

Yuji Mochizuki

List of Publications by Year in descending order

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179
papers

4,525
citations

94415

37
h-index

138468

58
g-index

185
all docs

185
docs citations

185
times ranked

1959
citing authors

#	ARTICLE	IF	CITATIONS
1	Electron-correlated fragment-molecular-orbital calculations for biomolecular and nano systems. <i>Physical Chemistry Chemical Physics</i> , 2014, 16, 10310-10344.	2.8	251
2	Large scale MP2 calculations with fragment molecular orbital scheme. <i>Chemical Physics Letters</i> , 2004, 396, 473-479.	2.6	186
3	A parallelized integral-direct second-order Møller-Plesset perturbation theory method with a fragment molecular orbital scheme. <i>Theoretical Chemistry Accounts</i> , 2004, 112, 442-452.	1.4	160
4	VISCANA: Visualized Cluster Analysis of Protein-Ligand Interaction Based on the ab Initio Fragment Molecular Orbital Method for Virtual Ligand Screening. <i>Journal of Chemical Information and Modeling</i> , 2006, 46, 221-230.	5.4	127
5	Molecular Interactions between Estrogen Receptor and Its Ligand Studied by the ab Initio Fragment Molecular Orbital Method. <i>Journal of Physical Chemistry B</i> , 2006, 110, 16102-16110.	2.6	118
6	Large scale FMO-MP2 calculations on a massively parallel-vector computer. <i>Chemical Physics Letters</i> , 2008, 457, 396-403.	2.6	113
7	Configuration interaction singles method with multilayer fragment molecular orbital scheme. <i>Chemical Physics Letters</i> , 2005, 406, 283-288.	2.6	112
8	Intra- and intermolecular interactions between cyclic-AMP receptor protein and DNA: Ab initio fragment molecular orbital study. <i>Journal of Computational Chemistry</i> , 2006, 27, 948-960.	3.3	107
9	A configuration analysis for fragment interaction. <i>Chemical Physics Letters</i> , 2005, 410, 247-253.	2.6	87
10	Fragment Molecular Orbital Based Interaction Analyses on COVID-19 Main Protease Inhibitor N3 Complex (PDB ID: 6LU7). <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 3593-3602.	5.4	84
11	Higher-order correlated calculations based on fragment molecular orbital scheme. <i>Theoretical Chemistry Accounts</i> , 2011, 130, 515-530.	1.4	78
12	Parallelized integral-direct CIS(D) calculations with multilayer fragment molecular orbital scheme. <i>Theoretical Chemistry Accounts</i> , 2007, 117, 541-553.	1.4	71
13	How Does an S_N2 Reaction Take Place in Solution? Full Ab Initio MD Simulations for the Hydrolysis of the Methyl Diazonium Ion. <i>Journal of the American Chemical Society</i> , 2008, 130, 2396-2397.	13.7	71
14	Self-Degradable Lipid-Like Materials Based on Hydrolysis accelerated by the intra-Particle Enrichment of Reactant (HyPER) for Messenger RNA Delivery. <i>Advanced Functional Materials</i> , 2020, 30, 1910575.	14.9	65
15	Fragment interaction analysis based on local MP2. <i>Theoretical Chemistry Accounts</i> , 2007, 118, 937-945.	1.4	62
16	Fragment molecular orbital-based molecular dynamics (FMO-MD), a quantum simulation tool for large molecular systems. <i>Computational and Theoretical Chemistry</i> , 2009, 898, 2-7.	1.5	59
17	Comparative study of dehydrogenation of methanol at Pt(111)/water and Pt(111)/vacuum interfaces. <i>Chemical Physics Letters</i> , 2003, 377, 236-242.	2.6	58
18	Fragment Molecular Orbital method-based Molecular Dynamics (FMO-MD) as a simulator for chemical reactions in explicit solvation. <i>Journal of Computational Chemistry</i> , 2009, 30, 40-50.	3.3	57

