

Marcus Elstner

List of Publications by Year in descending order

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50276

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times ranked

6528
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#	ARTICLE	IF	CITATIONS
1	Hydrogen bonding and stacking interactions of nucleic acid base pairs: A density-functional-theory based treatment. <i>Journal of Chemical Physics</i> , 2001, 114, 5149-5155.	3.0	978
2	DFTB3: Extension of the Self-Consistent-Charge Density-Functional Tight-Binding Method (SCC-DFTB). <i>Journal of Chemical Theory and Computation</i> , 2011, 7, 931-948.	5.3	828
3	Parametrization and Benchmark of DFTB3 for Organic Molecules. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 338-354.	5.3	743
4	A QM/MM Implementation of the Self-Consistent Charge Density Functional Tight Binding (SCC-DFTB) Method. <i>Journal of Physical Chemistry B</i> , 2001, 105, 569-585.	2.6	568
5	Semiempirical Quantum Mechanical Methods for Noncovalent Interactions for Chemical and Biochemical Applications. <i>Chemical Reviews</i> , 2016, 116, 5301-5337.	47.7	312
6	Development of Effective Quantum Mechanical/Molecular Mechanical (QM/MM) Methods for Complex Biological Processes. <i>Journal of Physical Chemistry B</i> , 2006, 110, 6458-6469.	2.6	290
7	Parameterization of DFTB3/3OB for Sulfur and Phosphorus for Chemical and Biological Applications. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 1518-1537.	5.3	275
8	Extension of the Self-Consistent-Charge Density-Functional Tight-Binding Method: Third-Order Expansion of the Density Functional Theory Total Energy and Introduction of a Modified Effective Coulomb Interaction. <i>Journal of Physical Chemistry A</i> , 2007, 111, 10861-10873.	2.5	265
9	Density functional tight binding. <i>Philosophical Transactions Series A, Mathematical, Physical, and Engineering Sciences</i> , 2014, 372, 20120483.	3.4	256
10	Comparison of a QM/MM force field and molecular mechanics force fields in simulations of alanine and glycine dipeptides (Ace-Ala-Nme and Ace-Gly-Nme) in water in relation to the problem of modeling the unfolded peptide backbone in solution. <i>Proteins: Structure, Function and Bioinformatics</i> , 2003, 50, 451-463.	2.6	250
11	Parameterization of the DFTB3 Method for Br, Ca, Cl, F, I, K, and Na in Organic and Biological Systems. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 332-342.	5.3	227
12	Implementation of the SCC-DFTB Method for Hybrid QM/MM Simulations within the Amber Molecular Dynamics Package. <i>Journal of Physical Chemistry A</i> , 2007, 111, 5655-5664.	2.5	213
13	Parameter Calibration of Transition-Metal Elements for the Spin-Polarized Self-Consistent-Charge Density-Functional Tight-Binding (DFTB) Method: Sc, Ti, Fe, Co, and Ni. <i>Journal of Chemical Theory and Computation</i> , 2007, 3, 1349-1367.	5.3	208
14	Color Tuning in Rhodopsins: The Mechanism for the Spectral Shift between Bacteriorhodopsin and Sensory Rhodopsin II. <i>Journal of the American Chemical Society</i> , 2006, 128, 10808-10818.	13.7	196
15	Electronic couplings for molecular charge transfer: Benchmarking CDFT, FODFT, and FODFTB against high-level <i>ab initio</i> calculations. <i>Journal of Chemical Physics</i> , 2014, 140, 104105.	3.0	175
16	An approximate DFT method for QM/MM simulations of biological structures and processes. <i>Computational and Theoretical Chemistry</i> , 2003, 632, 29-41.	1.5	172
17	Density functional tight binding: application to organic and biological molecules. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2014, 4, 49-61.	14.6	157
18	Modeling zinc in biomolecules with the self consistent charge-density functional tight binding (SCC-DFTB) method: Applications to structural and energetic analysis. <i>Journal of Computational Chemistry</i> , 2003, 24, 565-581.	3.3	150

