

Shunyang Wang

List of Publications by Year in descending order

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71
papers

7,858
citations

136950

32
h-index

91884

69
g-index

83
all docs

83
docs citations

83
times ranked

9262
citing authors

#	ARTICLE	IF	CITATIONS
1	Proposed Mechanism for the Biosynthesis of the [FeFe] Hydrogenase H-Cluster: Central Roles for the Radical SAM Enzymes HydG and HydE. <i>ACS Bio & Med Chem Au</i> , 2022, 2, 11-21.	3.7	6
2	Accumulation and Pulse Electron Paramagnetic Resonance Spectroscopic Investigation of the 4-Oxidobenzyl Radical Generated in the Radical S-Adenosyl-methionine Enzyme HydG. <i>Biochemistry</i> , 2022, 61, 107-116.	2.5	7
3	Quantum Chemical Prediction of Electron Ionization Mass Spectra of Trimethylsilylated Metabolites. <i>Analytical Chemistry</i> , 2022, , .	6.5	5
4	Evaluating the Accuracy of the QCEIMS Approach for Computational Prediction of Electron Ionization Mass Spectra of Purines and Pyrimidines. <i>Metabolites</i> , 2022, 12, 68.	2.9	4
5	Improving Force Field Accuracy by Training against Condensed-Phase Mixture Properties. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 3577-3592.	5.3	9
6	Open Force Field Evaluator: An Automated, Efficient, and Scalable Framework for the Estimation of Physical Properties from Molecular Simulation. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 3566-3576.	5.3	19
7	Source of Rate Acceleration for Carbocation Cyclization in Biomimetic Supramolecular Cages. <i>Journal of the American Chemical Society</i> , 2022, 144, 11413-11424.	13.7	15
8	Exploration and validation of force field design protocols through QM-to-MM mapping. <i>Physical Chemistry Chemical Physics</i> , 2022, 24, 17014-17027.	2.8	4
9	TeraChem: A graphical processing unit-accelerated electronic structure package for large-scale ab initio molecular dynamics. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2021, 11, e1494.	14.6	143
10	Reversible O-Acetyl Migration within the Sialic Acid Side Chain and Its Influence on Protein Recognition. <i>ACS Chemical Biology</i> , 2021, 16, 1951-1960.	3.4	19
11	Quantum Chemistry Calculations for Metabolomics. <i>Chemical Reviews</i> , 2021, 121, 5633-5670.	47.7	47
12	Resurrected Ancestors Reveal Origins of Metamorphism in XCL1. <i>Trends in Biochemical Sciences</i> , 2021, 46, 433-434.	7.5	0
13	Identification and characterization of metamorphic proteins: Current and future perspectives. <i>Biopolymers</i> , 2021, 112, e23473.	2.4	6
14	Development and Benchmarking of Open Force Field v1.0.0—the Parsley Small-Molecule Force Field. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 6262-6280.	5.3	80
15	Quantum Chemical Study of a Radical Relay Mechanism for the HydG-Catalyzed Synthesis of a Fe(II)(CO) ₂ (CN) ₂ cysteine Precursor to the H-Cluster of [FeFe] Hydrogenase. <i>Biochemistry</i> , 2021, 60, 3016-3026.	2.5	4
16	The automated optimisation of a coarse-grained force field using free energy data. <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 24842-24851.	2.8	3
17	Development and Validation of AMBER-FB15-Compatible Force Field Parameters for Phosphorylated Amino Acids. <i>Journal of Physical Chemistry B</i> , 2021, 125, 11927-11942.	2.6	8
18	Quantum Chemistry Common Driver and Databases (QCDB) and Quantum Chemistry Engine (QCEngine): Automation and interoperability among computational chemistry programs. <i>Journal of Chemical Physics</i> , 2021, 155, 204801.	3.0	15