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19	Development of the four-body corrected fragment molecular orbital (FMO4) method. <i>Chemical Physics Letters</i> , 2012, 523, 128-133.	2.6	56
20	Fragment molecular orbital calculations on large scale systems containing heavy metal atom. <i>Chemical Physics Letters</i> , 2006, 427, 159-165.	2.6	53
21	Three-body expansion and generalized dynamic fragmentation improve the fragment molecular orbital-based molecular dynamics (FMO-MD). <i>Chemical Physics Letters</i> , 2010, 484, 380-386.	2.6	53
22	A fully quantum mechanical simulation study on the lowest $\hat{\sigma}^*$ state of hydrated formaldehyde. <i>Chemical Physics Letters</i> , 2007, 437, 66-72.	2.6	52
23	Theoretical analysis of binding specificity of influenza viral hemagglutinin to avian and human receptors based on the fragment molecular orbital method. <i>Computational Biology and Chemistry</i> , 2008, 32, 198-211.	2.3	52
24	Acceleration of fragment molecular orbital calculations with Cholesky decomposition approach. <i>Chemical Physics Letters</i> , 2010, 490, 84-89.	2.6	51
25	Determination of valence band splitting parameters in GaN. <i>Journal of Applied Physics</i> , 1998, 83, 4542-4544.	2.5	49
26	Possibility of Mutation Prediction of Influenza Hemagglutinin by Combination of Hemadsorption Experiment and Quantum Chemical Calculation for Antibody Binding. <i>Journal of Physical Chemistry B</i> , 2009, 113, 4991-4994.	2.6	49
27	Three- and four-body corrected fragment molecular orbital calculations with a novel subdividing fragmentation method applicable to structure-based drug design. <i>Journal of Molecular Graphics and Modelling</i> , 2013, 41, 31-42.	2.4	48
28	Visualization analysis of inter-fragment interaction energies of CRP-cAMP-DNA complex based on the fragment molecular orbital method. <i>Biophysical Chemistry</i> , 2007, 130, 1-9.	2.8	47
29	Ab Initio Fragment Molecular Orbital Study of Molecular Interactions in Liganded Retinoid X Receptor: Specification of Residues Associated with Ligand Inducible Information Transmission. <i>Journal of Physical Chemistry B</i> , 2008, 112, 12081-12094.	2.6	47
30	Fragment Molecular Orbital Calculations with Implicit Solvent Based on the Poisson-Boltzmann Equation: II. Protein and Its Ligand-Binding System Studies. <i>Journal of Physical Chemistry B</i> , 2019, 123, 957-973.	2.6	46
31	Ab initio study on the structures of Th(IV) hydrate and its hydrolysis products in aqueous solution. <i>Chemical Physics Letters</i> , 2003, 375, 204-212.	2.6	44
32	Large-scale FMO-MP3 calculations on the surface proteins of influenza virus, hemagglutinin (HA) and neuraminidase (NA). <i>Chemical Physics Letters</i> , 2010, 493, 346-352.	2.6	44
33	Application of the fragment molecular orbital method for determination of atomic charges on polypeptides. II. Towards an improvement of force fields used for classical molecular dynamics simulations. <i>Chemical Physics Letters</i> , 2009, 467, 417-423.	2.6	42
34	Accuracy of fragmentation in ab initio calculations of hydrated sodium cation. <i>Chemical Physics Letters</i> , 2009, 478, 295-300.	2.6	41
35	Fragment Molecular Orbital Calculations on Red Fluorescent Proteins (DsRed and mFruits). <i>Journal of Physical Chemistry B</i> , 2009, 113, 1153-1161.	2.6	40
36	Implementation of Pair Interaction Energy Decomposition Analysis and Its Applications to Protein-Ligand Systems. <i>Journal of Computer Chemistry Japan</i> , 2015, 14, 1-9.	0.1	40

