

Carlos A Montanari

List of Publications by Year in descending order

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122
papers

2,027
citations

218381

26
h-index

329751

37
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130
all docs

130
docs citations

130
times ranked

2542
citing authors

#	ARTICLE	IF	CITATIONS
1	C�ncer e agentes antineopl�sicos ciclo-celular espec�ficos e ciclo-celular n�o espec�ficos que interagem com o DNA: uma introdu�o. <i>Quimica Nova</i> , 2005, 28, 118-129.	0.3	87
2	Inflation of correlation in the pursuit of drug-likeness. <i>Journal of Computer-Aided Molecular Design</i> , 2013, 27, 1-13.	1.3	72
3	Hologram QSAR model for the prediction of human oral bioavailability. <i>Bioorganic and Medicinal Chemistry</i> , 2007, 15, 7738-7745.	1.4	71
4	Discovery of novel <i>Trypanosoma cruzi</i> glyceraldehyde-3-phosphate dehydrogenase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2009, 17, 2476-2482.	1.4	67
5	Non-peptidic Cruzain Inhibitors with Trypanocidal Activity Discovered by Virtual Screening and In Vitro Assay. <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2370.	1.3	63
6	Ligand efficiency metrics considered harmful. <i>Journal of Computer-Aided Molecular Design</i> , 2014, 28, 699-710.	1.3	54
7	Two- and three-dimensional quantitative structure-activity relationships for a series of purine nucleoside phosphorylase inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 516-527.	1.4	50
8	Molecular Design, Synthesis and Trypanocidal Activity of Dipeptidyl Nitriles as Cruzain Inhibitors. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003916.	1.3	49
9	Hydrogen Bond Basicity Prediction for Medicinal Chemistry Design. <i>Journal of Medicinal Chemistry</i> , 2016, 59, 4278-4288.	2.9	48
10	Structural Basis for Selective Inhibition of Trypanosomatid Glyceraldehyde-3-Phosphate Dehydrogenase: Molecular Docking and 3D QSAR Studies. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 918-929.	2.5	47
11	Novel insights for dihydroorotate dehydrogenase class 1A inhibitors discovery. <i>European Journal of Medicinal Chemistry</i> , 2010, 45, 5899-5909.	2.6	47
12	Can Cysteine Protease Cross-Class Inhibitors Achieve Selectivity?. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 10497-10525.	2.9	47
13	Effects of organic solvents on the enzyme activity of <i>Trypanosoma cruzi</i> glyceraldehyde-3-phosphate dehydrogenase in calorimetric assays. <i>Analytical Biochemistry</i> , 2007, 370, 107-114.	1.1	45
14	Sele�o de vari�veis em QSAR. <i>Quimica Nova</i> , 2002, 25, 439-448.	0.3	43
15	A QSERR study on enantioselective separation of enantiomeric sulphoxides. <i>Analytica Chimica Acta</i> , 2000, 419, 93-100.	2.6	41
16	Conformational differences between the wild type and V30M mutant transthyretin modulate its binding to genistein: Implications to tetramer stability and ligand-binding. <i>Journal of Structural Biology</i> , 2010, 170, 522-531.	1.3	39
17	Experimental study and computational modelling of cruzain cysteine protease inhibition by dipeptidyl nitriles. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 24317-24328.	1.3	38
18	3D QSAR studies on binding affinities of coumarin natural products for glycosomal GAPDH of <i>Trypanosoma cruzi</i> . <i>Journal of Computer-Aided Molecular Design</i> , 2003, 17, 277-290.	1.3	35

