Klaus R. Liedl

List of Publications by Year in descending order

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Version: 2024-02-01

327 papers 9,587 citations

³⁸⁷⁴² 50 h-index

74163 75 g-index

356 all docs

 $\begin{array}{c} 356 \\ \\ \text{docs citations} \end{array}$

356 times ranked

10708 citing authors

#	Article	IF	CITATIONS
1	Paratope states in solution improve structure prediction and docking. Structure, 2022, 30, 430-440.e3.	3.3	8
2	Explicit solvation thermodynamics in ionic solution: extending grid inhomogeneous solvation theory to solvation free energy of salt–water mixtures. Journal of Computer-Aided Molecular Design, 2022, 36, 101-116.	2.9	8
3	Impact of different protonation states on virtual screening performance against cruzain. Chemical Biology and Drug Design, 2022, 99, 703-716.	3.2	1
4	CDR loop interactions can determine heavy and light chain pairing preferences in bispecific antibodies. MAbs, 2022, 14, 2024118.	5.2	4
5	Comparing Antibody Interfaces to Inform Rational Design of New Antibody Formats. Frontiers in Molecular Biosciences, 2022, 9, 812750.	3.5	4
6	Broadly neutralizing antibodies target a haemagglutinin anchor epitope. Nature, 2022, 602, 314-320.	27.8	78
7	Calcium current modulation by the \hat{I}^31 subunit depends on alternative splicing of CaV1.1. Journal of General Physiology, 2022, 154, .	1.9	4
8	Essential role of a conserved aspartate for the enzymatic activity of plasmanylethanolamine desaturase. Cellular and Molecular Life Sciences, 2022, 79, 214.	5.4	2
9	Grid inhomogeneous solvation theory for cross-solvation in rigid solvents. Journal of Chemical Physics, 2022, 156, .	3.0	3
10	Nanobody Paratope Ensembles in Solution Characterized by MD Simulations and NMR. International Journal of Molecular Sciences, 2022, 23, 5419.	4.1	6
11	Increase of Radiative Forcing through Midinfrared Absorption by Stable CO ₂ Dimers?. Journal of Physical Chemistry A, 2022, 126, 2966-2975.	2.5	3
12	Borylated Cymantrenes and Tromancenium Salts with Unusual Reactivity. Organometallics, 2022, 41, 1464-1473.	2.3	2
13	Hydration thermodynamics of cytosolic phospholipase A2 GIVA predict its membrane-associated parts and its highly hydrated binding site. Journal of Biomolecular Structure and Dynamics, 2021, 39, 953-959.	3.5	1
14	Conformational flexibility correlates with glucose tolerance for point mutations in β-glucosidases – a computational study. Journal of Biomolecular Structure and Dynamics, 2021, 39, 1621-1634.	3.5	10
15	Conformational Ensembles of Antibodies Determine Their Hydrophobicity. Biophysical Journal, 2021, 120, 143-157.	0.5	23
16	Ensembles in solution as a new paradigm for antibody structure prediction and design. MAbs, 2021, 13, 1923122.	5.2	19
17	Inverse relation between structural flexibility and IgE reactivity of Cor a 1 hazelnut allergens. Scientific Reports, 2021, 11, 4173.	3.3	14
18	Structural determinants of voltage-gating properties in calcium channels. ELife, 2021, 10, .	6.0	18

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19	<i>CACNA11</i> gain-of-function mutations differentially affect channel gating and cause neurodevelopmental disorders. Brain, 2021, 144, 2092-2106.	7.6	26
20	Conformational Shifts of Stacked Heteroaromatics: Vacuum vs. Water Studied by Machine Learning. Frontiers in Chemistry, 2021, 9, 641610.	3.6	1
21	Mutation of Framework Residue H71 Results in Different Antibody Paratope States in Solution. Frontiers in Immunology, 2021, 12, 630034.	4.8	17
22	X-Entropy: A Parallelized Kernel Density Estimator with Automated Bandwidth Selection to Calculate Entropy. Journal of Chemical Information and Modeling, 2021, 61, 1533-1538.	5.4	16
23	pyPolyBuilder: Automated Preparation of Molecular Topologies and Initial Configurations for Molecular Dynamics Simulations of Arbitrary Supramolecules. Journal of Chemical Information and Modeling, 2021, 61, 1539-1544.	5.4	8
24	Shark Antibody Variable Domains Rigidify Upon Affinity Maturationâ€"Understanding the Potential of Shark Immunoglobulins as Therapeutics. Frontiers in Molecular Biosciences, 2021, 8, 639166.	3.5	9
25	Correcting cisâ€ŧransâ€ŧransgressions in macromolecular structure models. FEBS Journal, 2021, , .	4.7	1
26	Implementation of the Freely Jointed Chain Model to Assess Kinetics and Thermodynamics of Thermosensitive Coil–Globule Transition by Markov States. Journal of Physical Chemistry B, 2021, 125, 4898-4909.	2.6	4
27	Pore mutation N617D in the skeletal muscle DHPR blocks Ca2+ influx due to atypical high-affinity Ca2+ binding. ELife, 2021, 10, .	6.0	5
28	Structure and Zeatin Binding of the Peach Allergen <i>Pru p $1 < i$. Journal of Agricultural and Food Chemistry, 2021, 69, 8120-8129.</i>	5.2	10
29	Germline-Dependent Antibody Paratope States and Pairing Specific VH-VL Interface Dynamics. Frontiers in Immunology, 2021, 12, 675655.	4.8	11
30	Inhibitors of Fumarylacetoacetate Hydrolase Domain Containing Protein 1 (FAHD1). Molecules, 2021, 26, 5009.	3.8	0
31	Energy penalties enhance flexible receptor docking in a model cavity. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	17
32	Ion-pair interactions between voltage-sensing domain IV and pore domain I regulate CaV1.1 gating. Biophysical Journal, 2021, 120, 4429-4441.	0.5	3
33	A Lightâ€Triggerable Nanoparticle Library for the Controlled Release of Nonâ€Coding RNAs. Angewandte Chemie - International Edition, 2020, 59, 1985-1991.	13.8	24
34	V _H â€V _L interdomain dynamics observed by computer simulations and NMR. Proteins: Structure, Function and Bioinformatics, 2020, 88, 830-839.	2.6	28
35	The interplay of VSCF/VCI calculations and matrix-isolation IR spectroscopy – Mid infrared spectrum of CH3CH2F and CD3CD2F. Journal of Molecular Spectroscopy, 2020, 367, 111224.	1.2	22
36	Alpha arbonic Acid Revisited: Carbonic Acid Monomethyl Ester as a Solid and its Conformational Isomerism in the Gas Phase. Chemistry - A European Journal, 2020, 26, 285-305.	3.3	9