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37	Ab initio quantum-chemical study on emission spectra of bioluminescent luciferases by fragment molecular orbital method. <i>Chemical Physics Letters</i> , 2009, 472, 118-123.	2.6	39
38	Fragment molecular orbital-based molecular dynamics (FMO-MD) method with MP2 gradient. <i>Chemical Physics Letters</i> , 2011, 504, 95-99.	2.6	38
39	Ab Initio Fragment Molecular Orbital Study of Molecular Interactions between Liganded Retinoid X Receptor and Its Coactivator; Part II: Influence of Mutations in Transcriptional Activation Function 2 Activating Domain Core on the Molecular Interactions. <i>Journal of Physical Chemistry A</i> , 2008, 112, 1986-1998.	2.5	37
40	Application of the fragment molecular orbital method for determination of atomic charges on polypeptides. <i>Chemical Physics Letters</i> , 2007, 449, 329-335.	2.6	36
41	Developments and applications of ABINIT-MP software based on the fragment molecular orbital method. , 2006, , 39-52.		36
42	Fragment molecular orbital-based molecular dynamics (FMO-MD) simulations on hydrated Zn(II) ion. <i>Chemical Physics Letters</i> , 2010, 490, 41-45.	2.6	35
43	Fragment Molecular Orbital Calculations with Implicit Solvent Based on the Poisson-Boltzmann Equation: Implementation and DNA Study. <i>Journal of Physical Chemistry B</i> , 2018, 122, 4457-4471.	2.6	35
44	Cm ³⁺ /Eu ³⁺ induced structural, mechanistic and functional implications for calmodulin. <i>Physical Chemistry Chemical Physics</i> , 2019, 21, 21213-21222.	2.8	34
45	Ab Initio Fragment Molecular Orbital Study of Molecular Interactions between Liganded Retinoid X Receptor and Its Coactivator: Roles of Helix 12 in the Coactivator Binding Mechanism. <i>Journal of Physical Chemistry B</i> , 2007, 111, 3525-3533.	2.6	33
46	Fragment molecular orbital calculations on red fluorescent protein (DsRed). <i>Chemical Physics Letters</i> , 2007, 433, 360-367.	2.6	33
47	Reflectance spectroscopy on GaN films under uniaxial stress. <i>Applied Physics Letters</i> , 1997, 71, 374-376.	3.3	31
48	Dynamic polarizability calculation with fragment molecular orbital scheme. <i>Chemical Physics Letters</i> , 2006, 418, 418-422.	2.6	31
49	An application of fragment interaction analysis based on local MP2. <i>Chemical Physics Letters</i> , 2008, 463, 189-194.	2.6	30
50	Modeling of peptide-silica interaction based on four-body corrected fragment molecular orbital (FMO4) calculations. <i>Chemical Physics Letters</i> , 2013, 566, 25-31.	2.6	30
51	Dissipative particle dynamics (DPD) simulations with fragment molecular orbital (FMO) based effective parameters for 1-Palmitoyl-2-oleoyl phosphatidyl choline (POPC) membrane. <i>Chemical Physics Letters</i> , 2017, 684, 427-432.	2.6	29
52	Development of the Fragment Molecular Orbital Method for Calculating Nonlocal Excitations in Large Molecular Systems. <i>Journal of Physical Chemistry A</i> , 2018, 122, 3886-3898.	2.5	28
53	Prolapses in four-component relativistic Gaussian basis sets. <i>Chemical Physics Letters</i> , 2003, 375, 399-405.	2.6	26
54	Theoretical study of hydration models of trivalent rare-earth ions using model core potentials. <i>Computational and Theoretical Chemistry</i> , 2010, 949, 28-35.	1.5	26