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19	Structure-Activity Relationships for the Design of Small-Molecule Inhibitors. <i>Mini-Reviews in Medicinal Chemistry</i> , 2005, 5, 585-593.	1.1	33
20	Pharmacophore Modeling for Anti-Chagas Drug Design Using the Fragment Molecular Orbital Method. <i>PLoS ONE</i> , 2015, 10, e0125829.	1.1	33
21	A comparative study of warheads for design of cysteine protease inhibitors. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2017, 27, 5031-5035.	1.0	32
22	Structure-activity relationships of novel inhibitors of glyceraldehyde-3-phosphate dehydrogenase. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2004, 14, 2199-2204.	1.0	30
23	ClogPalk: a method for predicting alkane/water partition coefficient. <i>Journal of Computer-Aided Molecular Design</i> , 2013, 27, 389-402.	1.3	30
24	Planejamento racional de fármacos baseado em produtos naturais. <i>Quimica Nova</i> , 2001, 24, .	0.3	30
25	2D QSAR and similarity studies on cruzain inhibitors aimed at improving selectivity over cathepsin L. <i>Bioorganic and Medicinal Chemistry</i> , 2008, 16, 838-853.	1.4	28
26	Assessment of the Cruzain Cysteine Protease Reversible and Irreversible Covalent Inhibition Mechanism. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 1666-1677.	2.5	26
27	Anti-trypanosomal activity of non-peptidic nitrile-based cysteine protease inhibitors. <i>PLoS Neglected Tropical Diseases</i> , 2017, 11, e0005343.	1.3	26
28	Determination of receptor-bound drug conformations by QSAR using flexible fitting to derive a molecular similarity index. <i>Journal of Computer-Aided Molecular Design</i> , 1996, 10, 67-73.	1.3	24
29	HEPT derivatives as non-nucleoside inhibitors of HIV-1 reverse transcriptase: QSAR studies agree with the crystal structures. <i>Journal of Computer-Aided Molecular Design</i> , 2002, 16, 287-295.	1.3	24
30	Multimilligram enantioresolution of sulfoxide proton pump inhibitors by liquid chromatography on polysaccharide-based chiral stationary phase. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2008, 47, 81-87.	1.4	23
31	Integration of Ligand- and Target-Based Virtual Screening for the Discovery of Cruzain Inhibitors. <i>Molecular Informatics</i> , 2011, 30, 565-578.	1.4	23
32	New Class of Antitrypanosomal Agents Based on Imidazopyridines. <i>ACS Medicinal Chemistry Letters</i> , 2017, 8, 766-770.	1.3	21
33	QSAR and molecular modelling studies on B-DNA recognition of minor groove binders. <i>European Journal of Medicinal Chemistry</i> , 2003, 38, 141-155.	2.6	20
34	In Silico Prediction of Human Plasma Protein Binding Using Hologram QSAR. <i>Letters in Drug Design and Discovery</i> , 2007, 4, 502-509.	0.4	20
35	Leveraging the cruzain S3 subsite to increase affinity for reversible covalent inhibitors. <i>Bioorganic Chemistry</i> , 2018, 79, 285-292.	2.0	20
36	The influence of hydrogen bonding on partition coefficients. <i>Journal of Computer-Aided Molecular Design</i> , 2017, 31, 163-181.	1.3	19

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37	Evaluating QM/MM Free Energy Surfaces for Ranking Cysteine Protease Covalent Inhibitors. Journal of Chemical Information and Modeling, 2020, 60, 880-889.	2.5	19
38	Calorimetry and structure-activity relationships for a series of antimicrobial hydrazides. Thermochimica Acta, 2004, 417, 283-294.	1.2	18
39	Synthesis and structure-activity relationship of nitrile-based cruzain inhibitors incorporating a trifluoroethylamine-based P2 amide replacement. Bioorganic and Medicinal Chemistry, 2019, 27, 115083.	1.4	18
40	Structural studies on some 1,3,4-thiadiazolium-2-aminides and their rearrangement isomers using ¹⁵ N and ¹³ C NMR spectroscopy. Journal of the Chemical Society Perkin Transactions II, 1994, , 2571.	0.9	17
41	Targeting the Shc-EGFR interaction with indomethacin inhibits MAP kinase pathway signalling. Cancer Letters, 2019, 457, 86-97.	3.2	17
42	QSAR based on biological microcalorimetry. Thermochimica Acta, 1999, 328, 91-97.	1.2	16
43	Organometallic Gold(III) Complex [Au(Hdamp)(L1 ₄)] ⁺ (L1 = <i>SNS</i> -Donating) Tj ETQq1 1 0.784314 rg 5T Diseases, 2019, 5, 1698-1707.	1.8	16
44	Predicting the affinity of halogenated reversible covalent inhibitors through relative binding free energy. Physical Chemistry Chemical Physics, 2019, 21, 24723-24730.	1.3	16
45	The role of molecular interaction fields on enantioselective and nonselective separation of chiral sulfoxides. Journal of Chromatography A, 2006, 1121, 64-75.	1.8	15
46	Novel Application of 2D and 3D-Similarity Searches To Identify Substrates among Cytochrome P450 2C9, 2D6, and 3A4. Journal of Chemical Information and Modeling, 2010, 50, 97-109.	2.5	15
47	Molecular design, synthesis and biological evaluation of 1,4-dihydro-4-oxoquinoline ribonucleosides as TcGAPDH inhibitors with trypanocidal activity. Bioorganic and Medicinal Chemistry Letters, 2013, 23, 4597-4601.	1.0	15
48	A neural networks study of quinone compounds with trypanocidal activity. Journal of Molecular Modeling, 2008, 14, 975-985.	0.8	14
49	Kinetic mechanism and catalysis of Trypanosoma cruzi dihydroorotate dehydrogenase enzyme evaluated by isothermal titration calorimetry. Analytical Biochemistry, 2010, 399, 13-22.	1.1	14
50	Estimation of the RP-HPLC Lipophilicity Parameters Log K', and Log K _W , A Comparison with the Hydrophobicity Index $\log P$. Journal of Liquid Chromatography and Related Technologies, 1997, 20, 1703-1715.	0.5	13
51	Similarity search combined with docking and molecular dynamics for novel hAChE inhibitor scaffolds. Journal of Molecular Modeling, 2018, 24, 41.	0.8	13
52	Discovery of small molecule inhibitors of <i>Leishmania braziliensis</i> Hsp90 chaperone. Journal of Enzyme Inhibition and Medicinal Chemistry, 2020, 35, 639-649.	2.5	13
53	QSAR Based on Biological Microcalorimetry. Journal of Medicinal Chemistry, 2000, 43, 3448-3452.	2.9	12
54	Identification of chiral selectors for improved enantioseparation based on molecular interaction fields. Analytica Chimica Acta, 2005, 545, 33-45.	2.6	12