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37	Thermosensitive Hydration of Four Acrylamide-Based Polymers in Coil and Globule Conformations. Journal of Physical Chemistry B, 2020, 124, 9745-9756.	2.6	10
38	<i>N</i> -Heterocyclic Carbene Gold(I) Complexes: Mechanism of the Ligand Scrambling Reaction and Their Oxidation to Gold(III) in Aqueous Solutions. Inorganic Chemistry, 2020, 59, 15312-15323.	4.0	27
39	Solvation Thermodynamics in Different Solvents: Water–Chloroform Partition Coefficients from Grid Inhomogeneous Solvation Theory. Journal of Chemical Information and Modeling, 2020, 60, 3843-3853.	5.4	11
40	Replacement of the Cobalt Center of Vitamin B ₁₂ by Nickel: Nibalamin and Nibyric Acid Prepared from Metalâ€Free B ₁₂ â€Ligands Hydrogenobalamin and Hydrogenobyric Acid. Angewandte Chemie - International Edition, 2020, 59, 20129-20136.	13.8	18
41	On the synergy of matrix-isolation infrared spectroscopy and vibrational configuration interaction computations. Theoretical Chemistry Accounts, 2020, 139, 174.	1.4	3
42	Polarizable and non-polarizable force fields: Protein folding, unfolding, and misfolding. Journal of Chemical Physics, 2020, 153, 185102.	3.0	26
43	Local and Global Rigidification Upon Antibody Affinity Maturation. Frontiers in Molecular Biosciences, 2020, 7, 182.	3.5	29
44	Profiling selectivity of chagasin mutants towards cysteine proteases cruzain or cathepsin L through molecular dynamics simulations. Journal of Biomolecular Structure and Dynamics, 2020, 39, 1-13.	3.5	1
45	T-Cell Receptor CDR3 Loop Conformations in Solution Shift the Relative $\hat{Vl}\pm\hat{Vl}^2$ Domain Distributions. Frontiers in Immunology, 2020, 11, 1440.	4.8	17
46	Decomposing anharmonicity and mode-coupling from matrix effects in the IR spectra of matrix-isolated carbon dioxide and methane. Physical Chemistry Chemical Physics, 2020, 22, 17932-17947.	2.8	9
47	Two CaV3.3 (CACNA1I) Gain-of-Function Mutations Linked to Epilepsy and Intellectual Disability Affect Gating Properties and the Window Current. Biophysical Journal, 2020, 118, 106a.	0.5	0
48	Polyreactive Broadly Neutralizing B cells Are Selected to Provide Defense against Pandemic Threat Influenza Viruses. Immunity, 2020, 53, 1230-1244.e5.	14.3	61
49	Antibodies exhibit multiple paratope states influencing VH–VL domain orientations. Communications Biology, 2020, 3, 589.	4.4	38
50	In silico Design of Phl p 6 Variants With Altered Fold-Stability Significantly Impacts Antigen Processing, Immunogenicity and Immune Polarization. Frontiers in Immunology, 2020, 11, 1824.	4.8	8
51	Replacement of the Cobalt Center of Vitamin B ₁₂ by Nickel: Nibalamin and Nibyric Acid Prepared from Metalâ€Free B ₁₂ â€Ligands Hydrogenobalamin and Hydrogenobyric Acid. Angewandte Chemie, 2020, 132, 20304-20311.	2.0	2
52	Surprisingly Fast Interface and Elbow Angle Dynamics of Antigen-Binding Fragments. Frontiers in Molecular Biosciences, 2020, 7, 609088.	3.5	11
53	The intermolecular anthracene-transfer in a regiospecific antipodal C ₆₀ difunctionalization. Organic and Biomolecular Chemistry, 2020, 18, 4090-4103.	2.8	1
54	Ni[B 2 (SO 4) 4] and Co[B 2 (SO 4) 4]: Unveiling Systematic Trends in Phyllosilicate Analogue Borosulfates. Chemistry - A European Journal, 2020, 26, 17405-17415.	3.3	12

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55	Macrocycle Cell Permeability Measured by Solvation Free Energies in Polar and Apolar Environments. Journal of Chemical Information and Modeling, 2020, 60, 3508-3517.	5.4	15
56	Dynamics Rationalize Proteolytic Susceptibility of the Major Birch Pollen Allergen Bet ν 1. Frontiers in Molecular Biosciences, 2020, 7, 18.	3.5	6
57	T-Cell Receptor Variable \hat{l}^2 Domains Rigidify During Affinity Maturation. Scientific Reports, 2020, 10, 4472.	3.3	20
58	STACKED – <u>S</u> olvation <u>T</u> heory of <u>A</u> romatic <u>C</u> omplexes as <u>K</u> ey for <u>E</u> stimating <u>D</u> rug Binding. Journal of Chemical Information and Modeling, 2020, 60, 2304-2313.	5.4	10
59	Protein-Protein Binding as a Two-Step Mechanism: Preselection of Encounter Poses during the Binding of BPTI and Trypsin. Biophysical Journal, 2020, 119, 652-666.	0.5	22
60	Local Unfolding Relates to Proteolytic Susceptibility of the Major Birch Pollen Allergen Bet ν 1. Biophysical Journal, 2020, 118, 502a.	0.5	0
61	Biophysical classification of a CACNA1D de novo mutation as a high-risk mutation for a severe neurodevelopmental disorder. Molecular Autism, 2020, 11, 4.	4.9	33
62	Catalytic Site p <i>K</i> _a Values of Aspartic, Cysteine, and Serine Proteases: Constant pH MD Simulations. Journal of Chemical Information and Modeling, 2020, 60, 3030-3042.	5.4	44
63	Antibody CDR loops as ensembles in solution vs. canonical clusters from X-ray structures. MAbs, 2020, 12, 1744328.	5.2	34
64	pH-Induced Local Unfolding of the Phl p 6 Pollen Allergen From cpH-MD. Frontiers in Molecular Biosciences, 2020, 7, 603644.	3.5	5
65	Charge Anisotropy of Nitrogen: Where Chemical Intuition Fails. Journal of Chemical Theory and Computation, 2020, 16, 4443-4453.	5.3	8
66	Sodium-induced population shift drives activation of thrombin. Scientific Reports, 2020, 10, 1086.	3.3	8
67	Zinc Substitution of Cobalt in Vitaminâ€B12: Zincobyric acid and Zincobalamin as Luminescent Structural B12â€Mimics. Angewandte Chemie - International Edition, 2019, 58, 14568-14572.	13.8	25
68	Zinc Substitution of Cobalt in Vitaminâ€B12: Zincobyric acid and Zincobalamin as Luminescent Structural B12â€Mimics. Angewandte Chemie, 2019, 131, 14710-14714.	2.0	4
69	Solvation Free Energy as a Measure of Hydrophobicity: Application to Serine Protease Binding Interfaces. Journal of Chemical Theory and Computation, 2019, 15, 5872-5882.	5.3	42
70	Die Hydrogenobyrsäreâ€Struktur enthüllt den Corrinâ€Liganden als entatisches Zustandsmodul zur Steigerung der Katalyseaktivitävon B ₁₂ â€Cofaktoren. Angewandte Chemie, 2019, 131, 10869-10873.	2.0	8
71	Novel Dual Ligands Targeting Sigma 1 Receptor and Acetylcholinesterase Endowed with Antioxidant Properties. Proceedings (mdpi), 2019, 22, .	0.2	0
72	Toward Elimination of Discrepancies between Theory and Experiment: Anharmonic Rotational–Vibrational Spectrum of Water in Solid Noble Gas Matrices. Journal of Physical Chemistry A, 2019, 123, 8234-8242.	2.5	9

