

Jean-Didier MarÃ©chal

List of Publications by Year in descending order

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127
papers

3,418
citations

109321

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docs citations

140
times ranked

3964
citing authors

#	ARTICLE	IF	CITATIONS
1	An Artificial Heme Enzyme for Cyclopropanation Reactions. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 7785-7789.	13.8	98
2	Targeting STAT1 by myricetin and delphinidin provides efficient protection of the heart from ischemia/reperfusion-induced injury. <i>FEBS Letters</i> , 2009, 583, 531-541.	2.8	80
3	Validation of Model of Cytochrome P450 2D6: An in Silico Tool for Predicting Metabolism and Inhibition. <i>Journal of Medicinal Chemistry</i> , 2004, 47, 5340-5346.	6.4	78
4	How Do Azoles Inhibit Cytochrome P450 Enzymes? A Density Functional Study. <i>Journal of Physical Chemistry A</i> , 2008, 112, 12911-12918.	2.5	76
5	Structural, Kinetic, and Docking Studies of Artificial Imine Reductases Based on Biotin-Streptavidin Technology: An Induced Lock-and-Key Hypothesis. <i>Journal of the American Chemical Society</i> , 2014, 136, 15676-15683.	13.7	75
6	Incorporation of Manganese Complexes into Xylanase: New Artificial Metalloenzymes for Enantioselective Epoxidation. <i>ChemBioChem</i> , 2012, 13, 240-251.	2.6	72
7	Filling a Hole in Cytochrome P450 BM3 Improves Substrate Binding and Catalytic Efficiency. <i>Journal of Molecular Biology</i> , 2007, 373, 633-651.	4.2	71
8	Phe120 contributes to the regioselectivity of cytochrome P450 2D6: mutation leads to the formation of a novel dextromethorphan metabolite. <i>Biochemical Journal</i> , 2004, 380, 353-360.	3.7	69
9	Design of an enantioselective artificial metallo-hydratase enzyme containing an unnatural metal-binding amino acid. <i>Chemical Science</i> , 2017, 8, 7228-7235.	7.4	69
10	Selective oxidation of aromatic sulfide catalyzed by an artificial metalloenzyme: new activity of hemozymes. <i>Organic and Biomolecular Chemistry</i> , 2009, 7, 3208.	2.8	66
11	Why Is Quinidine an Inhibitor of Cytochrome P450 2D6?. <i>Journal of Biological Chemistry</i> , 2005, 280, 38617-38624.	3.4	63
12	Three Dimensional Models of Cu ²⁺ -Al ²⁺ (16) Complexes from Computational Approaches. <i>Journal of the American Chemical Society</i> , 2011, 133, 15008-15014.	13.7	61
13	Amine conformational change and spin conversion induced by metal-assisted ligand oxidation: from the seven-coordinate iron(II)-TPAA complex to the two oxidized iron(II)-(py) ₃ tren isomers. Characterization, crystal structures, and density functional study. <i>Inorganica Chimica Acta</i> , 2000, 297, 338-350.	2.4	59
14	Normal mode analysis as a prerequisite for drug design: Application to matrix metalloproteinases inhibitors. <i>FEBS Letters</i> , 2006, 580, 5130-5136.	2.8	58
15	IN SILICO AND IN VITRO SCREENING FOR INHIBITION OF CYTOCHROME P450 CYP3A4 BY COMEDICATIONS COMMONLY USED BY PATIENTS WITH CANCER. <i>Drug Metabolism and Disposition</i> , 2006, 34, 534-538.	3.3	58
16	Insight into the apoptosis-inducing action of \pm -bisabolol towards malignant tumor cells: Involvement of lipid rafts and Bid. <i>Archives of Biochemistry and Biophysics</i> , 2008, 476, 113-123.	3.0	57
17	From α -hemoabzymes to α -hemozymes towards new biocatalysts for selective oxidations. <i>Chemical Communications</i> , 2015, 51, 2476-2494.	4.1	54
18	Prediction of the interaction of metallic moieties with proteins: An update for protein-ligand docking techniques. <i>Journal of Computational Chemistry</i> , 2018, 39, 42-51.	3.3	54