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55	Docking and molecular dynamics simulation of quinone compounds with trypanocidal activity. <i>Journal of Molecular Modeling</i> , 2009, 15, 1175-1184.	0.8	12
56	Expression, purification and kinetic characterization of His-tagged glyceraldehyde-3-phosphate dehydrogenase from <i>Trypanosoma cruzi</i> . <i>Protein Expression and Purification</i> , 2011, 76, 190-196.	0.6	12
57	The Open Form Inducer Approach for Structure-Based Drug Design. <i>PLoS ONE</i> , 2016, 11, e0167078.	1.1	12
58	Quantitative structure-retention relationships of antimicrobial hydrazides evaluated by reverse-phase liquid chromatography. <i>Chromatographia</i> , 2000, 51, 722-726.	0.7	11
59	Validação lateral em relações quantitativas entre estrutura e atividade farmacológica, QSAR. <i>Quimica Nova</i> , 2002, 25, 231-240.	0.3	11
60	Mapping the S1 and S1 TM subsites of cysteine proteases with new dipeptidyl nitrile inhibitors as trypanocidal agents. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0007755.	1.3	11
61	Nitrile-based peptoids as cysteine protease inhibitors. <i>Bioorganic and Medicinal Chemistry</i> , 2021, 41, 116211.	1.4	11
62	DETERMINATION OF PARTITION COEFFICIENTS FOR SEVERAL PROPRANOLOL ANALOGUES BY REVERSED PHASE HIGH PERFORMANCE LIQUID CHROMATOGRAPHY. <i>Journal of Liquid Chromatography and Related Technologies</i> , 1999, 22, 2139-2149.	0.5	10
63	Synthesis of Mesoionic 4-(para-substituted Phenyl)-2,4-dichlorophenyl-1,3,4-thiadiazolium-2-amminides by Direct Cyclization via Acylation of Thiosemicarbazides. <i>Synthetic Communications</i> , 2006, 36, 3359-3369.	1.1	10
64	Automated molecule editing in molecular design. <i>Journal of Computer-Aided Molecular Design</i> , 2013, 27, 655-664.	1.3	10
65	Integration of methods in cheminformatics and biocalorimetry for the design of trypanosomatid enzyme inhibitors. <i>Future Medicinal Chemistry</i> , 2014, 6, 17-33.	1.1	10
66	Quantitative structure-retention relationships of flavonoids unraveled by immobilized artificial membrane chromatography. <i>European Journal of Pharmaceutical Sciences</i> , 2016, 88, 147-157.	1.9	10
67	Determination of Log Papp for Drug Transfer to Microbial Cellular Systems via the Taylor-Aris Modified Diffusion Technique. <i>QSAR and Combinatorial Science</i> , 1998, 17, 102-108.	1.4	10
68	Predicting the Relative Binding Affinity for Reversible Covalent Inhibitors by Free Energy Perturbation Calculations. <i>Journal of Chemical Information and Modeling</i> , 2021, 61, 4733-4744.	2.5	9
69	Three-dimensional models of non-steroidal ligands: A comparative molecular field analysis. <i>Steroids</i> , 2006, 71, 417-428.	0.8	8
70	Synthesis, biochemical evaluation and molecular modeling studies of nonpeptidic nitrile-based fluorinated compounds. <i>Future Medicinal Chemistry</i> , 2021, 13, 25-43.	1.1	8
71	Structure Based Design, Synthesis, and Evaluation of Potential Inhibitors of Steroid Sulfatase. <i>Current Topics in Medicinal Chemistry</i> , 2014, 14, 1033-1044.	1.0	8
72	The Molecular Retention Mechanism in Reversed-Phase Liquid Chromatography of Meso-ionic Compounds by Quantitative Structure-Retention Relationships (QSRR). <i>Chemistry and Biodiversity</i> , 2005, 2, 1691-1700.	1.0	7