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73	pH-Dependent Protonation of the Phl p 6 Pollen Allergen Studied by NMR and cpH-aMD. Journal of Chemical Theory and Computation, 2019, 15, 5716-5726.	5.3	10
74	Molecular Dynamics Gives New Insights into the Glucose Tolerance and Inhibition Mechanisms on \hat{l}^2 -Glucosidases. Molecules, 2019, 24, 3215.	3.8	17
75	Hydration of Aromatic Heterocycles as an Adversary of π-Stacking. Journal of Chemical Information and Modeling, 2019, 59, 4209-4219.	5.4	11
76	Benzimidazole inhibitors of the major cysteine protease of <i>Trypanosoma brucei</i> . Future Medicinal Chemistry, 2019, 11, 1537-1551.	2.3	7
77	Coil–Globule Transition Thermodynamics of Poly(<i>N</i> Chemistry B, 2019, 123, 8838-8847.	2.6	45
78	CDR-H3 loop ensemble in solution – conformational selection upon antibody binding. MAbs, 2019, 11, 1077-1088.	5.2	49
79	The Hydrogenobyric Acid Structure Reveals the Corrin Ligand as an Entatic State Module Empowering B ₁₂ Cofactors for Catalysis. Angewandte Chemie - International Edition, 2019, 58, 10756-10760.	13.8	30
80	Mechanism of Olefin Metathesis with Neutral and Cationic Molybdenum Imido Alkylidene <i>N-</i> Heterocyclic Carbene Complexes. Journal of the American Chemical Society, 2019, 141, 8264-8276.	13.7	38
81	Structure Modelling of CaV1.1 Reveals Functional Trans-Domain Interactions Involved in Voltage Sensing. Biophysical Journal, 2019, 116, 112a.	0.5	0
82	Conformational selection of allergen-antibody complexesâ€"surface plasticity of paratopes and epitopes. Protein Engineering, Design and Selection, 2019, 32, 513-523.	2.1	17
83	Antibody humanization—the Influence of the antibody framework on the CDR-H3 loop ensemble in solution. Protein Engineering, Design and Selection, 2019, 32, 411-422.	2.1	17
84	Transitions of CDR-L3 Loop Canonical Cluster Conformations on the Micro-to-Millisecond Timescale. Frontiers in Immunology, 2019, 10, 2652.	4.8	18
85	Understanding Structure–Activity Relationships for Trypanosomal Cysteine Protease Inhibitors by Simulations and Free Energy Calculations. Journal of Chemical Information and Modeling, 2019, 59, 137-148.	5.4	17
86	Toward novel inhibitors against KdsB: a highly specific and selective broad-spectrum bacterial enzyme. Journal of Biomolecular Structure and Dynamics, 2019, 37, 1326-1345.	3.5	30
87	The fumarylacetoacetate hydrolase (FAH) superfamily of enzymes: multifunctional enzymes from microbes to mitochondria. Biochemical Society Transactions, 2018, 46, 295-309.	3.4	30
88	Cu[B ₂ (SO ₄) ₄] and Cu[B(SO ₄) ₂ (HSO ₄)]â€"Two Silicate Analogue Borosulfates Differing in their Dimensionality: A Comparative Study of Stability and Acidity. Angewandte Chemie - International Edition, 2018, 57, 9548-9552.	13.8	28
89	Peptidic Macrocycles - Conformational Sampling and Thermodynamic Characterization. Journal of Chemical Information and Modeling, 2018, 58, 982-992.	5.4	55
90	Pyroâ€Phyllobilins: Elusive Chlorophyll Catabolites Lacking a Critical Carboxylate Function of the Natural Chlorophylls. Chemistry - A European Journal, 2018, 24, 2987-2998.	3.3	11

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91	Cobaltocenylidene: A Mesoionic Metalloceno Carbene, Stabilized in a Gold(III) Complex. Chemistry - A European Journal, 2018, 24, 3165-3169.	3.3	17
92	Highly Electrophilic, Catalytically Active and Redoxâ€Responsive Cobaltoceniumyl and Ferrocenyl Triazolylidene Coinage Metal Complexes. Chemistry - A European Journal, 2018, 24, 3742-3753.	3.3	67
93	An unexpected switch in peptide binding mode: from simulation to substrate specificity. Journal of Biomolecular Structure and Dynamics, 2018, 36, 4072-4084.	3.5	7
94	Low frequency vibrational anharmonicity and nuclear spin effects of Cl–(H2) and Cl–(D2). Journal of Chemical Physics, 2018, 149, 174310.	3.0	9
95	Cu[B ₂ (SO ₄) ₄] und Cu[B(SO ₄) ₂ (HSO ₄)] – zwei silicatanaloge Borosulfate unterschiedlicher Dimensionalitä Vergleich von Stabilitäund Aziditä Angewandte Chemie, 2018, 130. 9693-9697.	2.0	12
96	Structural basis for the bi-functionality of human oxaloacetate decarboxylase FAHD1. Biochemical Journal, 2018, 475, 3561-3576.	3.7	13
97	Phase Diagram of a Stratum Corneum Lipid Mixture. Journal of Physical Chemistry B, 2018, 122, 10505-10521.	2.6	13
98	Identification of dual Sigma1 receptor modulators/acetylcholinesterase inhibitors with antioxidant and neurotrophic properties, as neuroprotective agents. European Journal of Medicinal Chemistry, 2018, 158, 353-370.	5.5	14
99	Electrostatic recognition in substrate binding to serine proteases. Journal of Molecular Recognition, 2018, 31, e2727.	2.1	13
100	Novel Types of Hypermodified Fluorescent Phyllobilins from Breakdown of Chlorophyll in Senescent Leaves of Grapevine (<i>Vitis vinifera</i>). Chemistry - A European Journal, 2018, 24, 17268-17279.	3.3	15
101	Innentitelbild: Cu[B2 (SO4)4] und Cu[B(SO4)2 (HSO4)] - zwei silicatanaloge Borosulfate unterschiedlicher Dimensionalit¤ Vergleich von Stabilit¤und Azidit¤(Angew. Chem. 30/2018). Angewandte Chemie, 2018, 130, 9330-9330.	2.0	0
102	Discovery of Retinoic Acid-Related Orphan Receptor \hat{l}^3t Inverse Agonists via Docking and Negative Image-Based Screening. ACS Omega, 2018, 3, 6259-6266.	3.5	10
103	Characterizing the Diversity of the CDR-H3 Loop Conformational Ensembles in Relationship to Antibody Binding Properties. Frontiers in Immunology, 2018, 9, 3065.	4.8	73
104	High-pressure synthesis and crystal structure of In ₃ B ₅ O ₁₂ . Zeitschrift Fur Naturforschung - Section B Journal of Chemical Sciences, 2017, 72, 69-76.	0.7	6
105	Binding Pose Flip Explained via Enthalpic and Entropic Contributions. Journal of Chemical Information and Modeling, 2017, 57, 345-354.	5.4	20
106	Structure of the Major Apple Allergen MalÂdÂ1. Journal of Agricultural and Food Chemistry, 2017, 65, 1606-1612.	5.2	50
107	Carbonic acid monoethyl ester as a pure solid and its conformational isomerism in the gas-phase. RSC Advances, 2017, 7, 22222-22233.	3.6	11
108	Ag[B(SO ₄) ₂] – Synthesis, Crystal Structure, and Characterization of the First Preciousâ€Metal Borosulfate. European Journal of Inorganic Chemistry, 2017, 2017, 3981-3989.	2.0	19

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109	Molecular dynamics simulation studies of novel \hat{l}^2 -lactamase inhibitor. Journal of Molecular Graphics and Modelling, 2017, 74, 143-152.	2.4	53
110	Crystal structure of Pla I 1 reveals both structural similarity and allergenic divergence within the Ole e 1–like protein family. Journal of Allergy and Clinical Immunology, 2017, 140, 277-280.	2.9	14
111	CaB ₂ S ₄ O ₁₆ : A Borosulfate Exhibiting a New Structure Type with Phyllosilicate Analogue Topology. Chemistry - A European Journal, 2017, 23, 16773-16781.	3.3	25
112	Mechanisms Responsible for $i\%$ -Pore Currents in Ca ν Calcium Channel Voltage-Sensing Domains. Biophysical Journal, 2017, 113, 1485-1495.	0.5	16
113	Balance between hydration enthalpy and entropy is important for ice binding surfaces in Antifreeze Proteins. Scientific Reports, 2017, 7, 11901.	3.3	21
114	Frontispiece: CaB ₂ S ₄ O ₁₆ : A Borosulfate Exhibiting a New Structure Type with Phyllosilicate Analogue Topology. Chemistry - A European Journal, 2017, 23, .	3.3	0
115	High-Pressure Synthesis and Characterization of the Ammonium Yttrium Borate (NH4)YB8O14. Inorganic Chemistry, 2017, 56, 14291-14299.	4.0	9
116	Gaining in pan-affinity towards sigma 1 and sigma 2 receptors. SAR studies on arylalkylamines. Bioorganic and Medicinal Chemistry, 2017, 25, 11-19.	3.0	15
117	Molecular Connectivity Predefines Polypharmacology: Aliphatic Rings, Chirality, and sp3 Centers Enhance Target Selectivity. Frontiers in Pharmacology, 2017, 8, 552.	3.5	16
118	Discovery and Characterization of Diazenylaryl Sulfonic Acids as Inhibitors of Viral and Bacterial Neuraminidases. Frontiers in Microbiology, 2017, 8, 205.	3.5	13
119	Determinants of Macromolecular Specificity from Proteomics-De rived Peptide Substrate Data. Current Protein and Peptide Science, 2017, 18, 905-913.	1.4	1
120	Conformational Flexibility Differentiates Naturally Occurring Bet v 1 Isoforms. International Journal of Molecular Sciences, 2017, 18, 1192.	4.1	18
121	Prediction of blood:air and fat:air partition coefficients of volatile organic compounds for the interpretation of data in breath gas analysis. Journal of Breath Research, 2016, 10, 017103.	3.0	15
122	Asymmetric arginine dimethylation of RelA provides a repressive mark to modulate TNFα/NF-κB response. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 4326-4331.	7.1	60
123	Orthorhombic HPâ€ <i>RE</i> OF (<i>RE</i> = Pr, Nd, Sm – Gd) – Highâ€Pressure Syntheses and Singleâ€Crys Structures (<i>RE</i> = Nd, Sm, Eu). Zeitschrift Fur Anorganische Und Allgemeine Chemie, 2016, 642, 1134-1142.	stal 1.2	11
124	DPPC-cholesterol phase diagram using coarse-grained Molecular Dynamics simulations. Biochimica Et Biophysica Acta - Biomembranes, 2016, 1858, 2846-2857.	2.6	95
125	Chlorophyllâ€Derived Yellow Phyllobilins of Higher Plants as Mediumâ€Responsive Chiral Photoswitches. Angewandte Chemie - International Edition, 2016, 55, 15760-15765.	13.8	24
126	Enthalpic and Entropic Contributions to Hydrophobicity. Journal of Chemical Theory and Computation, 2016, 12, 4600-4610.	5.3	68