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19	Efficient Calculation of Charge-Transfer Matrix Elements for Hole Transfer in DNA. <i>Journal of Physical Chemistry B</i> , 2008, 112, 7937-7947.	2.6	150
20	Quantum mechanics simulation of protein dynamics on long timescale. <i>Proteins: Structure, Function and Bioinformatics</i> , 2001, 44, 484-489.	2.6	140
21	Validation of the density-functional based tight-binding approximation method for the calculation of reaction energies and other data. <i>Journal of Chemical Physics</i> , 2005, 122, 114110.	3.0	140
22	Parametrization of DFTB3/3OB for Magnesium and Zinc for Chemical and Biological Applications. <i>Journal of Physical Chemistry B</i> , 2015, 119, 1062-1082.	2.6	138
23	Energetics and structure of glycine and alanine based model peptides: Approximate SCC-DFTB, AM1 and PM3 methods in comparison with DFT, HF and MP2 calculations. <i>Chemical Physics</i> , 2001, 263, 203-219.	1.9	132
24	Density functional tight binding: values of semi-empirical methods in an ab initio era. <i>Physical Chemistry Chemical Physics</i> , 2014, 16, 14368-14377.	2.8	125
25	Electronic couplings for molecular charge transfer: benchmarking CDFT, FODFT and FODFTB against high-level ab initio calculations. II. <i>Physical Chemistry Chemical Physics</i> , 2015, 17, 14342-14354.	2.8	119
26	From C2Molecules to Self-Assembled Fullerenes in Quantum Chemical Molecular Dynamics. <i>Nano Letters</i> , 2003, 3, 1657-1664.	9.1	87
27	Description of Phosphate Hydrolysis Reactions with the Self-Consistent-Charge Density-Functional-Tight-Binding (SCC-DFTB) Theory. 1. Parameterization. <i>Journal of Chemical Theory and Computation</i> , 2008, 4, 2067-2084.	5.3	87
28	Application of the SCC-DFTB Method to Neutral and Protonated Water Clusters and Bulk Water. <i>Journal of Physical Chemistry B</i> , 2011, 115, 6790-6805.	2.6	81
29	Simulating Water with the Self-Consistent-Charge Density Functional Tight Binding Method: From Molecular Clusters to the Liquid State. <i>Journal of Physical Chemistry A</i> , 2007, 111, 5685-5691.	2.5	79
30	Coarse-Grained Time-Dependent Density Functional Simulation of Charge Transfer in Complex Systems: Application to Hole Transfer in DNA. <i>Journal of Physical Chemistry B</i> , 2010, 114, 11221-11240.	2.6	79
31	Combined density functional theory and Landauer approach for hole transfer in DNA along classical molecular dynamics trajectories. <i>Journal of Chemical Physics</i> , 2009, 130, 215104.	3.0	78
32	A hybrid approach to simulation of electron transfer in complex molecular systems. <i>Journal of the Royal Society Interface</i> , 2013, 10, 20130415.	3.4	72
33	Automatized Parametrization of SCC-DFTB Repulsive Potentials: Application to Hydrocarbons. <i>Journal of Physical Chemistry A</i> , 2009, 113, 11866-11881.	2.5	69
34	Time-Dependent Extension of the Long-Range Corrected Density Functional Based Tight-Binding Method. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 1737-1747.	5.3	67
35	Effect of Polarization on the Opsin Shift in Rhodopsins. 1. A Combined QM/QM/MM Model for Bacteriorhodopsin and Pharaonis Sensory Rhodopsin II. <i>Journal of Physical Chemistry B</i> , 2008, 112, 11462-11467.	2.6	62
36	Towards an Understanding of Channelrhodopsin Function: Simulations Lead to Novel Insights of the Channel Mechanism. <i>Journal of Molecular Biology</i> , 2013, 425, 1795-1814.	4.2	62