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19	Data-Driven Mapping of Gas-Phase Quantum Calculations to General Force Field Lennard-Jones Parameters. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 1115-1127.	5.3	15
20	Sequence-Based Prediction of Metamorphic Behavior in Proteins. <i>Biophysical Journal</i> , 2020, 119, 1380-1390.	0.5	20
21	Data-driven analysis of the number of Lennard-Jones types needed in a force field. <i>Communications Chemistry</i> , 2020, 3, .	4.5	6
22	Linkage between Proximal and Distal Movements of P450cam Induced by Putidaredoxin. <i>Biochemistry</i> , 2020, 59, 2012-2021.	2.5	6
23	A combined NMR, MD and DFT conformational analysis of 9-O-acetyl sialic acid-containing GM3 ganglioside glycan and its 9-N-acetyl mimic. <i>Glycobiology</i> , 2020, 30, 787-801.	2.5	17
24	Driving torsion scans with wavefront propagation. <i>Journal of Chemical Physics</i> , 2020, 152, 244116.	3.0	22
25	Bond-Order Time Series Analysis for Detecting Reaction Events in <i>Ab Initio</i> Molecular Dynamics Simulations. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 1606-1617.	5.3	12
26	Non-bonded force field model with advanced restrained electrostatic potential charges (RESP2). <i>Communications Chemistry</i> , 2020, 3, .	4.5	98
27	Car-Parrinello Monitor for More Robust Born-Oppenheimer Molecular Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 4454-4467.	5.3	5
28	Systematic Optimization of Water Models Using Liquid/Vapor Surface Tension Data. <i>Journal of Physical Chemistry B</i> , 2019, 123, 7061-7073.	2.6	31
29	Capillary Effects on Groundwater Response to Earth Tides. <i>Water Resources Research</i> , 2019, 55, 6886-6895.	4.2	18
30	Binding Thermodynamics of Host-Guest Systems with SMIRNOFF99Frosst 1.0.5 from the Open Force Field Initiative. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 6225-6242.	5.3	21
31	Conformational Response of N-Terminally Truncated Cytochrome P450 3A4 to Ligand Binding in Solution. <i>Biochemistry</i> , 2019, 58, 3903-3910.	2.5	12
32	An Intermediate Conformational State of Cytochrome P450cam-CN in Complex with Putidaredoxin. <i>Biochemistry</i> , 2019, 58, 2353-2361.	2.5	12
33	Force Field Development and Nanoreactor Chemistry. <i>Challenges and Advances in Computational Chemistry and Physics</i> , 2019, , 127-159.	0.6	1
34	Quantum chemical studies of redox properties and conformational changes of a four-center iron CO ₂ reduction electrocatalyst. <i>Chemical Science</i> , 2018, 9, 2645-2654.	7.4	6
35	Advanced models for water simulations. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2018, 8, e1355.	14.6	42
36	Assimilating Radial Distribution Functions To Build Water Models with Improved Structural Properties. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 1766-1778.	5.4	22

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37	Polarizable Molecular Simulations Reveal How Silicon-Containing Functional Groups Govern the Desalination Mechanism in Nanoporous Graphene. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 4279-4290.	5.3	8
38	An Estimation of Hybrid Quantum Mechanical Molecular Mechanical Polarization Energies for Small Molecules Using Polarizable Force-Field Approaches. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 679-695.	5.3	19
39	A Chemical Biology Solution to Problems with Studying Biologically Important but Unstable 9-O-Acetyl Sialic Acids. <i>ACS Chemical Biology</i> , 2017, 12, 214-224.	3.4	37
40	Building a More Predictive Protein Force Field: A Systematic and Reproducible Route to AMBER-FB15. <i>Journal of Physical Chemistry B</i> , 2017, 121, 4023-4039.	2.6	192
41	OpenMM 7: Rapid development of high performance algorithms for molecular dynamics. <i>PLoS Computational Biology</i> , 2017, 13, e1005659.	3.2	1,561
42	Geometry optimization made simple with translation and rotation coordinates. <i>Journal of Chemical Physics</i> , 2016, 144, 214108.	3.0	137
43	Advanced Potential Energy Surfaces for Molecular Simulation. <i>Journal of Physical Chemistry B</i> , 2016, 120, 9811-9832.	2.6	77
44	Large earthquakes create vertical permeability by breaching aquitards. <i>Water Resources Research</i> , 2016, 52, 5923-5937.	4.2	75
45	Training and Validation of a Liquid-Crystalline Phospholipid Bilayer Force Field. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 5960-5967.	5.3	14
46	Automated Code Engine for Graphical Processing Units: Application to the Effective Core Potential Integrals and Gradients. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 92-106.	5.3	55
47	Automated Discovery and Refinement of Reactive Molecular Dynamics Pathways. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 638-649.	5.3	95
48	Efficient implementation of effective core potential integrals and gradients on graphical processing units. <i>Journal of Chemical Physics</i> , 2015, 143, 014114.	3.0	17
49	Tensor Hypercontraction Second-Order Møller-Plesset Perturbation Theory: Grid Optimization and Reaction Energies. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 3042-3052.	5.3	47
50	Revised Parameters for the AMOEBA Polarizable Atomic Multipole Water Model. <i>Journal of Physical Chemistry B</i> , 2015, 119, 9423-9437.	2.6	183
51	Why many semiempirical molecular orbital theories fail for liquid water and how to fix them. <i>Journal of Computational Chemistry</i> , 2015, 36, 934-939.	3.3	16
52	United polarizable multipole water model for molecular mechanics simulation. <i>Journal of Chemical Physics</i> , 2015, 143, 014504.	3.0	36
53	MDTraj: A Modern Open Library for the Analysis of Molecular Dynamics Trajectories. <i>Biophysical Journal</i> , 2015, 109, 1528-1532.	0.5	1,576
54	What Can Density Functional Theory Tell Us about Artificial Catalytic Water Splitting?. <i>Inorganic Chemistry</i> , 2014, 53, 6386-6397.	4.0	126