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127	Interaction mechanisms of a melatonergic inhibitor in the melatonin synthesis pathway. Journal of Molecular Liquids, 2016, 221, 507-517.	4.9	39
128	Von Chlorophyll abstammende gelbe Phyllobiline h $\tilde{A}\P$ herer Pflanzen als umgebungsgesteuerte, chirale Photoschalter. Angewandte Chemie, 2016, 128, 15992-15997.	2.0	4
129	Innentitelbild: Von Chlorophyll abstammende gelbe Phyllobiline h \tilde{A} herer Pflanzen als umgebungsgesteuerte, chirale Photoschalter (Angew. Chem. 51/2016). Angewandte Chemie, 2016, 128, 15912-15912.	2.0	0
130	Sequence diversity of NanA manifests in distinct enzyme kinetics and inhibitor susceptibility. Scientific Reports, 2016, 6, 25169.	3.3	14
131	Chlorophyll Catabolites in Fall Leaves of the Wych Elm Tree Present a Novel Glycosylation Motif. Chemistry - A European Journal, 2016, 22, 9498-9503.	3.3	23
132	Protease Inhibitors in View of Peptide Substrate Databases. Journal of Chemical Information and Modeling, 2016, 56, 1228-1235.	5.4	4
133	Kinetic barriers in the isomerization of substituted ureas: implications for computer-aided drug design. Journal of Computer-Aided Molecular Design, 2016, 30, 391-400.	2.9	10
134	Localization of Millisecond Dynamics: Dihedral Entropy from Accelerated MD. Journal of Chemical Theory and Computation, 2016, 12, 3449-3455.	5.3	23
135	Fold stability during endolysosomal acidification is a key factor for allergenicity and immunogenicity of the major birch pollen allergen. Journal of Allergy and Clinical Immunology, 2016, 137, 1525-1534.	2.9	69
136	Quantitative Correlation of Conformational Binding Enthalpy with Substrate Specificity of Serine Proteases. Journal of Physical Chemistry B, 2016, 120, 299-308.	2.6	5
137	New High-Pressure Gallium Borate Ga ₂ B ₃ O ₇ (OH) with Photocatalytic Activity. Inorganic Chemistry, 2016, 55, 676-681.	4.0	36
138	Highâ€Pressure Synthesis of Cd(NH ₃) ₂ [B ₃ O ₅ (NH ₃)] ₂ : Pioneering the Way to the Substance Class of Ammine Borates. Angewandte Chemie - International Edition, 2015, 54, 6360-6363.	13.8	33
139	Ice nucleation by water-soluble macromolecules. Atmospheric Chemistry and Physics, 2015, 15, 4077-4091.	4.9	198
140	Interface dynamics explain assembly dependency of influenza neuraminidase catalytic activity. Journal of Biomolecular Structure and Dynamics, 2015, 33, 104-120.	3.5	24
141	Independent Metrics for Protein Backbone and Side-Chain Flexibility: Time Scales and Effects of Ligand Binding. Journal of Chemical Theory and Computation, 2015, 11, 851-860.	5.3	24
142	Structure, Thermal Behavior, and Vibrational Spectroscopy of the Silver Borate AgB _{0₅. European Journal of Inorganic Chemistry, 2015, 2015, 527-533.}	2.0	10
143	CACNA1D De Novo Mutations in Autism Spectrum Disorders Activate Cav1.3 L-Type Calcium Channels. Biological Psychiatry, 2015, 77, 816-822.	1.3	147
144	Unique Macrocycles in the Taiwan Traditional Chinese Medicine Database. Planta Medica, 2015, 81, 459-466.	1.3	6

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145	Antipneumococcal activity of neuraminidase inhibiting artocarpin. International Journal of Medical Microbiology, 2015, 305, 289-297.	3.6	32
146	Complementary assays helping to overcome challenges for identifying neuraminidase inhibitors. Future Virology, 2015, 10, 77-88.	1.8	23
147	Identification of FAH Domain-containing Protein 1 (FAHD1) as Oxaloacetate Decarboxylase. Journal of Biological Chemistry, 2015, 290, 6755-6762.	3.4	37
148	Strong Nonadditivity as a Key Structure–Activity Relationship Feature: Distinguishing Structural Changes from Assay Artifacts. Journal of Chemical Information and Modeling, 2015, 55, 483-494.	5.4	38
149	Matched Peptides: Tuning Matched Molecular Pair Analysis for Biopharmaceutical Applications. Journal of Chemical Information and Modeling, 2015, 55, 2315-2323.	5.4	6
150	FAH Domain Containing Protein 1 (FAHD-1) Is Required for Mitochondrial Function and Locomotion Activity in C. elegans. PLoS ONE, 2015, 10, e0134161.	2.5	13
151	Dynamics Govern Specificity of a Protein-Protein Interface: Substrate Recognition by Thrombin. PLoS ONE, 2015, 10, e0140713.	2.5	24
152	A Cross-Reactive Human Single-Chain Antibody for Detection of Major Fish Allergens, Parvalbumins, and Identification of a Major IgE-Binding Epitope. PLoS ONE, 2015, 10, e0142625.	2.5	12
153	Characterizing Protease Specificity: How Many Substrates Do We Need?. PLoS ONE, 2015, 10, e0142658.	2.5	25
154	The Impact of Nitration on the Structure and Immunogenicity of the Major Birch Pollen Allergen Bet ν 1.0101. PLoS ONE, 2014, 9, e104520.	2.5	70
155	A gatekeeper helix determines the substrate specificity of Sjögren–Larsson Syndrome enzyme fatty aldehyde dehydrogenase. Nature Communications, 2014, 5, 4439.	12.8	55
156	Ligand Binding Modulates the Structural Dynamics and Compactness of the Major Birch Pollen Allergen. Biophysical Journal, 2014, 107, 2972-2981.	0.5	35
157	Precursors for cytochrome P450 profiling breath tests from an in silico screening approach. Journal of Breath Research, 2014, 8, 046001.	3.0	1
158	Specificity of a protein–protein interface: Local dynamics direct substrate recognition of effector caspases. Proteins: Structure, Function and Bioinformatics, 2014, 82, 546-555.	2.6	20
159	Non- 13 CO 2 targeted breath tests: a feasibility study. Journal of Breath Research, 2014, 8, 046005.	3.0	11
160	Dynamic regulation of phenylalanine hydroxylase. Pteridines, 2014, 25, 33-39.	0.5	7
161	Matched Molecular Pair Analysis: Significance and the Impact of Experimental Uncertainty. Journal of Medicinal Chemistry, 2014, 57, 3786-3802.	6.4	62
162	Heteroaromatic π-Stacking Energy Landscapes. Journal of Chemical Information and Modeling, 2014, 54, 1371-1379.	5.4	144