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55	Prediction of probable mutations in influenza virus hemagglutinin protein based on large-scale ab initio fragment molecular orbital calculations. <i>Journal of Molecular Graphics and Modelling</i> , 2011, 30, 110-119.	2.4	26
56	Theoretical analyses on water cluster structures in polymer electrolyte membrane by using dissipative particle dynamics simulations with fragment molecular orbital based effective parameters. <i>RSC Advances</i> , 2018, 8, 34582-34595.	3.6	26
57	Four-component relativistic calculations on the complexes between a water molecule and trivalent lanthanoid and actinoid ions. <i>Chemical Physics</i> , 2001, 273, 135-148.	1.9	24
58	Counterpoise-corrected interaction energy analysis based on the fragment molecular orbital scheme. <i>Chemical Physics Letters</i> , 2011, 509, 67-71.	2.6	24
59	A theoretical study of the bent form of CuO ₂ . <i>Chemical Physics Letters</i> , 1989, 164, 225-230.	2.6	23
60	FMO-MD Simulations on the Hydration of Formaldehyde in Water Solution with Constraint Dynamics. <i>Chemistry - A European Journal</i> , 2012, 18, 9714-9721.	3.3	23
61	Vapor-Liquid Equilibrium Data for the Four Binary Systems Containing Fluorocarbon, Hydrofluorocarbon, and Fluorinated Ethers at 101.3 kPa. <i>Journal of Chemical & Engineering Data</i> , 2001, 46, 913-917.	1.9	22
62	Comments on relativistic basis sets. <i>Theoretical Chemistry Accounts</i> , 2003, 109, 40-42.	1.4	22
63	Modification for spin-adapted version of configuration interaction singles with perturbative doubles. <i>Chemical Physics Letters</i> , 2007, 443, 389-397.	2.6	22
64	Electronic structure of the linear form of OCuO. <i>Chemical Physics</i> , 1991, 151, 11-20.	1.9	21
65	Does Amination of Formaldehyde Proceed Through a Zwitterionic Intermediate in Water? Fragment Molecular Orbital Molecular Dynamics Simulations by Using Constraint Dynamics. <i>Chemistry - A European Journal</i> , 2010, 16, 6430-6433.	3.3	21
66	Flexible ligand recognition of peroxisome proliferator-activated receptor- β (PPAR β). <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010, 20, 3344-3347.	2.2	21
67	Differences in hydration between cis- and trans-platin: Quantum insights by ab initio fragment molecular orbital-based molecular dynamics (FMO-MD). <i>Computational and Theoretical Chemistry</i> , 2012, 986, 30-34.	2.5	21
68	Size, Order, and Dimensional Relations for Silicon Cluster Polarizabilities. <i>Journal of Physical Chemistry A</i> , 2002, 106, 395-399.	2.5	20
69	Explicit solvation modulates intra- and inter-molecular interactions within DNA: Electronic aspects revealed by the ab initio fragment molecular orbital (FMO) method. <i>Computational and Theoretical Chemistry</i> , 2015, 1054, 29-37.	2.5	20
70	Modeling of hydroxyapatite-peptide interaction based on fragment molecular orbital method. <i>Chemical Physics Letters</i> , 2015, 629, 58-64.	2.6	20
71	Fragment Molecular Orbital Based Parametrization Procedure for Mesoscopic Structure Prediction of Polymeric Materials. <i>Journal of Physical Chemistry B</i> , 2018, 122, 338-347.	2.6	20
72	Interaction analyses of SARS-CoV-2 spike protein based on fragment molecular orbital calculations. <i>RSC Advances</i> , 2021, 11, 3272-3279.	3.6	20