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37	Parametrization and Benchmark of Long-Range Corrected DFTB2 for Organic Molecules. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 115-125.	5.3	60
38	Size-Consistent Multipartitioning QM/MM: A Stable and Efficient Adaptive QM/MM Method. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 4242-4252.	5.3	55
39	Parametrization of the SCC-DFTB Method for Halogens. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 2939-2949.	5.3	54
40	Multi-Scale Approach to Non-Adiabatic Charge Transport in High-Mobility Organic Semiconductors. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 5068-5082.	5.3	53
41	Formation of Fullerene Molecules from Carbon Nanotubes: A Quantum Chemical Molecular Dynamics Study. <i>Nano Letters</i> , 2003, 3, 465-470.	9.1	50
42	Computational photochemistry of retinal proteins. <i>Journal of Computer-Aided Molecular Design</i> , 2006, 20, 511-518.	2.9	50
43	The Protonation State of Glu181 in Rhodopsin Revisited: Interpretation of Experimental Data on the Basis of QM/MM Calculations. <i>Journal of Physical Chemistry B</i> , 2010, 114, 11338-11352.	2.6	50
44	Effect of Polarization on the Opsin Shift in Rhodopsins. 2. Empirical Polarization Models for Proteins. <i>Journal of Physical Chemistry B</i> , 2008, 112, 11468-11478.	2.6	49
45	Reaction dynamics of the chimeric channelrhodopsin C1C2. <i>Scientific Reports</i> , 2017, 7, 7217.	3.3	48
46	Improving intermolecular interactions in DFTB3 using extended polarization from chemical-potential equalization. <i>Journal of Chemical Physics</i> , 2015, 143, 084123.	3.0	47
47	Electronic Coupling Calculations for Bridge-Mediated Charge Transfer Using Constrained Density Functional Theory (CDFT) and Effective Hamiltonian Approaches at the Density Functional Theory (DFT) and Fragment-Orbital Density Functional Tight Binding (FODFTB) Level. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 4793-4805.	5.3	46
48	Quantum Chemical Molecular Dynamics Model Study of Fullerene Formation from Open-Ended Carbon Nanotubes. <i>Journal of Physical Chemistry A</i> , 2004, 108, 3182-3194.	2.5	43
49	Extended Polarization in Third-Order SCC-DFTB from Chemical-Potential Equalization. <i>Journal of Physical Chemistry A</i> , 2012, 116, 9131-9141.	2.5	42
50	Quantum Effects in Cation Interactions with First and Second Coordination Shell Ligands in Metalloproteins. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 4992-5001.	5.3	42
51	QM/QM Approach to Model Energy Disorder in Amorphous Organic Semiconductors. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 560-567.	5.3	40
52	Active site structure and absorption spectrum of channelrhodopsin-2 wild-type and C128T mutant. <i>Chemical Science</i> , 2016, 7, 3879-3891.	7.4	40
53	Benchmarking density functional tight binding models for barrier heights and reaction energetics of organic molecules. <i>Journal of Computational Chemistry</i> , 2017, 38, 2171-2185.	3.3	39
54	Benchmark and performance of long-range corrected time-dependent density functional tight binding (LC-TD-DFTB) on rhodopsins and light-harvesting complexes. <i>Physical Chemistry Chemical Physics</i> , 2020, 22, 10500-10518.	2.8	36

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55	DFTB/MM Molecular Dynamics Simulations of the FMO Light-Harvesting Complex. <i>Journal of Physical Chemistry Letters</i> , 2020, 11, 8660-8667.	4.6	34
56	A Modified QM/MM Hamiltonian with the Self-Consistent-Charge Density-Functional-Tight-Binding Theory for Highly Charged QM Regions. <i>Journal of Chemical Theory and Computation</i> , 2012, 8, 4293-4304.	5.3	30
57	DFTB3 Parametrization for Copper: The Importance of Orbital Angular Momentum Dependence of Hubbard Parameters. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 4205-4219.	5.3	30
58	Charge and Exciton Transfer Simulations Using Machine-Learned Hamiltonians. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 4061-4070.	5.3	30
59	Color Tuning in Binding Pocket Models of the Chlamydomonas-Type Channelrhodopsins. <i>Journal of Physical Chemistry B</i> , 2011, 115, 15119-15128.	2.6	28
60	Simulation of Temperature-Dependent Charge Transport in Organic Semiconductors with Various Degrees of Disorder. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 3087-3096.	5.3	27
61	Modeling charge transport in DNA using multi-scale methods. <i>Physica Status Solidi (B): Basic Research</i> , 2013, 250, 2277-2287.	1.5	26
62	Multiscale QM/MM molecular dynamics simulations of the trimeric major light-harvesting complex II. <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 7407-7417.	2.8	24
63	Simulation of Singlet Exciton Diffusion in Bulk Organic Materials. <i>Journal of Chemical Theory and Computation</i> , 2016, 12, 4209-4221.	5.3	22
64	On the mechanism of spontaneous thiol-disulfide exchange in proteins. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 16222-16230.	2.8	22
65	Performance of Mixed Quantum-Classical Approaches on Modeling the Crossover from Hopping to Bandlike Charge Transport in Organic Semiconductors. <i>Journal of Chemical Theory and Computation</i> , 2020, 16, 2071-2084.	5.3	21
66	Interplay of structural dynamics and electronic effects in an engineered assembly of pentacene in a metal-organic framework. <i>Chemical Science</i> , 2021, 12, 4477-4483.	7.4	18
67	Accurate Free Energies for Complex Condensed-Phase Reactions Using an Artificial Neural Network Corrected DFTB/MM Methodology. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 1213-1226.	5.3	18
68	Molecular Insights into Variable Electron Transfer in Amphibian Cryptochrome. <i>Biophysical Journal</i> , 2018, 114, 2563-2572.	0.5	17
69	Improved Electronic Properties from Third-Order SCC-DFTB with Cost Efficient Post-SCF Extensions. <i>Journal of Physical Chemistry A</i> , 2012, 116, 11927-11937.	2.5	16
70	Polaron Effects on Charge Transport through Molecular Wires: A Multiscale Approach. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 286-296.	5.3	16
71	Mechanism of a proton pump analyzed with computer simulations. <i>Theoretical Chemistry Accounts</i> , 2010, 125, 353-363.	1.4	15
72	Coupled-perturbed DFTB-QM/MM metadynamics: Application to proton-coupled electron transfer. <i>Journal of Chemical Physics</i> , 2018, 149, 072328.	3.0	14