#	Article	IF	Citations
163	Computer-Guided Approach to Access the Anti-influenza Activity of Licorice Constituents. Journal of Natural Products, 2014, 77, 563-570.	3.0	38
164	A Conserved Acidic Residue in Phenylalanine Hydroxylase Contributes to Cofactor Affinity and Catalysis. Biochemistry, 2014, 53, 6834-6848.	2.5	7
165	Charge Anisotropy: Where Atomic Multipoles Matter Most. Journal of Chemical Theory and Computation, 2014, 10, 4488-4496.	5.3	54
166	Limits to molecular matched-pair analysis: the experimental uncertainty case. Journal of Cheminformatics, 2014, 6, O6.	6.1	1
167	Entropy in specificity and thermodynamics of binding. Journal of Cheminformatics, 2014, 6, O8.	6.1	0
168	(How to) Profit from Molecular Dynamics-based Ensemble Docking. Challenges and Advances in Computational Chemistry and Physics, 2014, , 501-538.	0.6	3
169	Linking ethnopharmacology with phenotypic and virtual screening for the identification of antiviral agents from natural sources. Planta Medica, 2014, 80, .	1.3	0
170	An additional substrate binding site in a bacterial phenylalanine hydroxylase. European Biophysics Journal, 2013, 42, 691-708.	2.2	10
171	Substrate Sequences Tell Similar Stories as Binding Cavities: Commentary. Journal of Chemical Information and Modeling, 2013, 53, 3115-3116.	5.4	4
172	Deriving Static Atomic Multipoles from the Electrostatic Potential. Journal of Chemical Information and Modeling, 2013, 53, 3410-3417.	5.4	25
173	A Novel Paramagnetic Relaxation Enhancement Tag for Nucleic Acids: A Tool to Study Structure and Dynamics of RNA. ACS Chemical Biology, 2013, 8, 2697-2706.	3.4	31
174	Matrix Isolation Studies of Carbonic Acidâ€"The Vapor Phase above the β-Polymorph. Journal of the American Chemical Society, 2013, 135, 7732-7737.	13.7	33
175	Entropy from State Probabilities: Hydration Entropy of Cations. Journal of Physical Chemistry B, 2013, 117, 6466-6472.	2.6	23
176	Characterisation of Nox4 Inhibitors from Edible Plants. Planta Medica, 2013, 79, 244-252.	1.3	15
177	Substrate-Driven Mapping of the Degradome by Comparison of Sequence Logos. PLoS Computational Biology, 2013, 9, e1003353.	3.2	23
178	Cleavage Entropy as Quantitative Measure of Protease Specificity. PLoS Computational Biology, 2013, 9, e1003007.	3.2	49
179	Reciprocal regulation of PKA and Rac signaling. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 8531-8536.	7.1	42
180	New insights into the anti-influenza activity of licorice constituents. Planta Medica, 2013, 79, .	1.3	2

#	Article	IF	Citations
181	Bioprospecting diverse plant species for antiviral agents against upper respiratory tract infections. Planta Medica, 2013, 79, .	1.3	O
182	Influenza neuraminidase: A druggable target for natural products. Natural Product Reports, 2012, 29, 11-36.	10.3	146
183	Porphyrin-LEGO®: synthesis of a hexafullereno-diporphyrin using porphyrins programmed for [4+2]-cycloaddition. Chemical Communications, 2012, 48, 4359.	4.1	17
184	Identification of PPARgamma Partial Agonists of Natural Origin (I): Development of a Virtual Screening Procedure and In Vitro Validation. PLoS ONE, 2012, 7, e50816.	2.5	48
185	Dynamic Regulation of Phenylalanine Hydroxylase by Simulated Redox Manipulation. PLoS ONE, 2012, 7, e53005.	2.5	27
186	Effects of Pooling Samples on the Performance of Classification Algorithms: A Comparative Study. Scientific World Journal, The, 2012, 2012, 1-10.	2.1	13
187	Identification of Novel Liver X Receptor Activators by Structure-Based Modeling. Journal of Chemical Information and Modeling, 2012, 52, 1391-1400.	5.4	16
188	Local structural order in carbonic acid polymorphs: Raman and FTâ€IR spectroscopy. Journal of Raman Spectroscopy, 2012, 43, 108-115.	2.5	26
189	Identification of Novel Functional Inhibitors of Acid Sphingomyelinase. PLoS ONE, 2011, 6, e23852.	2.5	145
190	Minor Groove Binders and Drugs Targeting Proteins Cover Complementary Regions in Chemical Shape Space. Journal of Chemical Information and Modeling, 2011, 51, 2223-2232.	5.4	14
191	A GRID-Derived Water Network Stabilizes Molecular Dynamics Computer Simulations of a Protease. Journal of Chemical Information and Modeling, 2011, 51, 2860-2867.	5.4	13
192	Qualitative prediction of blood–brain barrier permeability on a large and refined dataset. Journal of Computer-Aided Molecular Design, 2011, 25, 1095-1106.	2.9	97
193	Backbone flexibility controls the activity and specificity of a protein-protein interface – specificity in snake venom metalloproteases (SVMPs). Journal of Cheminformatics, 2011, 3, .	6.1	0
194	A challenging system: Free energy prediction for factor Xa. Journal of Computational Chemistry, 2011, 32, 1743-1752.	3.3	75
195	Rýcktitelbild: Spektroskopische Beobachtung von matrixisolierter KohlensÃ u re, abgeschieden aus der Gasphase (Angew. Chem. 8/2011). Angewandte Chemie, 2011, 123, 1988-1988.	2.0	0
196	Spectroscopic Observation of Matrixâ€Isolated Carbonic Acid Trapped from the Gas Phase. Angewandte Chemie - International Edition, 2011, 50, 1939-1943.	13.8	50
197	Back Cover: Spectroscopic Observation of Matrix-Isolated Carbonic Acid Trapped from the Gas Phase (Angew. Chem. Int. Ed. 8/2011). Angewandte Chemie - International Edition, 2011, 50, 1946-1946.	13.8	2
198	Novel neuraminidase inhibitors: identification, biological evaluation and investigations of the binding mode. Future Medicinal Chemistry, 2011, 3, 437-450.	2.3	34