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73	Theoretical investigation on the GaH molecule and its positive ion. <i>Theoretical Chemistry Accounts</i> , 1998, 99, 88-94.	1.4	18
74	On the electronic structure of $\text{Cm}(\text{H}_2\text{O})_n^{3+}$ ($n=1,2,4,6$) by all-electron Dirac-Hartree-Fock calculations. <i>Journal of Chemical Physics</i> , 2002, 116, 8838-8842.	3.0	18
75	Fragment molecular orbital (FMO) study on stabilization mechanism of neuro-oncological ventral antigen (NOVA)-RNA complex system. <i>Computational and Theoretical Chemistry</i> , 2010, 962, 45-55.	1.5	18
76	Partial geometry optimization with FMO-MP2 gradient: Application to TrpCage. <i>Chemical Physics Letters</i> , 2012, 535, 157-162.	2.6	18
77	A theoretical investigation of sulphur K-shell X-ray absorption of cysteine. <i>Chemical Physics Letters</i> , 1999, 309, 241-248.	2.6	17
78	Interaction energy analysis on specific binding of influenza virus hemagglutinin to avian and human sialosaccharide receptors: Importance of mutation-induced structural change. <i>Journal of Molecular Graphics and Modelling</i> , 2014, 53, 48-58.	2.4	17
79	Folding simulation of small proteins by dissipative particle dynamics (DPD) with non-empirical interaction parameters based on fragment molecular orbital calculations. <i>Applied Physics Express</i> , 2020, 13, 017002.	2.4	17
80	Dynamic Cooperativity of Ligand-Residue Interactions Evaluated with the Fragment Molecular Orbital Method. <i>Journal of Physical Chemistry B</i> , 2021, 125, 6501-6512.	2.6	17
81	Antigen-antibody interactions of influenza virus hemagglutinin revealed by the fragment molecular orbital calculation. <i>Theoretical Chemistry Accounts</i> , 2011, 130, 1197-1202.	1.4	16
82	Fragment molecular orbital calculations for excitation energies of blue- and yellow-fluorescent proteins. <i>Chemical Physics Letters</i> , 2011, 504, 76-82.	2.6	16
83	Accuracy of the fragment molecular orbital (FMO) calculations for DNA: Total energy, molecular orbital, and inter-fragment interaction energy. <i>Computational and Theoretical Chemistry</i> , 2014, 1034, 7-16.	2.5	16
84	Interaction Analysis on the SARS-CoV-2 Spike Protein Receptor Binding Domain Using Visualization of the Interfacial Electrostatic Complementarity. <i>Journal of Physical Chemistry Letters</i> , 2021, 12, 11267-11272.	4.6	16
85	A CSF-based multi-reference coupled pair approximation. <i>Theoretical Chemistry Accounts</i> , 1998, 98, 165-170.	1.4	15
86	Application of fragment molecular orbital scheme to silicon-containing systems. <i>Chemical Physics Letters</i> , 2006, 430, 361-366.	2.6	15
87	Fragment molecular orbital (FMO) calculations on DNA by a scaled third-order Møller-Plesset perturbation (MP2.5) scheme. <i>Computational and Theoretical Chemistry</i> , 2017, 1101, 46-54.	2.5	15
88	Stabilization Mechanism for a Nonfibrillar Amyloid β^2 Oligomer Based on Formation of a Hydrophobic Core Determined by Dissipative Particle Dynamics. <i>ACS Chemical Neuroscience</i> , 2020, 11, 385-394.	3.5	15
89	HF-STEX and RASSCF calculations on nitrogen K-shell X-ray absorption of purine base and its derivative. <i>Journal of Synchrotron Radiation</i> , 2001, 8, 1003-1005.	2.4	14
90	On the electronic structures of Th^{4+} and Ac^{3+} hydrate models. <i>Chemical Physics Letters</i> , 2003, 372, 114-120.	2.6	14