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19	Toward the Computational Design of Artificial Metalloenzymes: From Proteinâ€™Ligand Docking to Multiscale Approaches. ACS Catalysis, 2015, 5, 2469-2480.	11.2	51
20	Multiple Substrate Binding by Cytochrome P450 3A4: Estimation of the Number of Bound Substrate Molecules. Drug Metabolism and Disposition, 2008, 36, 2136-2144.	3.3	50
21	Natural mutations of the anti-Mullerian hormone type II receptor found in persistent Mullerian duct syndrome affect ligand binding, signal transduction and cellular transport. Human Molecular Genetics, 2009, 18, 3002-3013.	2.9	49
22	Defective AMH signaling disrupts GnRH neuron development and function and contributes to hypogonadotropic hypogonadism. ELife, 2019, 8, .	6.0	49
23	Directed Evolution of an Artificial Imine Reductase. Angewandte Chemie - International Edition, 2018, 57, 1863-1868.	13.8	47
24	The design and synthesis of novel 3-[2-indol-1-yl-ethyl]-1H-indole derivatives as selective inhibitors of CDK4. Tetrahedron Letters, 2005, 46, 1423-1425.	1.4	44
25	Hemozymes Peroxidase Activity Of Artificial Hemoproteins Constructed From theStreptomyces lividansXylanase A and Iron(III)-Carboxy-Substituted Porphyrins. Bioconjugate Chemistry, 2008, 19, 899-910.	3.6	43
26	Design, synthesis and biological activity of new CDK4-specific inhibitors, based on faspaplysin. Organic and Biomolecular Chemistry, 2006, 4, 787.	2.8	42
27	Insights into drug metabolism by cytochromes P450 from modelling studies of CYP2D6â€™drug interactions. British Journal of Pharmacology, 2008, 153, S82-9.	5.4	42
28	IN SILICO PREDICTION OF DRUG BINDING TO CYP2D6: IDENTIFICATION OF A NEW METABOLITE OF METOCLOPRAMIDE. Drug Metabolism and Disposition, 2006, 34, 1386-1392.	3.3	41
29	Elucidation of Binding Site and Chiral Specificity of Oxidovanadium Drugs with Lysozyme through Theoretical Calculations. Inorganic Chemistry, 2017, 56, 12938-12951.	4.0	40
30	Microsolvation and Encapsulation Effects on Supramolecular Catalysis: Câ€™C Reductive Elimination inside [Ga₄L₆]^{12â€™} Metallo cage. Journal of the American Chemical Society, 2019, 141, 13114-13123.	13.7	40
31	Theoretical modeling of the heme group with a hybrid QM/MM method. Journal of Computational Chemistry, 2000, 21, 282-294.	3.3	39
32	Flexible N,N,N-chelates as supports for iron and cobalt chloride complexes; synthesis, structures, DFT calculations and ethylene oligomerisation studies. Dalton Transactions, 2004, , 3231-3240.	3.3	38
33	In Silico Analysis of the Apolipoprotein E and the Amyloid Î² Peptide Interaction: Misfolding Induced by Frustration of the Salt Bridge Network. PLoS Computational Biology, 2010, 6, e1000663.	3.2	38
34	GaudiMM: A modular multiâ€™objective platform for molecular modeling. Journal of Computational Chemistry, 2017, 38, 2118-2126.	3.3	37
35	Artificial Metalloenzymes with the Neocarzinostatin Scaffold: Toward a Biocatalyst for the Dielsâ€™Alder Reaction. ChemBioChem, 2016, 17, 433-440.	2.6	36
36	Validation and Applications of Proteinâ€™Ligand Docking Approaches Improved for Metalloligands with Multiple Vacant Sites. Inorganic Chemistry, 2019, 58, 294-306.	4.0	35

