

Unsupervised Learning Methods for Molecular Simulation

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Citation Report

#	ARTICLE	IF	CITATIONS
2	Physics-Inspired Structural Representations for Molecules and Materials. <i>Chemical Reviews</i> , 2021, 121, 9759-9815.	23.0	247
3	Markov State Models to Study the Functional Dynamics of Proteins in the Wake of Machine Learning. <i>Jacs Au</i> , 2021, 1, 1330-1341.	3.6	56
5	Computational strategies for protein conformational ensemble detection. <i>Current Opinion in Structural Biology</i> , 2022, 72, 79-87.	2.6	6
6	Variational embedding of protein folding simulations using Gaussian mixture variational autoencoders. <i>Journal of Chemical Physics</i> , 2021, 155, 194108.	1.2	11
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20	From Data to Knowledge: Systematic Review of Tools for Automatic Analysis of Molecular Dynamics Output. <i>Frontiers in Pharmacology</i> , 2022, 13, 844293.	1.6	1

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