#	Article	IF	Citations
199	Development of Anti-Viral Agents Using Molecular Modeling and Virtual Screening Techniques. Infectious Disorders - Drug Targets, 2011, 11, 64-93.	0.8	43
200	Backbone Flexibility Controls the Activity and Specificity of a Proteinâ^'Protein Interface: Specificity in Snake Venom Metalloproteases. Journal of the American Chemical Society, 2010, 132, 10330-10337.	13.7	51
201	Dispersion dominated halogenâ€"ï€ interactions: energies and locations of minima. Physical Chemistry Chemical Physics, 2010, 12, 14941.	2.8	73
202	Stabilizing of a Globular Protein by a Highly Complex Water Network: A Molecular Dynamics Simulation Study on Factor Xa. Journal of Physical Chemistry B, 2010, 114, 7405-7412.	2.6	59
203	One Concept, Three Implementations of 3D Pharmacophore-Based Virtual Screening: Distinct Coverage of Chemical Search Space. Journal of Chemical Information and Modeling, 2010, 50, 1241-1247.	5.4	54
204	Antiviral Potential and Molecular Insight into Neuraminidase Inhibiting Diarylheptanoids from <i>Alpinia katsumadai</i> . Journal of Medicinal Chemistry, 2010, 53, 778-786.	6.4	114
205	Challenges for Computer Simulations in Drug Design. Challenges and Advances in Computational Chemistry and Physics, 2010, , 431-463.	0.6	2
206	Raman Spectroscopic Study of the Phase Transition of Amorphous to Crystalline β arbonic Acid. Angewandte Chemie - International Edition, 2009, 48, 2690-2694.	13.8	33
207	Hydrogen-Bonding Patterns of Minor Groove-Binderâ^'DNA Complexes Reveal Criteria for Discovery of New Scaffolds. Journal of Chemical Information and Modeling, 2009, 49, 1063-1069.	5.4	11
208	How To Optimize Shape-Based Virtual Screening: Choosing the Right Query and Including Chemical Information. Journal of Chemical Information and Modeling, 2009, 49, 678-692.	5.4	178
209	Theoretical Prediction of Hydrogen Bond Strength for Use in Molecular Modeling. Journal of Chemical Information and Modeling, 2009, 49, 2067-2076.	5.4	37
210	Sequenceâ€Specific Positions of Water Molecules at the Interface between DNA and Minor Groove Binders. ChemPhysChem, 2008, 9, 2766-2771.	2.1	14
211	The Protein Data Bank (PDB), Its Related Services and Software Tools as Key Components for In Silico Guided Drug Discovery. Journal of Medicinal Chemistry, 2008, 51, 7021-7040.	6.4	91
212	DNA Minor Groove Pharmacophores Describing Sequence Specific Properties. Journal of Chemical Information and Modeling, 2007, 47, 1580-1589.	5.4	27
213	Modeling the heterogeneous reaction probability for chlorine nitrate hydrolysis on ice. Journal of Geophysical Research, 2006, 111 , .	3.3	3
214	M.Taql facilitates the base flipping via an unusual DNA backbone conformation. Biopolymers, 2005, 79, 128-138.	2.4	2
215	Towards an Understanding of DNA Recognition by the Methyl-CpG Binding Domain 1. Journal of Biomolecular Structure and Dynamics, 2005, 22, 695-706.	3.5	12
216	Z-DNA's Conformer Substates Revealed by FT-IR Difference Spectroscopy of Nonoriented Left-Handed Double Helical Poly(dG-dC). Journal of Biomolecular Structure and Dynamics, 2005, 22, 595-614.	3.5	18

#	Article	IF	CITATIONS
217	TheN6-Methyl Group of Adenine Further Increases the BI Stability of DNA Compared toC5-Methyl Groups. Journal of Physical Chemistry B, 2005, 109, 557-564.	2.6	7
218	Daunomycin Intercalation Stabilizes Distinct Backbone Conformations of DNA. Journal of Biomolecular Structure and Dynamics, 2004, 21, 713-724.	3.5	24
219	Cooperative effects on the formation of intercalation sites. Nucleic Acids Research, 2004, 32, 4696-4703.	14.5	38
220	Water-mediated contacts in thetrp-repressor operator complex recognition process. Biopolymers, 2004, 73, 668-681.	2.4	12
221	Sulfurous acid (H2SO3) on lo?. Icarus, 2004, 169, 242-249.	2.5	20
222	The ground-state tunneling splitting of various carboxylic acid dimers. Journal of Chemical Physics, 2004, 120, 631-637.	3.0	71
223	Dynamics of DNA:  BI and BII Phosphate Backbone Transitions. Journal of Physical Chemistry B, 2004, 108, 2470-2476.	2.6	21
224	On the Formation of the Sulfonate Ion from Hydrated Sulfur Dioxide. Journal of Physical Chemistry A, 2004, 108, 3859-3864.	2.5	35
225	Double hydrogen tunneling revisited: The breakdown of experimental tunneling criteria. Journal of Chemical Physics, 2004, 120, 11650-11657.	3.0	54
226	Mechanism of the Cisâ^'Trans Isomerization of Bis(glycinato)copper(II). Journal of Physical Chemistry B, 2004, 108, 2098-2102.	2.6	27
227	Extended method for adiabatic mode reordering. Journal of Computational Chemistry, 2003, 24, 386-395.	3.3	1
228	A QM-MM interface between CHARMM and TURBOMOLE: Implementation and application to systems in bulk phase and biologically active systems. Journal of Computational Chemistry, 2003, 24, 1240-1249.	3.3	23
229	Untersuchung der HBr-Ionisierung auf molekularer Ebene. Angewandte Chemie, 2003, 115, 2162-2164.	2.0	4
230	About the Stability of Sulfurous Acid (H2SO3) and Its Dimer ChemInform, 2003, 34, no.	0.0	0
231	Exploring HBr Ionization at the Molecular Level. ChemInform, 2003, 34, no.	0.0	O
232	Exploring HBr Ionization at the Molecular Level. Angewandte Chemie - International Edition, 2003, 42, 2114-2116.	13.8	14
233	Stepwise induced fit in the pico- to nanosecond time scale governs the complexation of theeven-skipped transcriptional repressor homeodomain to DNA. Biopolymers, 2003, 68, 139-149.	2.4	11
234	Reactions of HOBr + HCl + nH2O and HOBr + HBr + nH2O. Chemical Physics Letters, 2003, 372, 569-576.	2.6	7

#	Article	IF	Citations
235	The ground state tunneling splitting of the 2-pyridone2-hydroxypyridine dimer. Chemical Physics, 2003, 292, 47-52.	1.9	26
236	C5-Methylation of Cytosine in B-DNA Thermodynamically and Kinetically Stabilizes BI. Journal of the American Chemical Society, 2003, 125, 14990-14991.	13.7	35
237	Molecular Docking Studies of Natural Cholinesterase-Inhibiting Steroidal Alkaloids fromSarcococcasaligna. Journal of Medicinal Chemistry, 2003, 46, 5087-5090.	6.4	42
238	Hydration of Hydroxypyrrole Influences Binding of ImHpPyPy- \hat{l}^2 -Dp Polyamide to DNA. Journal of the American Chemical Society, 2003, 125, 1088-1095.	13.7	24
239	Energetic and Stereochemical Effects of the Protein Environment on Substrate:Â A Theoretical Study of Methylmalonyl-CoA Mutase. Journal of the American Chemical Society, 2003, 125, 1072-1078.	13.7	18
240	About the Kinetic Feasibility of the Lipscomb Mechanism in Human Carbonic Anhydrase II. Journal of Physical Chemistry B, 2003, 107, 12013-12020.	2.6	27
241	Modeling Anhydrous and Aqua Copper(II) Amino Acid Complexes:Â A New Molecular Mechanics Force Field Parametrization Based on Quantum Chemical Studies and Experimental Crystal Data. Inorganic Chemistry, 2003, 42, 2268-2279.	4.0	69
242	Toward elimination of discrepancies between theory and experiment: The gas-phase reaction of N2O5 with H2O. Physical Chemistry Chemical Physics, 2003, 5, 487-495.	2.8	27
243	The Conformer Substates of Nonoriented B-type DNA in Double Helical Poly(dG-dC). Journal of Biomolecular Structure and Dynamics, 2003, 20, 547-559.	3.5	1
244	Influence of Backbone Conformations of Human Carbonic Anhydrase II on Carbon Dioxide Hydration:  Hydration Pathways and Binding of Bicarbonate. Journal of the American Chemical Society, 2003, 125, 8921-8927.	13.7	28
245	The optimal tunneling path for the proton transfer in malonaldehyde. Journal of Chemical Physics, 2002, 117, 1962-1966.	3.0	80
246	An accurate semiclassical method to predict ground-state tunneling splittings. Journal of Chemical Physics, 2002, 117, 1967-1974.	3.0	47
247	Reactions of HOCl + HCl +nH2O and HOCl + HBr +nH2O. Journal of Physical Chemistry A, 2002, 106, 7850-7857.	2.5	19
248	Restructuring of Hydration Shells Rules the Low-Temperature Dynamics of B-DNA via Its Two Conformer Substates. Journal of Physical Chemistry B, 2002, 106, 3263-3274.	2.6	10
249	Indirect Readout of thetrp-Repressorâ^'Operator Complex by B-DNA's Backbone Conformation Transitionsâ€. Biochemistry, 2002, 41, 4088-4095.	2.5	26
250	The structure, modelling and dynamics of 2,7-diisopropoxy-1,8-diarylnaphthalenes. Perkin Transactions II RSC, 2002, , 1510-1519.	1.1	26
251	Characterization of the Vitamin E-Binding Properties of Human Plasma Afamin. Biochemistry, 2002, 41, 14532-14538.	2,5	103
252	Pvu II-Endonuclease Induces Structural Alterations at the Scissile Phosphate Group of its Cognate DNA. Journal of Molecular Biology, 2002, 324, 491-500.	4.2	7