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37	Neocarzinostatin-based hybrid biocatalysts for oxidation reactions. Dalton Transactions, 2014, 43, 8344-8354.	3.3	32
38	Design and evolution of chimeric streptavidin for protein-enabled dual gold catalysis. Nature Catalysis, 2021, 4, 643-653.	34.4	32
39	Consensus modes, a robust description of protein collective motions from multiple-minima normal mode analysisâ€™ application to the HIV-1 protease. Physical Chemistry Chemical Physics, 2010, 12, 2850.	2.8	31
40	Factors affecting imine coordination in (iminoterpyridine)MX ₂ (M = Fe, Co, Ni, Zn): synthesis, structures, DFT calculations and ethylene oligomerisation studies. New Journal of Chemistry, 2007, 31, 75-85.	2.8	30
41	Insights into the Mechanism of Binding of Arachidonic Acid to Mammalian 15-Lipoxygenases. Journal of Physical Chemistry B, 2010, 114, 7037-7046.	2.6	30
42	Stereoselective Formation of Chiral Metallopeptides. Chemistry - A European Journal, 2012, 18, 7030-7035.	3.3	30
43	3D Structures and Redox Potentials of Cu ²⁺ â€™Al ²⁺ (1â€™16) Complexes at Different pH: A Computational Study. Journal of Physical Chemistry B, 2014, 118, 4840-4850.	2.6	30
44	Programmed stereoselective assembly of DNA-binding helical metallopeptides. Chemical Communications, 2014, 50, 11097-11100.	4.1	30
45	DFT Protocol for EPR Prediction of Paramagnetic Cu(II) Complexes and Application to Protein Binding Sites. Magnetochemistry, 2018, 4, 55.	2.4	30
46	Copper(II) <i>N</i>,<i>N</i>,<i>O</i>-Chelating Complexes as Potential Anticancer Agents. Inorganic Chemistry, 2021, 60, 2939-2952.	4.0	30
47	New fascaplysin-based CDK4-specific inhibitors: design, synthesis and biological activity. Chemical Communications, 2004, , 1696-1697.	4.1	29
48	Mono- vs. bi-metallic assembly on a bulky bis(imino)terpyridine framework: a combined experimental and theoretical study. Dalton Transactions, 2006, , 2350-2361.	3.3	29
49	Molecular Modeling for Artificial Metalloenzyme Design and Optimization. Accounts of Chemical Research, 2020, 53, 896-905.	15.6	29
50	Dynamic Stereoselection of Peptide Helicates and Their Selective Labeling of DNA Replication Foci in Cells**. Angewandte Chemie - International Edition, 2021, 60, 8859-8866.	13.8	29
51	Decoding Surface Interaction of V ^{IV} Metallodrug Candidates with Lysozyme. Inorganic Chemistry, 2018, 57, 4456-4469.	4.0	28
52	Interaction of Vanadium(IV) Species with Ubiquitin: A Combined Instrumental and Computational Approach. Inorganic Chemistry, 2019, 58, 8064-8078.	4.0	28
53	CA224, a non-planar analogue of fascaplysin, inhibits Cdk4 but not Cdk2 and arrests cells at G0/G1 inhibiting pRB phosphorylation. Bioorganic and Medicinal Chemistry Letters, 2006, 16, 4272-4278.	2.2	27
54	Computational Insights on an Artificial Imine Reductase Based on the Biotinâ€™Streptavidin Technology. ACS Catalysis, 2014, 4, 833-842.	11.2	27