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91	A practical use of self-energy shift for the description of orbital relaxation. <i>Chemical Physics Letters</i> , 2008, 453, 109-116.	2.6	14
92	Theoretical investigation of the GaF molecule and its positive ion. <i>Theoretical Chemistry Accounts</i> , 1999, 101, 257-261.	1.4	13
93	Multi-reference calculations of nitric oxide dimer. <i>Chemical Physics Letters</i> , 2008, 451, 31-36.	2.6	13
94	4f-in-core model core potentials for trivalent lanthanides. <i>Chemical Physics Letters</i> , 2011, 510, 261-266.	2.6	13
95	Development of an Analysis Toolkit, AnalysisFMO, to Visualize Interaction Energies Generated by Fragment Molecular Orbital Calculations. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 25-30.	5.4	13
96	A theoretical study of the CuOH molecule. <i>Chemical Physics Letters</i> , 1991, 185, 535-543.	2.6	12
97	Theoretical study of the Cl desorption reaction induced by H ₂ in the chloride atomic layer epitaxy. <i>Journal of Crystal Growth</i> , 1994, 135, 259-268.	1.5	12
98	Multireference coupled-pair approximation study of the CuSi molecule. <i>Theoretical Chemistry Accounts</i> , 1999, 101, 332-335.	1.4	12
99	A size-extensive modification of super-CI for orbital relaxation. <i>Chemical Physics Letters</i> , 2005, 410, 165-171.	2.6	12
100	Hydration of ligands of influenza virus neuraminidase studied by the fragment molecular orbital method. <i>Journal of Molecular Graphics and Modelling</i> , 2016, 69, 144-153.	2.4	12
101	Destabilization of DNA through interstrand crosslinking by UO ₂ ²⁺ . <i>Chemical Communications</i> , 2019, 55, 2015-2018.	4.1	12
102	Statistical interaction analyses between SARS-CoV-2 main protease and inhibitor N3 by combining molecular dynamics simulation and fragment molecular orbital calculation. <i>Applied Physics Express</i> , 2021, 14, 027003.	2.4	12
103	Theoretical study of As ₂ desorption from the Ga dangling-bond site. <i>Physical Review B</i> , 1994, 49, 4658-4667.	3.2	11
104	On the Reaction Scheme for Ti/TiN Chemical Vapor Deposition (CVD) Process Using TiCl ₄ . <i>Japanese Journal of Applied Physics</i> , 1995, 34, L326-L329.	1.5	11
105	Polarizability of silicon clusters. <i>Chemical Physics Letters</i> , 2001, 336, 451-456.	2.6	11
106	Recent Advances in Fragment Molecular Orbital-Based Molecular Dynamics (FMO-MD) Simulations. , 0, ..		11
107	Variational quantum eigensolver simulations with the multireference unitary coupled cluster ansatz: a case study of the C ₂ v ⁻ quasi-reaction pathway of beryllium insertion into a H ₂ molecule. <i>Physical Chemistry Chemical Physics</i> , 2022, 24, 8439-8452.	2.8	11
108	Ab initio cas SCF/MRSDCI study of the CuCH ₂ cluster. <i>Chemical Physics Letters</i> , 1988, 152, 457-463.	2.6	10

