

Shunyang Wang

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

70
papers

4,977
citations

29
h-index

70
g-index

83
ext. papers

6,520
ext. citations

6.4
avg. IF

5.78
L-index

| # | Paper | IF | Citations |
|----|--|------|-----------|
| 70 | Accumulation and Pulse Electron Paramagnetic Resonance Spectroscopic Investigation of the 4-Oxidobenzyl Radical Generated in the Radical -Adenosyl-l-methionine Enzyme HydG.. <i>Biochemistry</i> , 2022 , | 3.2 | 2 |
| 69 | Quantum Chemical Prediction of Electron Ionization Mass Spectra of Trimethylsilylated Metabolites.. <i>Analytical Chemistry</i> , 2022 , | 7.8 | 1 |
| 68 | 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 | | 0 |
| 67 | 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 | 3.9 | 3 |
| 66 | The automated optimisation of a coarse-grained force field using free energy data. <i>Physical Chemistry Chemical Physics</i> , 2021 , 23, 24842-24851 | 3.6 | 2 |
| 65 | 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 | 3.4 | 1 |
| 64 | Reversible -Acetyl Migration within the Sialic Acid Side Chain and Its Influence on Protein Recognition. <i>ACS Chemical Biology</i> , 2021 , 16, 1951-1960 | 4.9 | 5 |
| 63 | Quantum Chemistry Calculations for Metabolomics. <i>Chemical Reviews</i> , 2021 , 121, 5633-5670 | 68.1 | 18 |
| 62 | Resurrected Ancestors Reveal Origins of Metamorphism in XCL1. <i>Trends in Biochemical Sciences</i> , 2021 , 46, 433-434 | 10.3 | |
| 61 | 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 | 7.9 | 49 |
| 60 | Identification and characterization of metamorphic proteins: Current and future perspectives. <i>Biopolymers</i> , 2021 , 112, e23473 | 2.2 | 1 |
| 59 | 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 | 6.4 | 12 |
| 58 | Quantum Chemical Study of a Radical Relay Mechanism for the HydG-Catalyzed Synthesis of a Fe(II)(CO)(CN)cysteine Precursor to the H-Cluster of [FeFe] Hydrogenase. <i>Biochemistry</i> , 2021 , 60, 3016-3026 | 3.2 | 3 |
| 57 | Linkage between Proximal and Distal Movements of P450cam Induced by Putidaredoxin. <i>Biochemistry</i> , 2020 , 59, 2012-2021 | 3.2 | 4 |
| 56 | 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 | 5.8 | 9 |
| 55 | Driving torsion scans with wavefront propagation. <i>Journal of Chemical Physics</i> , 2020 , 152, 244116 | 3.9 | 10 |
| 54 | Bond-Order Time Series Analysis for Detecting Reaction Events in Molecular Dynamics Simulations. <i>Journal of Chemical Theory and Computation</i> , 2020 , 16, 1606-1617 | 6.4 | 4 |