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55	Quantitative prediction of electronic absorption spectra of copper(II)â€™bioligand systems: Validation and applications. <i>Journal of Inorganic Biochemistry</i> , 2020, 204, 110953.	3.5	27
56	Internal cavities and ligand passageways in human hemoglobin characterized by molecular dynamics simulations. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2005, 1724, 385-393.	2.4	26
57	Progress in cytochrome P450 active site modeling. <i>Archives of Biochemistry and Biophysics</i> , 2005, 433, 361-368.	3.0	26
58	What can molecular modelling bring to the design of artificial inorganic cofactors?. <i>Faraday Discussions</i> , 2011, 148, 137-159.	3.2	26
59	An Artificial Heme Enzyme for Cyclopropanation Reactions. <i>Angewandte Chemie</i> , 2018, 130, 7911-7915.	2.0	26
60	BioMetAll: Identifying Metal-Binding Sites in Proteins from Backbone Preorganization. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 311-323.	5.4	25
61	Integrated ESI-MS/EPR/computational characterization of the binding of metal species to proteins: vanadium drugâ€™myoglobin application. <i>Inorganic Chemistry Frontiers</i> , 2019, 6, 1561-1578.	6.0	24
62	Cofactor Binding Dynamics Influence the Catalytic Activity and Selectivity of an Artificial Metalloenzyme. <i>ACS Catalysis</i> , 2020, 10, 11783-11790.	11.2	24
63	Integrated experimental/computational approaches to characterize the systems formed by vanadium with proteins and enzymes. <i>Inorganic Chemistry Frontiers</i> , 2021, 8, 1951-1974.	6.0	24
64	Unveiling V^{IV}O²⁺ Binding Modes to Human Serum Albumins by an Integrated Spectroscopicâ€™Computational Approach. <i>Chemistry - A European Journal</i> , 2020, 26, 11316-11326.	3.3	23
65	Assessing protein-ligand docking for the binding of organometallic compounds to proteins. <i>Journal of Computational Chemistry</i> , 2014, 35, 192-198.	3.3	22
66	Computational insight into the catalytic implication of head/tail-first orientation of arachidonic acid in human 5-lipoxygenase: consequences for the positional specificity of oxygenation. <i>Physical Chemistry Chemical Physics</i> , 2016, 18, 23017-23035.	2.8	22
67	Effect of secondary interactions, steric hindrance and electric charge on the interaction of V^{IV}O species with proteins. <i>New Journal of Chemistry</i> , 2019, 43, 17647-17660.	2.8	22
68	Elucidating the 3D structures of Al(<i>scp>iii</scp></i>)â€™Al ² complexes: a template free strategy based on the pre-organization hypothesis. <i>Chemical Science</i> , 2017, 8, 5041-5049.	7.4	21
69	Functionalized Artificial Bidomain Proteins Based on an Î±-Solenoid Protein Repeat Scaffold: A New Class of Artificial Dielsâ€™Alderses. <i>ACS Omega</i> , 2019, 4, 4437-4447.	3.5	21
70	Investigating Potential Inhibitory Effect of <i>Uncaria tomentosa</i> (Catâ€™s Claw) against the Main Protease 3CLpro of SARS-CoV-2 by Molecular Modeling. <i>Evidence-based Complementary and Alternative Medicine</i> , 2020, 2020, 1-14.	1.2	21
71	Discovery of processive catalysis by an exo-hydrolase with a pocket-shaped active site. <i>Nature Communications</i> , 2019, 10, 2222.	12.8	20
72	Simple Coordination Geometry Descriptors Allow to Accurately Predict Metal-Binding Sites in Proteins. <i>ACS Omega</i> , 2019, 4, 3726-3731.	3.5	20

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73	ESigen: Electronic Supporting Information Generator for Computational Chemistry Publications. <i>Journal of Chemical Information and Modeling</i> , 2018, 58, 561-564.	5.4	19
74	Inhibition of cancer cell growth by cyclin dependent kinase 4 inhibitors synthesized based on the structure of foscarnin. <i>Bioorganic Chemistry</i> , 2006, 34, 287-297.	4.1	18
75	PyChimera: use UCSF Chimera modules in any Python 2.7 project. <i>Bioinformatics</i> , 2018, 34, 1784-1785.	4.1	18
76	Reaction Rate Inside the Cavity of [Ga ₄ L ₆] ¹²⁺ Supramolecular Metallogage is Regulated by the Encapsulated Solvent. <i>Chemistry - A European Journal</i> , 2020, 26, 6988-6992.	3.3	18
77	Crystal structure of c5321: a protective antigen present in uropathogenic <i>Escherichia coli</i> strains displaying an SLR fold. <i>BMC Structural Biology</i> , 2013, 13, 19.	2.3	17
78	Modeling Cu ²⁺ -A ²⁺ complexes from computational approaches. <i>AIP Advances</i> , 2015, 5, 092402.	1.3	17
79	Accurate prediction of vertical electronic transitions of Ni(II) coordination compounds via time dependent density functional theory. <i>International Journal of Quantum Chemistry</i> , 2018, 118, e25655.	2.0	16
80	Monitoring lactoferrin iron levels by fluorescence resonance energy transfer: a combined chemical and computational study. <i>Journal of Biological Inorganic Chemistry</i> , 2014, 19, 439-447.	2.6	15
81	Differential response to Cadmium exposure by expression of a two and a three-domain metallothionein isoform in the land snail <i>Pomatias elegans</i> : Valuating the marine heritage of a land snail. <i>Science of the Total Environment</i> , 2019, 648, 561-571.	8.0	15
82	Stepwise Oxidations in a Cofacial Copper(II) Porphyrin Dimer: Through-Space Spin-Coupling and Interplay between Metal and Radical Spins. <i>Chemistry - A European Journal</i> , 2020, 26, 7869-7880.	3.3	15
83	Direct interaction of natural and synthetic catechins with signal transducer activator of transcription 1 affects both its phosphorylation and activity. <i>FEBS Journal</i> , 2014, 281, 724-738.	4.7	14
84	GPathFinder: Identification of Ligand-Binding Pathways by a Multi-Objective Genetic Algorithm. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3155.	4.1	14
85	Use of normal modes for structural modeling of proteins: the case study of rat heme oxygenase 1. <i>European Biophysics Journal</i> , 2008, 37, 1157-1165.	2.2	11
86	Crystal Structure of Two Anti-Porphyrin Antibodies with Peroxidase Activity. <i>PLoS ONE</i> , 2012, 7, e51128.	2.5	11
87	Synthesis of Novel Nucleoside Analogues Built on a Bicyclo[4.1.0]heptane Scaffold. <i>Journal of Organic Chemistry</i> , 2015, 80, 9495-9505.	3.2	11
88	Directed Self-Assembly of Trimeric DNA-Binding chiral Mini-protein Helicates. <i>Frontiers in Chemistry</i> , 2018, 6, 520.	3.6	11
89	Studying the reactivity of α -old-Cu(II) complexes for α -novel-anticancer purposes. <i>Journal of Inorganic Biochemistry</i> , 2019, 195, 51-60.	3.5	11
90	Biospeciation of Potential Vanadium Drugs of Acetylacetonate in the Presence of Proteins. <i>Frontiers in Chemistry</i> , 2020, 8, 345.	3.6	11