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55	Discovering chemistry with an ab initio nanoreactor. <i>Nature Chemistry</i> , 2014, 6, 1044-1048.	13.6	286
56	Building Force Fields: An Automatic, Systematic, and Reproducible Approach. <i>Journal of Physical Chemistry Letters</i> , 2014, 5, 1885-1891.	4.6	400
57	Modeling Organochlorine Compounds and the Īf-Hole Effect Using a Polarizable Multipole Force Field. <i>Journal of Physical Chemistry B</i> , 2014, 118, 6456-6465.	2.6	69
58	Systematic Improvement of a Classical Molecular Model of Water. <i>Journal of Physical Chemistry B</i> , 2013, 117, 9956-9972.	2.6	279
59	Calculations of the Electric Fields in Liquid Solutions. <i>Journal of Physical Chemistry B</i> , 2013, 117, 16236-16248.	2.6	83
60	A pathway to diphosphorus from the dissociation of photoexcited tetraphosphorus. <i>RSC Advances</i> , 2013, 3, 23166.	3.6	14
61	OpenMM 4: A Reusable, Extensible, Hardware Independent Library for High Performance Molecular Simulation. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 461-469.	5.3	583
62	Systematic Parametrization of Polarizable Force Fields from Quantum Chemistry Data. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 452-460.	5.3	164
63	Basinâ€scale transport of heat and fluid induced by earthquakes. <i>Geophysical Research Letters</i> , 2013, 40, 3893-3897.	4.0	41
64	A Polarizable QM/MM Explicit Solvent Model for Computational Electrochemistry in Water. <i>Journal of Chemical Theory and Computation</i> , 2012, 8, 610-617.	5.3	71
65	Simulation of Solution Phase Electron Transfer in a Compact Donorâ€Acceptor Dyad. <i>Journal of Physical Chemistry B</i> , 2011, 115, 12135-12144.	2.6	27
66	Molecular Insight Into the Energy Levels at the Organic Donor/Acceptor Interface: A Quantum Mechanics/Molecular Mechanics Study. <i>Journal of Physical Chemistry C</i> , 2011, 115, 14431-14436.	3.1	83
67	Direct-Coupling O ₂ Bond Forming a Pathway in Cobalt Oxide Water Oxidation Catalysts. <i>Journal of Physical Chemistry Letters</i> , 2011, 2, 2200-2204.	4.6	177
68	Communication: Hybrid ensembles for improved force matching. <i>Journal of Chemical Physics</i> , 2010, 133, 231101.	3.0	19
69	The Diabatic Picture of Electron Transfer, Reaction Barriers, and Molecular Dynamics. <i>Annual Review of Physical Chemistry</i> , 2010, 61, 149-170.	10.8	280
70	Electronic Properties of Disordered Organic Semiconductors via QM/MM Simulations. <i>Accounts of Chemical Research</i> , 2010, 43, 995-1004.	15.6	90
71	Acidâ€Base Mechanism for Ruthenium Water Oxidation Catalysts. <i>Inorganic Chemistry</i> , 2010, 49, 4543-4553.	4.0	139