

András Farkas

List of Publications by Year in descending order

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45
papers

1,922
citations

304743

22
h-index

265206

42
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46
all docs

46
docs citations

46
times ranked

1865
citing authors

#	ARTICLE	IF	CITATIONS
1	Geometry optimization with QM/MM, ONIOM, and other combined methods. I. Microiterations and constraints. <i>Journal of Computational Chemistry</i> , 2003, 24, 760-769.	3.3	560
2	Methods for optimizing large molecules. II. Quadratic search. <i>Journal of Chemical Physics</i> , 1999, 111, 10806-10814.	3.0	169
3	Effects of the Protein Environment on the Structure and Energetics of Active Sites of Metalloenzymes. ONIOM Study of Methane Monooxygenase and Ribonucleotide Reductase. <i>Journal of the American Chemical Society</i> , 2002, 124, 192-193.	13.7	124
4	Methods for optimizing large molecules. Part III. An improved algorithm for geometry optimization using direct inversion in the iterative subspace (GDIIIS). <i>Physical Chemistry Chemical Physics</i> , 2002, 4, 11-15.	2.8	114
5	Methods for geometry optimization of large molecules. I. An O(N ²) algorithm for solving systems of linear equations for the transformation of coordinates and forces. <i>Journal of Chemical Physics</i> , 1998, 109, 7100-7104.	3.0	65
6	Toward a computed peptide structure database: The role of a universal atomic numbering system of amino acids in peptides and internal hierarchy of database. <i>International Journal of Quantum Chemistry</i> , 2002, 90, 933-968.	2.0	54
7	Peptide models. XXXIII. Extrapolation of low-level Hartree-Fock data of peptide conformation to large basis set SCF, MP2, DFT, and CCSD(T) results. The Ramachandran surface of alanine dipeptide computed at various levels of theory. <i>Journal of Computational Chemistry</i> , 2003, 24, 1026-1042.	3.3	54
8	Peptide Models. 18. Hydroxymethyl Side-Chain Induced Backbone Conformational Shifts of L-Serine Amide. All ab Initio Conformers of For-L-Ser-NH ₂ . <i>Journal of the American Chemical Society</i> , 1996, 118, 7809-7817.	13.7	49
9	Peptide models XXIV: An ab initio study on N-formyl-L-prolinamide with trans peptide bond. The existence or non-existence of ¹ L and ¹ uL conformations. <i>Computational and Theoretical Chemistry</i> , 1999, 465, 79-91.	1.5	48
10	Phenylalanine Ammonia-Lyase-Catalyzed Deamination of an Acyclic Amino Acid: Enzyme Mechanistic Studies Aided by a Novel Microreactor Filled with Magnetic Nanoparticles. <i>ChemBioChem</i> , 2015, 16, 2283-2288.	2.6	46
11	Peptide models XXII. A conformational model for aromatic amino acid residues in proteins. A comprehensive analysis of all the RHF/6-31+G* conformers of For-L-Phe-NH ₂ . <i>Computational and Theoretical Chemistry</i> , 1998, 455, 303-314.	1.5	44
12	Peptide models XX. Aromatic side-chain-backbone interaction in phenylalanine-containing diamide model system. A systematic search for the identification of all the ab initio conformers of N-formyl-L-phenylalanine-amide. <i>Canadian Journal of Chemistry</i> , 1997, 75, 1120-1130.	1.1	43
13	Peptide models XIX: Side-chain conformational energy surface and amide I vibrational frequencies of N-formyl-L-phenylalaninamide (For-Phe-NH ₂) in its ¹ L or ¹ inv or C7eq backbone conformation. <i>Computational and Theoretical Chemistry</i> , 1996, 369, 105-114.	1.5	41
14	Peptide models XVI. The identification of selected HCO-L-SER-NH ₂ conformers via a systematic grid search using ab initio potential energy surfaces. <i>Journal of Computational Chemistry</i> , 1996, 17, 821-834.	3.3	37
15	Peptide models XXIII. Conformational model for polar side-chain containing amino acid residues: A comprehensive analysis of RHF, DFT, and MP2 properties of HCO-L-SER-NH ₂ . <i>Journal of Computational Chemistry</i> , 2000, 21, 626-655.	3.3	37
16	Mechanism of the Tyrosine Ammonia Lyase Reaction-Tandem Nucleophilic and Electrophilic Enhancement by a Proton Transfer. <i>Chemistry - A European Journal</i> , 2012, 18, 7793-7802.	3.3	37
17	Mechanism for the Oxidation of Sulfides and Sulfoxides with Periodates: Reactivity of the Oxidizing Species. <i>European Journal of Organic Chemistry</i> , 2009, 2009, 2102-2111.	2.4	35
18	Characteristics of Ramachandran maps of L-alanine diamides as computed by various molecular mechanics, semiempirical and ab initio MO methods.. <i>Computational and Theoretical Chemistry</i> , 1998, 455, 275-301.	1.5	34