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91	Ab initio calculations predict a very low barrier for the rotation of the axial ligand in Fe(P)(Im). Chemical Physics Letters, 2002, 353, 379-382.	2.6	10
92	A DFT study on the relative affinity for oxygen of the $\hat{1}\alpha$ and $\hat{1}\beta$ subunits of hemoglobin. Journal of Computational Chemistry, 2006, 27, 1446-1453.	3.3	10
93	Unravelling novel synergies between organometallic and biological partners: a quantum mechanics/molecular mechanics study of an artificial metalloenzyme. Journal of the Royal Society Interface, 2014, 11, 20140090.	3.4	10
94	Neocarzinostatin-based hybrid biocatalysts with a RNase like activity. Bioorganic and Medicinal Chemistry, 2014, 22, 5678-5686.	3.0	10
95	New Insights into the Reactivity of Cisplatin with Free and Restrained Nucleophiles: Microsolvation Effects and Base Selectivity in Cisplatin-DNA Interactions. ChemPhysChem, 2016, 17, 3932-3947.	2.1	10
96	Computational insight into the interaction of oxaliplatin with insulin. Metallomics, 2019, 11, 765-773.	2.4	10
97	Integrated Computational Study of the Cu-Catalyzed Hydration of Alkenes in Water Solvent and into the Context of an Artificial Metallohydratase. ACS Catalysis, 2019, 9, 4616-4626.	11.2	10
98	Modeling Kinetics and Thermodynamics of Guest Encapsulation into the [M ₄ L ₆]12-Supramolecular Organometallic Cage. Journal of Chemical Information and Modeling, 2021, 61, 4370-4381.	5.4	10
99	The folding of a metallopeptide. Dalton Transactions, 2016, 45, 881-885.	3.3	9
100	The Effect of Cofactor Binding on the Conformational Plasticity of the Biological Receptors in Artificial Metalloenzymes: The Case Study of LmrR. Frontiers in Chemistry, 2019, 7, 211.	3.6	9
101	An Artificial Hemoprotein with Inducible Peroxidase- and Monooxygenase-Like Activities. Chemistry - A European Journal, 2020, 26, 14929-14937.	3.3	9
102	Origin of the Rate Acceleration in the C ¹³ Reductive Elimination from Pt(IV)-complex in a [Ga ₄ L ₆] ¹²⁺ Supramolecular Metallocage. Chemistry - A European Journal, 2021, 27, 15973-15980.	3.3	9
103	Understanding the interaction of an antitumoral platinum(II) 7-azaindolate complex with proteins and DNA. BioMetals, 2014, 27, 1159-1177.	4.1	8
104	Directed Evolution of an Artificial Imine Reductase. Angewandte Chemie, 2018, 130, 1881-1886.	2.0	8
105	Synthesis, Anti-HIV Activity Studies, and in-silico Rationalization of Cyclobutane-Fused Nucleosides. ChemMedChem, 2012, 7, 1044-1056.	3.2	7
106	Synthesis, Antiviral Evaluation, and Computational Studies of Cyclobutane and Cyclobutene Nucleoside Analogues. European Journal of Organic Chemistry, 2013, 2013, 7761-7775.	2.4	7
107	Various strategies for obtaining oxidative artificial hemoproteins with a catalytic oxidative activity: from "Hemoabzymes" to "Hemozymes"?. Journal of Porphyrins and Phthalocyanines, 2014, 18, 1063-1092.	0.8	7
108	Dynamic Stereoselection of Peptide Helicates and Their Selective Labeling of DNA Replication Foci in Cells**. Angewandte Chemie, 2021, 133, 8941-8948.	2.0	7