#	Article	IF	CITATIONS
253	Towards the Experimental Decomposition Rate of Carbonic Acid (H2CO3) in Aqueous Solution. Chemistry - A European Journal, 2002, 8, 66-73.	3.3	84
254	About the Stability of Sulfurous Acid (H2SO3) and Its Dimer. Chemistry - A European Journal, 2002, 8, 5644-5651.	3.3	43
255	Influence of netropsin's charges on the minor groove width of d(CGCGAATTCGCG)2. Biopolymers, 2002, 61, 276-286.	2.4	4
256	Estimation of the BII Conformer Substate Population in Nonoriented Hydrated B-DNA via Curve Resolution of Infrared Spectra. Applied Spectroscopy, 2001, 55, 9-22.	2.2	4
257	Significance of Ligand Tails for Interaction with the Minor Groove of B-DNA. Biophysical Journal, 2001, 81, 1588-1599.	0.5	26
258	The structure, modelling and dynamics of hindered 5,6-diarylacenaphthenes. Perkin Transactions II RSC, 2001, , 459-467.	1.1	17
259	Structural Flexibility of the d(CCAGTACTGG)2B-DNA Decamer and Its Complex with Two Polyamides. Journal of Physical Chemistry B, 2001, 105, 3135-3142.	2.6	12
260	Complex of B-DNA with Polyamides Freezes DNA Backbone Flexibility. Journal of the American Chemical Society, 2001, 123, 5044-5049.	13.7	41
261	Bl⇌ BIISubstate Transitions Induce Changes in the Hydration of B-DNA, Potentially Mediating Signal Transduction from the Minor to Major Groove. Journal of Physical Chemistry B, 2001, 105, 10379-10387.	2.6	19
262	Water-Mediated Proton Transfer:Â A Mechanistic Investigation on the Example of the Hydration of Sulfur Oxides. Journal of Physical Chemistry A, 2001, 105, 5137-5145.	2.5	69
263	Calculation of sequence-dependent free energies of hydration of dipeptides formed by alanine and glycine. Journal of Computational Chemistry, 2001, 22, 846-860.	3.3	2
264	The Reaction Rate Constant of Chlorine Nitrate Hydrolysis. Chemistry - A European Journal, 2001, 7, 1662-1669.	3.3	11
265	Prediction of the structure of human Janus kinase 2 (JAK2) comprising the two carboxy-terminal domains reveals a mechanism for autoregulation. Protein Engineering, Design and Selection, 2001, 14, 27-37.	2.1	154
266	On the Surprising Kinetic Stability of Carbonic Acid (H2CO3). Angewandte Chemie - International Edition, 2000, 39, 891-894.	13.8	152
267	B-DNA's dynamics and conformational substates revealed by calorimetric enthalpy relaxation and fourier transform infrared spectroscopy. Journal of Molecular Liquids, 2000, 86, 137-149.	4.9	7
268	The role of water in B-DNAs BI to BII conformer substates interconversion: a combined study by calorimetry, FT-IR spectroscopy and computer simulation. Chemical Physics, 2000, 258, 391-404.	1.9	20
269	Toward elimination of discrepancies between theory and experiment: The rate constant of the atmospheric conversion of SO3 to H2SO4. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 8874-8878.	7.1	91
270	Automated Docking of Ligands to Antibodies: Methods and Applications. Methods, 2000, 20, 280-291.	3.8	72

#	Article	IF	CITATIONS
271	Elbow Flexibility and Ligand-Induced Domain Rearrangements in Antibody Fab NC6.8: Large Effects of a Small Hapten. Biophysical Journal, 2000, 79, 614-628.	0.5	31
272	On the competing hydrations of sulfur dioxide and sulfur trioxide in our atmosphere. Chemical Communications, 2000, , 999-1000.	4.1	25
273	B-DNA's BIIConformer Substate Population Increases with Decreasing Water Activity. 1. A Molecular Dynamics Study of d(CGCGAATTCGCG)2. Journal of Physical Chemistry B, 2000, 104, 11349-11353.	2.6	13
274	Nonoriented d(CGCGAATTCGCG)2Dodecamer Persists in the B-Form Even at Low Water Activity. Journal of the American Chemical Society, 2000, 122, 716-717.	13.7	23
275	B-DNA's BIIConformer Substate Population Increases with Decreasing Water Activity. 2. A Fourier Transform Infrared Spectroscopic Study of Nonoriented d(CGCGAATTCGCG)2. Journal of Physical Chemistry B, 2000, 104, 11354-11359.	2.6	8
276	Simulation of EcoRI Dodecamer Netropsin Complex Confirms Class I Complexation Mode. Journal of the American Chemical Society, 2000, 122, 3927-3931.	13.7	31
277	Application of multivariate data analysis methods to comparative molecular field analysis (CoMFA) data: proton affinities and pKa prediction for nucleic acids components. Journal of Computer-Aided Molecular Design, 1999, 13, 611-623.	2.9	24
278	Helix Morphology Changes in B-DNA Induced By Spontaneous B _I ⇌B _{II} Substate Interconversion. Journal of Biomolecular Structure and Dynamics, 1999, 17, 223-235.	3.5	27
279	Investigation of Cu2+ Hydration and the Jahnâ^'Teller Effect in Solution by QM/MM Monte Carlo Simulations. Journal of Physical Chemistry A, 1999, 103, 11387-11393.	2.5	69
280	Unexpected B II Conformer Substate Population in Unoriented Hydrated Films of the d(CGCGAATTCGCG) 2 Dodecamer and of Native B-DNA from Salmon Testes. Biophysical Journal, 1999, 77, 398-409.	0.5	33
281	Ligand Binding by Antibody IgE Lb4: Assessment of Binding Site Preferences Using Microcalorimetry, Docking, and Free Energy Simulations. Biophysical Journal, 1999, 76, 2966-2977.	0.5	12
282	Why Are Copper(II) Amino Acid Complexes Not Planar in Their Crystal Structures? An ab Initio and Molecular Mechanics Study. Inorganic Chemistry, 1999, 38, 2764-2774.	4.0	50
283	Temperature-Dependent Ways of Proton TransferA Benchmark Study on Cyclic HF Oligomers. Journal of Physical Chemistry A, 1999, 103, 9022-9028.	2.5	8
284	The hydration shell structure of Li+ investigated by Born–Oppenheimer ab initio QM/MM dynamics. Chemical Physics Letters, 1998, 286, 56-64.	2.6	89
285	Improving the predictive quality of CoMFA models. Journal of Computer - Aided Molecular Design, 1998, 12/14, 41-56.	1.0	6
286	Comparative molecular field analysis of artemisinin derivatives: ab initio versus semiempirical optimized structures. Journal of Computer-Aided Molecular Design, 1998, 12, 397-397.	2.9	14
287	Carbonic Acid in the Gas Phase and Its Astrophysical Relevance. Science, 1998, 279, 1332-1335.	12.6	125
288	Reactivity and Regioselectivity of Hydroxyl Radical Addition to Halogenated Ethenes. Journal of Physical Chemistry A, 1998, 102, 1583-1594.	2.5	81