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19	Conformations of ethylbenzene (CH ₃ —CH ₂ —Ph). An ab initio study. Computational and Theoretical Chemistry, 1996, 367, 25-31.	1.5	32
20	Mechanism of nucleophilic substitutions at phenacyl bromides with pyridines. A computational study of intermediate and transition state. Journal of Physical Organic Chemistry, 2008, 21, 988-996.	1.9	28
21	Concerted S _N ² mechanism for the hydrolysis of acid chlorides: comparisons of reactivities calculated by the density functional theory with experimental data. Journal of Physical Organic Chemistry, 2011, 24, 480-491.	1.9	26
22	A DFT study of transition structures and reactivity in solvolyses of tert-butyl chloride, cumyl chlorides, and benzyl chlorides. Journal of Physical Organic Chemistry, 2008, 21, 53-61.	1.9	25
23	Molecular modelling of xylose isomerase catalysis: the role of electrostatics and charge transfer to metals. Protein Engineering, Design and Selection, 1995, 8, 925-933.	2.1	21
24	Effect of Substituents on Activation Parameters in Aliphatic SN ₂ Reactions. A DFT Study. Journal of Organic Chemistry, 2006, 71, 3409-3416.	3.2	21
25	Peptide models. XIV. Ab initio study on the role of side-chain backbone interaction stabilizing the building unit of right- and left-handed helices in peptides and proteins. International Journal of Quantum Chemistry, 1997, 61, 797-814.	2.0	20
26	Peptide models XXI. Side-chain/backbone conformational interconversions in HCO-I-Ser-NH ₂ . Tracing relaxation paths by ab initio modeling. An exploratory study. Computational and Theoretical Chemistry, 1998, 455, 315-338.	1.5	16
27	Computational Study of Reactivity and Transition Structures in Nucleophilic Substitutions on Benzyl Bromides. European Journal of Organic Chemistry, 2006, 2006, 5570-5580.	2.4	16
28	Peptide models XII Topological features of molecular mechanics and ab-initio 8D-Ramachandran maps. Conformational data for Ac-(l-Ala) ₄ -NHMe and For-(l-Ala) ₄ -NH ₂ . Computational and Theoretical Chemistry, 1995, 331, 11-26.	1.5	15
29	Peptide models VII The ending of the right-handed helices in oligopeptides [For-(Ala) _n -NH ₂ for 2 ≤ n ≤ 4] and in proteins. Computational and Theoretical Chemistry, 1995, 331, 5-10.	1.5	14
30	Conformational effects of one glycine residue on the other glycine residues in the Ac-Gly-Gly-Gly-NHMe tripeptide motif: an ab initio exploratory study. Computational and Theoretical Chemistry, 2002, 588, 187-200.	1.5	14
31	Exploration of the Four-Dimensional-Conformational Potential Energy Hypersurface of N-Acetyl-l-aspartic Acid N-Methylamide with Its Internally Hydrogen Bonded Side-Chain Orientation. Journal of Physical Chemistry A, 2002, 106, 6999-7009.	2.5	13
32	Generation and analysis of the conformational potential energy surfaces of N-acetyl-N-methyl-l-alanine-N-methylamide. An exploratory ab initio study. Computational and Theoretical Chemistry, 2003, 625, 121-136.	1.5	11
33	Mechanism of the Gibbs Reaction. 3. Indophenol Formation via Radical Electrophilic Aromatic Substitution (SREAr) on Phenols. Journal of Organic Chemistry, 1994, 59, 6543-6557.	3.2	10
34	Reverse turn conformation of N-thioacetyl thioprolyl glycine N-methylamide in the crystal and in solution. Tetrahedron, 1993, 49, 6661-6668.	1.9	9
35	Geometry optimization of Kringle 1 of plasminogen using the PM3 semiempirical method. International Journal of Quantum Chemistry, 2000, 77, 82-89.	2.0	9
36	Mechanism for the reactions of sulfides and sulfoxides with hypochlorites: racemization and oxygen exchange of oxysulfonium salts and sulfoxides. Journal of Physical Organic Chemistry, 2012, 25, 1086-1096.	1.9	9

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37	Antibody Recognition and Conformational Flexibility of a Plaque-Specific β -Amyloid Epitope Modulated by Non-native Peptide Flanking Regions. <i>Journal of Medicinal Chemistry</i> , 2008, 51, 1150-1161.	6.4	6
38	Quantum chemical (QM:MM) investigation of the mechanism of enzymatic reaction of tryptamine and N,N-dimethyltryptamine with monoamine oxidase A. <i>Organic and Biomolecular Chemistry</i> , 2020, 18, 9660-9674.	2.8	6
39	Mechanism for the reactions of sulfides with hypochlorous acid and N-chlorosulfonamides: Formation of solvated chlorosulfonium cation and λ 4-sulfane intermediates. <i>Journal of Physical Organic Chemistry</i> , 2019, 32, e4005.	1.9	4
40	Parameter-free linear relationship (PFLR) and its application to 3D QSAR. <i>Journal of Mathematical Chemistry</i> , 2009, 45, 598-606.	1.5	2
41	Helix compactness and stability: Electron structure calculations of conformer dependent thermodynamic functions. <i>Chemical Physics Letters</i> , 2013, 563, 80-87.	2.6	2
42	An exploratory study on the oxo-enol tautomerization of selected dioxopiperazines and their sulphur-containing analogues. <i>Computational and Theoretical Chemistry</i> , 1993, 286, 131-148.	1.5	1
43	Peptide models. XIV. Ab initio study on the role of side-chain backbone interaction stabilizing the building unit of right- and left-handed helices in peptides and proteins. , 1997, 61, 797.		1
44	Synthesis, structure and conformational analysis of imidazo-thiazines. <i>Journal of Molecular Structure</i> , 1996, 377, 277-288.	3.6	0
45	Quantum chemical study of the hydrolysis of oxidized endogenous psychedelic N,N-dimethyltryptamine. <i>Computational and Theoretical Chemistry</i> , 2022, , 113789.	2.5	0