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| 53 | 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 | 6.4 | 8 |
| 52 | Sequence-Based Prediction of Metamorphic Behavior in Proteins. <i>Biophysical Journal</i> , 2020 , 119, 1380-1390 | 6.3 | 6 |
| 51 | Data-driven analysis of the number of Lennard-Jones types needed in a force field. <i>Communications Chemistry</i> , 2020 , 3, | 6.3 | 1 |
| 50 | Non-bonded force field model with advanced restrained electrostatic potential charges (RESP2). <i>Communications Chemistry</i> , 2020 , 3, | 6.3 | 33 |
| 49 | Conformational Response of N-Terminally Truncated Cytochrome P450 3A4 to Ligand Binding in Solution. <i>Biochemistry</i> , 2019 , 58, 3903-3910 | 3.2 | 8 |
| 48 | An Intermediate Conformational State of Cytochrome P450cam-CN in Complex with Putidaredoxin. <i>Biochemistry</i> , 2019 , 58, 2353-2361 | 3.2 | 8 |
| 47 | Force Field Development and Nanoreactor Chemistry. <i>Challenges and Advances in Computational Chemistry and Physics</i> , 2019 , 127-159 | 0.7 | |
| 46 | Car-Parrinello Monitor for More Robust Born-Oppenheimer Molecular Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2019 , 15, 4454-4467 | 6.4 | 5 |
| 45 | Systematic Optimization of Water Models Using Liquid/Vapor Surface Tension Data. <i>Journal of Physical Chemistry B</i> , 2019 , 123, 7061-7073 | 3.4 | 18 |
| 44 | Capillary Effects on Groundwater Response to Earth Tides. <i>Water Resources Research</i> , 2019 , 55, 6886-6895 | 3.4 | 12 |
| 43 | 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 | 6.4 | 11 |
| 42 | Quantum chemical studies of redox properties and conformational changes of a four-center iron CO reduction electrocatalyst. <i>Chemical Science</i> , 2018 , 9, 2645-2654 | 9.4 | 4 |
| 41 | Assimilating Radial Distribution Functions To Build Water Models with Improved Structural Properties. <i>Journal of Chemical Information and Modeling</i> , 2018 , 58, 1766-1778 | 6.1 | 12 |
| 40 | 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 | 6.4 | 6 |
| 39 | Advanced models for water simulations. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2018 , 8, e1355 | 7.9 | 30 |
| 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 | 6.4 | 14 |
| 37 | 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 | 4.9 | 27 |
| 36 | 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 | 3.4 | 147 |

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|----|---|------|-----|
| 35 | OpenMM 7: Rapid development of high performance algorithms for molecular dynamics. <i>PLoS Computational Biology</i> , 2017 , 13, e1005659 | 5 | 686 |
| 34 | Large earthquakes create vertical permeability by breaching aquitards. <i>Water Resources Research</i> , 2016 , 52, 5923-5937 | 5.4 | 59 |
| 33 | Training and Validation of a Liquid-Crystalline Phospholipid Bilayer Force Field. <i>Journal of Chemical Theory and Computation</i> , 2016 , 12, 5960-5967 | 6.4 | 12 |
| 32 | 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 | 6.4 | 46 |
| 31 | Automated Discovery and Refinement of Reactive Molecular Dynamics Pathways. <i>Journal of Chemical Theory and Computation</i> , 2016 , 12, 638-49 | 6.4 | 73 |
| 30 | Geometry optimization made simple with translation and rotation coordinates. <i>Journal of Chemical Physics</i> , 2016 , 144, 214108 | 3.9 | 71 |
| 29 | Advanced Potential Energy Surfaces for Molecular Simulation. <i>Journal of Physical Chemistry B</i> , 2016 , 120, 9811-32 | 3.4 | 66 |
| 28 | Why many semiempirical molecular orbital theories fail for liquid water and how to fix them. <i>Journal of Computational Chemistry</i> , 2015 , 36, 934-9 | 3.5 | 12 |
| 27 | United polarizable multipole water model for molecular mechanics simulation. <i>Journal of Chemical Physics</i> , 2015 , 143, 014504 | 3.9 | 30 |
| 26 | MDTraj: A Modern Open Library for the Analysis of Molecular Dynamics Trajectories. <i>Biophysical Journal</i> , 2015 , 109, 1528-32 | 2.9 | 834 |
| 25 | Efficient implementation of effective core potential integrals and gradients on graphical processing units. <i>Journal of Chemical Physics</i> , 2015 , 143, 014114 | 3.9 | 13 |
| 24 | Tensor Hypercontraction Second-Order Møller-Plesset Perturbation Theory: Grid Optimization and Reaction Energies. <i>Journal of Chemical Theory and Computation</i> , 2015 , 11, 3042-52 | 6.4 | 38 |
| 23 | Revised Parameters for the AMOEBA Polarizable Atomic Multipole Water Model. <i>Journal of Physical Chemistry B</i> , 2015 , 119, 9423-9437 | 3.4 | 149 |
| 22 | What can density functional theory tell us about artificial catalytic water splitting?. <i>Inorganic Chemistry</i> , 2014 , 53, 6386-97 | 5.1 | 97 |
| 21 | Discovering chemistry with an ab initio nanoreactor. <i>Nature Chemistry</i> , 2014 , 6, 1044-8 | 17.6 | 209 |
| 20 | Building Force Fields: An Automatic, Systematic, and Reproducible Approach. <i>Journal of Physical Chemistry Letters</i> , 2014 , 5, 1885-91 | 6.4 | 294 |
| 19 | Modeling organochlorine compounds and the Ehole effect using a polarizable multipole force field. <i>Journal of Physical Chemistry B</i> , 2014 , 118, 6456-65 | 3.4 | 65 |
| 18 | Systematic improvement of a classical molecular model of water. <i>Journal of Physical Chemistry B</i> , 2013 , 117, 9956-72 | 3.4 | 235 |