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109	Inhibition of the Human Hsc70 System by Small Ligands as a Potential Anticancer Approach. <i>Cancers</i> , 2021, 13, 2936.	3.7	7
110	Impact of Cu(II) and Al(III) on the conformational landscape of amyloid β_{1-42} . <i>Physical Chemistry Chemical Physics</i> , 2021, 23, 13023-13032.	2.8	7
111	Insights into Drug Metabolism from Modelling Studies of Cytochrome P450-Drug Interactions. <i>Current Topics in Medicinal Chemistry</i> , 2006, 6, 1619-1626.	2.1	6
112	GARLEEK: Adding an extra flavor to ONIOM. <i>Journal of Computational Chemistry</i> , 2019, 40, 381-386.	3.3	6
113	Hybrid Cyclobutane/Proline-Containing Peptidomimetics: The Conformational Constraint Influences Their Cell-Penetration Ability. <i>International Journal of Molecular Sciences</i> , 2021, 22, 5092.	4.1	5
114	Selective recognition of A/T-rich DNA 3-way junctions with a three-fold symmetric tripeptide. <i>Chemical Communications</i> , 0, , .	4.1	5
115	Computational insights on the possibility of tri-coordinated cisplatinated adducts with protein models. <i>Journal of Inorganic Biochemistry</i> , 2012, 117, 230-236.	3.5	4
116	Metal Complexation of a D-Ribose-Based Ligand Decoded by Experimental and Theoretical Studies. <i>European Journal of Inorganic Chemistry</i> , 2012, 2012, 3308-3319.	2.0	4
117	Chiral Cyclobutane-Containing Cell-Penetrating Peptides as Selective Vectors for Anti-Leishmania Drug Delivery Systems. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7502.	4.1	4
118	Bioelectrochemical monitoring of soluble guanylate cyclase inhibition by the natural β -carboline canthin-6-one. <i>Journal of Molecular Structure</i> , 2017, 1134, 661-667.	3.6	3
119	Stereoselective Self-Assembly of DNA Binding Helicates Directed by the Viral β -Annulus Trimeric Peptide Motif. <i>Bioconjugate Chemistry</i> , 2021, 32, 1564-1569.	3.6	3
120	Receptor Flexibility in Ligand Docking and Virtual Screening. , 2011, , 99-117.		2
121	Theoretical modeling of the heme group with a hybrid QM/MM method. <i>Journal of Computational Chemistry</i> , 2000, 21, 282.	3.3	1
122	Chapter 15. Enzyme Design. <i>RSC Theoretical and Computational Chemistry Series</i> , 2016, , 481-521.	0.7	1
123	Successes and Challenges in Multiscale Modelling of Artificial Metalloenzymes: the Case Study of POP-Rh2 Cyclopropanase. <i>Faraday Discussions</i> , 2022, , .	3.2	1
124	Controlling oncogenic KRAS signaling pathways with a Palladium-responsive peptide. <i>Communications Chemistry</i> , 2022, 5, .	4.5	1
125	The Design and Synthesis of Novel 3-[2-Indol-1-yl-ethyl]-1H-indole Derivatives as Selective Inhibitors of CDK4.. <i>ChemInform</i> , 2005, 36, no.	0.0	0
126	Computational simulations of copper complexes relevant to Alzheimer's disease. , 2014, , .		0

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127	Influence of Association on Binding of Disaccharides to YKL-39 and hHyal-1 Enzymes. International Journal of Molecular Sciences, 2022, 23, 7705.	4.1	0