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109	New Concerted Mechanism of the Cl-Removal Reaction Induced by H ₂ in Chloride Atomic Layer Epitaxy. Japanese Journal of Applied Physics, 1993, 32, L197-L199.	1.5	10
110	Theoretical spectroscopic constants of the GaN molecule. Theoretical Chemistry Accounts, 1999, 101, 292-296.	1.4	10
111	Ab initio path integral Monte Carlo simulations for water trimer with electron correlation effects. Computational and Theoretical Chemistry, 2012, 997, 7-13.	2.5	10
112	The ABINIT-MP Program. , 2021, , 53-67.		10
113	Theoretical study of the gallium chloride molecule and its interaction with arsenic dangling bonds. Physical Review B, 1993, 47, 13420-13431.	3.2	9
114	Computational study of the spectroscopic constants of the ground state of the As ₂ molecule. Chemical Physics Letters, 1997, 274, 264-268.	2.6	9
115	An automated framework to evaluate effective interaction parameters for dissipative particle dynamics simulations based on the fragment molecular orbital (FMO) method. Journal of Computer Chemistry Japan, 2018, 17, 102-109.	0.1	9
116	Electronic structure of CO adsorbed on small Cu clusters: Theoretical study on excited states. Physical Review B, 1989, 39, 11907-11913.	3.2	8
117	Electronic structure of lower singlet states of binuclear copper acetate monohydrate. Canadian Journal of Chemistry, 1992, 70, 393-398.	1.1	8
118	Theoretical study on the direct reaction between AsH ₃ and surface-adsorbed GaCl. Journal of Crystal Growth, 1995, 148, 96-105.	1.5	8
119	Theoretical study of hydrolysis reactions of tetravalent thorium ion. Chemical Physics Letters, 2003, 373, 213-217.	2.6	8
120	On the electronic structure of CmFn (n=1~4) by all-electron Dirac-Hartree-Fock calculations. Journal of Chemical Physics, 2003, 118, 9201-9207.	3.0	8
121	A graphical symmetric group approach for a spin adapted full configuration interaction: partitioning of a configuration graph into sets of closed-shell and open-shell graphs. Theoretical Chemistry Accounts, 2007, 117, 397-405.	1.4	8
122	Application of Dyson-corrected second-order perturbation theories. Chemical Physics Letters, 2009, 472, 143-148.	2.6	8
123	Explicit solvation of a single-stranded DNA, a binding protein, and their complex: a suitable protocol for a fragment molecular orbital calculation. Chem-Bio Informatics Journal, 2017, 17, 72-84.	0.3	8
124	Current Status of ABINIT-MP as a FMO Program and Related Works with Machine Learning. Journal of Computer Chemistry Japan, 2017, 16, 119-122.	0.1	8
125	A New Treatment for Water near the Interface between Lipid Membrane and Silica in Dissipative Particle Dynamics Simulation. Journal of Computer Chemistry Japan, 2017, 16, 28-31.	0.1	8
126	Fragment molecular orbital based interaction analyses on complexes between SARS-CoV-2 RBD variants and ACE2. Japanese Journal of Applied Physics, 2021, 60, 090901.	1.5	8

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127	Theoretical studies on the chloride ALE process. Applied Surface Science, 1994, 82-83, 200-207.	6.1	7
128	Modification of nonrelativistic Gaussian basis sets for relativistic calculations. Journal of Chemical Physics, 2001, 115, 9160-9164.	3.0	7
129	Interaction between a Single-Stranded DNA and a Binding Protein Viewed by the Fragment Molecular Orbital Method. Bulletin of the Chemical Society of Japan, 2018, 91, 1596-1605.	3.2	7
130	Collective residue interactions in trimer complexes of SARS-CoV-2 spike proteins analyzed by fragment molecular orbital method. Applied Physics Express, 2022, 15, 017001.	2.4	7
131	Theoretical Study of the Surface Reaction Mechanism of GaN with HCl. Japanese Journal of Applied Physics, 1996, 35, L1641-L1643.	1.5	6
132	Excited state calculation for free-base and metalloporphyrins with the partially renormalized polarization propagator approach. Chemical Physics Letters, 2012, 525-526, 144-149.	2.6	6
133	Hydration effects on enzyme-substrate complex of nylon oligomer hydrolase: inter-fragment interaction energy study by the fragment molecular orbital method. Molecular Physics, 2015, 113, 319-326.	1.7	6
134	Formation Mechanism of Lipid Membrane and Vesicle Using Small Angle X-ray Scattering and Dissipative Particle Dynamics (DPD) Method. Journal of Computer Chemistry Japan, 2018, 17, 172-179.	0.1	6
135	RI-MP3 calculations of biomolecules based on the fragment molecular orbital method. Journal of Computational Chemistry, 2018, 39, 1970-1978.	3.3	6
136	Application of TensorFlow to recognition of visualized results of fragment molecular orbital (FMO) calculations. Chem-Bio Informatics Journal, 2018, 18, 58-69.	0.3	6
137	Interaction between calcite and adsorptive peptide analyzed by fragment molecular orbital method. Japanese Journal of Applied Physics, 2019, 58, 120906.	1.5	6
138	Developments of FMO Methodology and Graphical User Interface in ABINIT-MP. , 2009, , 37-62.		6
139	Deeper Level Parallelization and Performance Evaluation of FMO Program ABINIT-MP on Oakforest-PACS. Journal of Computer Chemistry Japan, 2018, 17, 147-149.	0.1	6
140	Fragment molecular orbital-based molecular dynamics (FMO-MD) simulations on hydrated Cu(II) ion. Chem-Bio Informatics Journal, 2014, 14, 1-13.	0.3	5
141	Current Status of ABINIT-MP Open Series. Journal of Computer Chemistry Japan, 2019, 18, 129-131.	0.1	5
142	Generalized doubly symbolic formulation for integral-driven direct configuration interaction method. Theoretica Chimica Acta, 1996, 93, 211-233.	0.8	4
143	MULTI-REFERENCE COUPLED PAIR APPROXIMATION: a state-universal approach of a CEPA type variant of MRSDCI. Recent Advances in Computational, 1999, , 95-130.	0.8	4
144	Vectorization of Direct Fock Matrix Construction in DIRAC-DHF Calculations. Journal of Nuclear Science and Technology, 2002, 39, 195-199.	1.3	4

