566-1577.  | 3.5  | 18        |
| 17 | Characterization of the Three-Dimensional Free Energy Manifold for the Uracil Ribonucleoside from Asynchronous Replica Exchange Simulations. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 373-377.               | 5.3  | 10        |
| 18 | Multiscale Methods for Computational RNA Enzymology. <i>Methods in Enzymology</i> , 2015, 553, 335-374.   | 1.0  | 16        |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 19 | Roadmaps through Free Energy Landscapes Calculated Using the Multidimensional vFEP Approach. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 24-34.  | 5.3 | 58        |
| 20 | A Variational Linear-Scaling Framework to Build Practical, Efficient Next-Generation Orbital-Based Quantum Force Fields. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 1417-1427.   | 5.3 | 55        |
| 21 | A New Maximum Likelihood Approach for Free Energy Profile Construction from Molecular Simulations. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 153-164.   | 5.3 | 76        |
| 22 | Molecular Simulations of RNA 2'-O <sup>2</sup> -Transesterification Reaction Models in Solution. <i>Journal of Physical Chemistry B</i> , 2013, 117, 94-103.   | 2.6 | 21        |
| 23 | Experimental and computational analysis of the transition state for ribonuclease A-catalyzed RNA 2'-O <sup>2</sup> -transphosphorylation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 13002-13007. | 7.1 | 62        |
| 24 | Modeling Reactive Scattering of F <sub>2</sub> at a Liquid Squalane Interface: A Hybrid QM/MM Molecular Dynamics Study. <i>Journal of Physical Chemistry A</i> , 2009, 113, 7218-7226.   | 2.5 | 13        |
| 25 | Modeling Ion Channels Using Poisson-Nernst-Planck Theory as an Integrated Approach To Introducing Nanotechnology Concepts: The PNP Cyclic Peptide Ion Channel Model. <i>Journal of Chemical Education</i> , 2008, 85, 744.                                 | 2.3 | 2         |