#	Article	IF	Citations
289	Predictions of rate constants and estimates for tunneling splittings of concerted proton transfer in small cyclic water clusters. Journal of Chemical Physics, 1998, 109, 2672-2679.	3.0	57
290	Toward Elimination of Discrepancies between Theory and Experiment:  Double Proton Transfer in Dimers of Carboxylic Acids. Journal of the American Chemical Society, 1998, 120, 12595-12600.	13.7	72
291	Large Curvature Tunneling Effects Reveal Concerted Hydrogen Exchange Rates in Cyclic Hydrogen Fluoride Clusters Comparable to Carboxylic Acid Dimers. Journal of the American Chemical Society, 1998, 120, 404-412.	13.7	29
292	Conformationally Induced Changes in the Electronic Structures of Some Flexible Benzenes. A Molecular Orbital Model. Journal of the American Chemical Society, 1998, 120, 12573-12582.	13.7	57
293	B-DNA's Bl→ BIIConformer Substate Dynamics Is Coupled with Water Migration. Journal of Physical Chemistry B, 1998, 102, 8934-8940.	2.6	57
294	Ligand-induced domain movement in an antibody fab: molecular dynamics studies confirm the unique domain movement observed experimentally for fab NC6.8 upon complexation and reveal its segmental flexibility 1 1Edited by I. Wilson. Journal of Molecular Biology, 1998, 278, 301-306.	4.2	25
295	Highly Accurate Estimates of Hydrogen-Bond Energies Relying on Basis Set Convergence Patterns. Journal of Physical Chemistry A, 1998, 102, 1832-1836.	2.5	16
296	Structure, Reaction Enthalpies, Entropies, and Free Energies of Cationâ^'Molecule Complexes. A Theoretical Study by Means of the ab Initio Complete Basis Set CBS-Q Method. Journal of Physical Chemistry A, 1998, 102, 771-777.	2.5	38
297	Bornâ^'Oppenheimer ab Initio QM/MM Dynamics Simulations of Na+and K+in Water:Â From Structure Making to Structure Breaking Effects. Journal of Physical Chemistry A, 1998, 102, 10340-10347.	2.5	218
298	Quantitative Analysis of the Structural Requirements for Blockade of the N-Methyl-d-aspartate Receptor at the Phencyclidine Binding Site. Journal of Medicinal Chemistry, 1998, 41, 393-400.	6.4	38
299	Dangers of counterpoise corrected hypersurfaces. Advantages of basis set superposition improvement. Journal of Chemical Physics, 1998, 108, 3199-3204.	3.0	51
300	Improving the Predictive Quality of CoMFA Models. , 1998, , 41-56.		2
301	Has the Dimer of Carbonic Acid a Lower Energy Than Its Constituents Water and Carbon Dioxide?. Journal of the American Chemical Society, 1997, 119, 3782-3784.	13.7	70
302	Reaction-Path Dynamics of Hydroxyl Radical Reactions with Ethane and Haloethanes. Journal of Physical Chemistry A, 1997, 101, 4245-4253.	2.5	43
303	New Insights into the Dynamics of Concerted Proton Tunneling in Cyclic Water and Hydrogen Fluoride Clusters. Journal of Physical Chemistry A, 1997, 101, 4707-4716.	2.5	37
304	Solvation of Ca2+in Water Studied by Bornâ^'Oppenheimer ab Initio QM/MM Dynamics. Journal of Physical Chemistry A, 1997, 101, 6299-6309.	2.5	140
305	How acidic are thiocarboxylic S-acids and thiosilanoic S-acids (X^y YOSH, $X = H$, F, Cl, CH3, NH2; $Y = C$, Si)?. Computational and Theoretical Chemistry, 1997, 418, 179-187.	1.5	17
306	A critical analysis of electronic density functionals for structural, energetic, dynamic, and magnetic properties of hydrogen fluoride clusters. Journal of Computational Chemistry, 1997, 18, 1695-1719.	3.3	85

#	Article	IF	Citations
307	Comparative Molecular Field Analysis of Haptens Docked to the Multispecific Antibody IgE(Lb4). Journal of Medicinal Chemistry, 1996, 39, 3882-3888.	6.4	19
308	Heteroligation of a mouse monoclonal IgE antibody (La2) with small molecules, analysed by computer-aided automated docking. Molecular Immunology, 1996, 33, 129-144.	2.2	7
309	Structure and gas-phase acidity of oxalic acid and its disila derivative. A theoretical study by means of the DFT quantum theoretical method. Journal of the Chemical Society Perkin Transactions II, 1996, , 1743.	0.9	22
310	Different electrostatic descriptors in comparative molecular field analysis: A comparison of molecular electrostatic and coulomb potentials. Journal of Computational Chemistry, 1996, 17, 1296-1308.	3.3	36
311	Bidirectional molecular dynamics: Interpretation in terms of a modern formulation of classical mechanics. Journal of Computational Chemistry, 1996, 17, 1564-1570.	3.3	2
312	Prediction of IgE(Lb4)–ligand complex structures by automated docking. , 1996, 9, 239-246.		3
313	A QM/MM simulation method applied to the solution of Li+ in liquid ammonia. Chemical Physics, 1996, 211, 313-323.	1.9	175
314	Gas-phase acidities of HM(=X)XH (Mî—»C, Si; Xî—»O, S) acids calculated by ab initio molecular orbital methods at the G2 level of theory. Chemical Physics Letters, 1996, 263, 379-384.	2.6	28
315	Comparative docking studies on ligand binding to the multispecific antibodies IgE-La2 and IgE-Lb4. Journal of Computer-Aided Molecular Design, 1996, 10, 305-320.	2.9	23
316	Different electrostatic descriptors in comparative molecular field analysis: A comparison of molecular electrostatic and coulomb potentials. Journal of Computational Chemistry, 1996, 17, 1296-1308.	3.3	0
317	Salt induced peptide formation: on the selectivity of the copper induced peptide formation under possible prebiotic conditions. Inorganica Chimica Acta, 1995, 228, 207-214.	2.4	47
318	Hydrogen transitions between (HF)n Cnh structures (n=2â^'5) via Dnh transition states as models for hydrogen tunneling in hydrogen fluoride clusters. Chemical Physics Letters, 1995, 246, 455-462.	2.6	22
319	Cation binding effect on hydrogen bonding and the energetics of proton transfer in the system (CH3)3NH+…â^'OCOH. Computational and Theoretical Chemistry, 1995, 336, 7-15.	1.5	3
320	Interaction of local anaesthetics with an anionic receptor site. An ab initio SCF study on procaine, lidocaine, tocainide and mexiletine and their HCO2â ^{-/-} complexes. Computational and Theoretical Chemistry, 1995, 343, 141-147.	1.5	5
321	Migratory aptitude of hydrogen and stabilization of the transition-state by the non-migrating substituent in 1,2-migrations to electron deficient nitrogen. Chemical Physics Letters, 1994, 231, 289-294.	2.6	1
322	1,2-Shift of Carbon to Electron-Deficient Nitrogen Is Not a Nucleophilic Rearrangement. Ab Initio Study on a 1,2-Rearrangement in 1,2,4-Triazolium Salts. Journal of the American Chemical Society, 1994, 116, 6277-6283.	13.7	14
323	1,2-Carbon to nitrogen migrations. Part 2. Ab initio study on the rearrangement of (\hat{l}_{\pm} -methylazo)alkyl isocyanates. Journal of the Chemical Society Perkin Transactions II, 1994, , 2129-2135.	0.9	5
324	Evaporation cycle experiments â€" A simulation of salt-induced peptide synthesis under possible prebiotic conditions. Origins of Life and Evolution of Biospheres, 1993, 23, 167-176.	1.9	56

#	Article	IF	CITATIONS
325	Structure and acidity of carbamic acid, and its thio, dithio and sila dervatives. Journal of the Chemical Society, Faraday Transactions, 1993, 89, 2375.	1.7	20
326	Ab initio calculations concerning the reaction mechanism of the copper(II) catalyzed glycine condensation in aqueous sodium chloride solution. Chemical Physics Letters, 1992, 197, 181-186.	2.6	34
327	Challenge Accepted $\hat{a} \in ``Paratope States in Solution Improve Structure Prediction and Docking. SSRN Electronic Journal, 0, , .$	0.4	2