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145	Fragmentation at sp ² carbon atoms in fragment molecular orbital method. Journal of Computational Chemistry, 2020, 41, 1416-1420.	3.3	4
146	Four-component relativistic calculations on the mono-ammine complexes of trivalent f0, f7, and f14 ions. Chemical Physics Letters, 2002, 359, 331-336.	2.6	3
147	Intra- and intermolecular interactions between cyclic-AMP receptor protein and DNA: Abinitio fragment molecular orbital study. Journal of Computational Chemistry, 2007, 28, 2237-2239.	3.3	3
148	Application of the FMO Method to Specific Molecular Recognition of Biomacromolecules. , 2009, , 133-170.		3
149	Fragment Molecular Orbital (FMO) Calculations of Peptoids. Journal of Computer Chemistry Japan, 2016, 15, 51-52.	0.1	3
150	Development Status of ABINIT-MP in 2020. Journal of Computer Chemistry Japan, 2020, 19, 142-145.	0.1	3
151	ON MECHANISM OF ENHANCED FLUORESCENCE IN GREEN FLUORESCENT PROTEIN. Biophysical Reviews and Letters, 2007, 02, 221-227.	0.8	2
152	Development and Performance Evaluation of a Simulation Code for Dissipative Particle Dynamics (DPD) CAMUS. Journal of Computer Chemistry Japan, 2017, 16, 126-128.	0.1	2
153	Effects of Point Mutations on the Binding Energies of Estrogen Receptor with Estradiol. Journal of Computer Chemistry Japan, 2007, 6, 33-46.	0.1	2
154	Development Status of ABINIT-MP in 2021. Journal of Computer Chemistry Japan, 2021, 20, 132-136.	0.1	2
155	Ab initio MO studies on the hydration of trivalent curium ion. Journal of Nuclear Science and Technology, 2002, 39, 418-421.	1.3	1
156	Application of Fragment Molecular Orbital (FMO) Method to Nano-Bio Field. Journal of Computer Chemistry Japan, 2007, 6, 173-184.	0.1	1
157	Performance Evaluations of Parallelized DFT Calculations with SMASH on Intel Xeon Phi Processor. Journal of Computer Chemistry Japan, 2016, 15, 92-96.	0.1	1
158	<i>Ab Initio</i> Fragment Molecular Orbital-Based Molecular Dynamics (FMO-MD) Simulations of (NH ₃) ₃₂ Cluster: Effects of Electron Correlation. Bulletin of the Chemical Society of Japan, 2020, 93, 553-560.	3.2	1
159	Acceleration of Environmental Electrostatic Potential Using Cholesky Decomposition with Adaptive Metric (CDAM) for Fragment Molecular Orbital (FMO) Method. Bulletin of the Chemical Society of Japan, 2021, 94, 91-96.	3.2	1
160	Dynamic fragmentation with static fragments (DF/SF) algorithm designed for <i>ab initio</i> fragment molecular orbital-based molecular dynamics (FMO-MD) simulations of polypeptides. Chem-Bio Informatics Journal, 2013, 13, 45-57.	0.3	1
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