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| 17 | Calculations of the electric fields in liquid solutions. <i>Journal of Physical Chemistry B</i> , 2013 , 117, 16236-48 | 3.4 | 73 |
| 16 | A pathway to diphosphorus from the dissociation of photoexcited tetraphosphorus. <i>RSC Advances</i> , 2013 , 3, 23166 | 3.7 | 11 |
| 15 | 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 | 6.4 | 440 |
| 14 | Systematic Parametrization of Polarizable Force Fields from Quantum Chemistry Data. <i>Journal of Chemical Theory and Computation</i> , 2013 , 9, 452-60 | 6.4 | 124 |
| 13 | Basin-scale transport of heat and fluid induced by earthquakes. <i>Geophysical Research Letters</i> , 2013 , 40, 3893-3897 | 4.9 | 34 |
| 12 | A Polarizable QM/MM Explicit Solvent Model for Computational Electrochemistry in Water. <i>Journal of Chemical Theory and Computation</i> , 2012 , 8, 610-7 | 6.4 | 63 |
| 11 | Simulation of solution phase electron transfer in a compact donor-acceptor dyad. <i>Journal of Physical Chemistry B</i> , 2011 , 115, 12135-44 | 3.4 | 24 |
| 10 | 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 | 3.8 | 82 |
| 9 | Direct-Coupling O2 Bond Forming a Pathway in Cobalt Oxide Water Oxidation Catalysts. <i>Journal of Physical Chemistry Letters</i> , 2011 , 2, 2200-2204 | 6.4 | 159 |
| 8 | Electronic properties of disordered organic semiconductors via QM/MM simulations. <i>Accounts of Chemical Research</i> , 2010 , 43, 995-1004 | 24.3 | 83 |
| 7 | Acid-base mechanism for ruthenium water oxidation catalysts. <i>Inorganic Chemistry</i> , 2010 , 49, 4543-53 | 5.1 | 134 |
| 6 | Communication: Hybrid ensembles for improved force matching. <i>Journal of Chemical Physics</i> , 2010 , 133, 231101 | 3.9 | 17 |
| 5 | The diabatic picture of electron transfer, reaction barriers, and molecular dynamics. <i>Annual Review of Physical Chemistry</i> , 2010 , 61, 149-70 | 15.7 | 256 |
| 4 | Force Field Partial Charges with Restrained Electrostatic Potential 2 (RESP2) | | 3 |
| 3 | OpenMM 7: Rapid Development of High Performance Algorithms for Molecular Dynamics | | 1 |
| 2 | The Automated Optimisation of a Coarse-Grained Force Field Using Free Energy Data | | 2 |
| 1 | Capturing non-local through-bond effects when fragmenting molecules for quantum chemical torsion scans | | 3 |