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73	HAB79: A New Molecular Dataset for Benchmarking DFT and DFTB Electronic Couplings Against High-Level Ab-initio Calculations. <i>Journal of Chemical Physics</i> , 2021, 155, 234115.	3.0	14
74	Exploring the applicability of density functional tight binding to transition metal ions. Parameterization for nickel with the spin-polarized DFTB3 model. <i>Journal of Computational Chemistry</i> , 2019, 40, 400-413.	3.3	13
75	Mechanism by which Untwisting of Retinal Leads to Productive Bacteriorhodopsin Photocycle States. <i>Journal of Physical Chemistry B</i> , 2015, 119, 2229-2240.	2.6	12
76	Best of Two Worlds? How MD Simulations of Amphiphilic Helical Peptides in Membranes Can Complement Data from Oriented Solid-State NMR. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 6002-6014.	5.3	12
77	Computational Study of Synthetic Agonist Ligands of Ionotropic Glutamate Receptors. <i>PLoS ONE</i> , 2013, 8, e58774.	2.5	11
78	Relation between Dephasing Time and Energy Gap Fluctuations in Biomolecular Systems. <i>Journal of Physical Chemistry Letters</i> , 2016, 7, 1102-1108.	4.6	10
79	Analytical Time-Dependent Long-Range Corrected Density Functional Tight Binding (TD-LC-DFTB) Gradients in DFTB+: Implementation and Benchmark for Excited-State Geometries and Transition Energies. <i>Journal of Chemical Theory and Computation</i> , 2021, 17, 2266-2282.	5.3	10
80	Molecular Dynamics Investigation of gluazo, a Photo-Switchable Ligand for the Glutamate Receptor GluK2. <i>PLoS ONE</i> , 2015, 10, e0135399.	2.5	8
81	Ligand Photo-Isomerization Triggers Conformational Changes in iGluR2 Ligand Binding Domain. <i>PLoS ONE</i> , 2014, 9, e92716.	2.5	8
82	Electrostatic interactions contribute to the control of intramolecular thiol-disulfide isomerization in a protein. <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 26366-26375.	2.8	6
83	Geometry dependence of excitonic couplings and the consequences for configuration-space sampling. <i>Journal of Computational Chemistry</i> , 2021, 42, 1402-1418.	3.3	5
84	O to bR transition in bacteriorhodopsin occurs through a proton hole mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	5
85	siRNA traffic lights: arabino-configured 2'-anchors for fluorescent dyes are key for dual color readout in cell imaging. <i>Organic and Biomolecular Chemistry</i> , 2018, 16, 3726-3731.	2.8	4
86	Efficient Surface Hopping Approach for Modeling Charge Transport in Organic Semiconductors. <i>Journal of Chemical Theory and Computation</i> , 2022, 18, 1264-1274.	5.3	4
87	Origin of the Solvatochromism in Organic Fluorophores with Flexible Side Chains: A Case Study of Flugi-2. <i>Journal of Physical Chemistry A</i> , 2019, 123, 4581-4587.	2.5	3
88	Improvement of d-d interactions in density functional tight binding for transition metal ions with a ligand field model: assessment of a DFTB3+U model on nickel coordination compounds. <i>Physical Chemistry Chemical Physics</i> , 2020, 22, 27084-27095.	2.8	3
89	Unravelling the mechanism of glucose binding in a protein-based fluorescence probe: molecular dynamics simulation with a tailor-made charge model. <i>Physical Chemistry Chemical Physics</i> , 2022, 24, 2441-2453.	2.8	2
90	Understanding excited state properties of host materials in OLEDs: simulation of absorption spectrum of amorphous 4,4-bis(carbazol-9-yl)-2,2-biphenyl (CBP). <i>Physical Chemistry Chemical Physics</i> , 2022, , .	2.8	2

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91	Hydrogen Storage in Single-Walled and Multi-Walled Carbon Nanotubes. Materials Research Society Symposia Proceedings, 1999, 593, 187.	0.1	1