

Unbiased Rare Event Sampling in Spatial Stochastic Systems Using a Weighted Ensemble of Trajectories

PLoS Computational Biology

12, e1004611

DOI: [10.1371/journal.pcbi.1004611](https://doi.org/10.1371/journal.pcbi.1004611)

Citation Report

#	ARTICLE	IF	CITATIONS
1	Efficient Atomistic Simulation of Pathways and Calculation of Rate Constants for a Protein–Peptide Binding Process: Application to the MDM2 Protein and an Intrinsically Disordered p53 Peptide. <i>Journal of Physical Chemistry Letters</i> , 2016, 7, 3440-3445.	2.1	94
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8	MCell-R: A Particle-Resolution Network-Free Spatial Modeling Framework. <i>Methods in Molecular Biology</i> , 2019, 1945, 203-229.	0.4	17
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15	A first-passage approach to diffusion-influenced reversible binding and its insights into nanoscale signaling at the presynapse. <i>Scientific Reports</i> , 2021, 11, 5377.	1.6	15
18	Quantifying the roles of space and stochasticity in computer simulations for cell biology and cellular biochemistry. <i>Molecular Biology of the Cell</i> , 2021, 32, 186-210.	0.9	18
19	Wepy: A Flexible Software Framework for Simulating Rare Events with Weighted Ensemble Resampling. <i>ACS Omega</i> , 2020, 5, 31608-31623.	1.6	21
21	A Suite of Tutorials for the WESTPA Rare-Events Sampling Software [Article v1.0]. <i>Living Journal of Computational Molecular Science</i> , 2019, 1, .	2.2	16

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25	A gentle introduction to the non-equilibrium physics of trajectories: Theory, algorithms, and biomolecular applications. American Journal of Physics, 2021, 89, 1048-1061.	0.3	7
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