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73	In silico screening of HIV-1 non-nucleoside reverse transcriptase and protease inhibitors. <i>European Journal of Medicinal Chemistry</i> , 2008, 43, 1412-1422.	2.6	7
74	A new bianthron glycoside as inhibitor of <i>Trypanosoma cruzi</i> glyceraldehyde 3-phosphate dehydrogenase activity. <i>Journal of the Brazilian Chemical Society</i> , 2009, 20, 947-953.	0.6	7
75	On the intrinsic reactivity of highly potent trypanocidal cruzain inhibitors. <i>RSC Medicinal Chemistry</i> , 2020, 11, 1275-1284.	1.7	7
76	Optimization strategy of single-digit nanomolar cross-class inhibitors of mammalian and protozoa cysteine proteases. <i>Bioorganic Chemistry</i> , 2020, 101, 104039.	2.0	7
77	Structural, thermodynamic and functional studies of human 71 kDa heat shock cognate protein (HSPA8/hHsc70). <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2021, 1869, 140719.	1.1	7
78	Molecular Recognition of B-DNA Minor-Groove Binders: The Rigid Analogue Approach to Synthesis of Antileishmaniasis Compounds: A Molecular Modeling Study. <i>Journal of the Brazilian Chemical Society</i> , 1998, 9, 175-180.	0.6	6
79	ON THE PARTITIONING OF SOME NEWLY SYNTHESIZED MESOIONIC 1,3,4-THIADIAZOLIUM-2-AMINIDE AND PRECURSORS EVALUATED BY RP-HPLC. <i>Journal of Liquid Chromatography and Related Technologies</i> , 1999, 22, 357-366.	0.5	6
80	Evaluation of the leishmanicidal and cytotoxic effects of inhibitors for microorganism metabolic pathway enzymes. <i>Biomedicine and Pharmacotherapy</i> , 2015, 74, 95-100.	2.5	6
81	Novel anti-prostate cancer scaffold identified by the combination of in silico and cell-based assays targeting the PI3K-AKT-mTOR pathway. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2017, 27, 4001-4006.	1.0	6
82	Ligand-induced conformational selection predicts the selectivity of cysteine protease inhibitors. <i>PLoS ONE</i> , 2019, 14, e0222055.	1.1	6
83	Crystal structure of <i>Leishmania mexicana</i> cysteine protease B in complex with a high-affinity azadipeptide nitrile inhibitor. <i>Bioorganic and Medicinal Chemistry</i> , 2020, 28, 115743.	1.4	6
84	Design, synthesis and stepwise optimization of nitrile-based inhibitors of cathepsins B and L. <i>Bioorganic and Medicinal Chemistry</i> , 2021, 29, 115827.	1.4	6
85	Biological Activity and Physicochemical Properties of Dipeptidyl Nitrile Derivatives Against Pancreatic Ductal Adenocarcinoma Cells. <i>Anti-Cancer Agents in Medicinal Chemistry</i> , 2019, 19, 112-120.	0.9	6
86	The GRID/CPCA approach in drug discovery. <i>Expert Opinion on Drug Discovery</i> , 2010, 5, 333-346.	2.5	5
87	3D-WHIM pattern recognition study for bisamidines. A structure-property relationship study. <i>Journal of the Brazilian Chemical Society</i> , 2000, 11, 393-397.	0.6	5
88	Bioactivity of a series of substituted N(3-phenyl-2-propenylidene)benzeneamines: a microcalorimetric study. <i>International Journal of Pharmaceutics</i> , 1992, 85, 199-202.	2.6	4
89	On Flexible Fitting Using The Molecular Similarity Index. A neural network analysis for antileishmaniasis compounds. <i>QSAR and Combinatorial Science</i> , 1997, 16, 480-481.	1.4	4
90	Sistemas transportadores de drogas. <i>Quimica Nova</i> , 1998, 21, 470-476.	0.3	4

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91	Sobre o uso de métodos químicos em química combinatória. <i>Química Nova</i> , 2000, 23, 178-184.	0.3	4
92	Hologram Quantitative Structure-Activity Relationships for a Class of Inhibitors of HIV-1 Protease. <i>Letters in Drug Design and Discovery</i> , 2007, 4, 356-364.	0.4	4
93	Comparative Molecular Field Analysis of a Series of Inhibitors of HIV-1 Protease. <i>Medicinal Chemistry</i> , 2011, 7, 71-79.	0.7	4
94	In silico selection and cell-based characterization of selective and bioactive compounds for androgen-dependent prostate cancer cell. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2017, 27, 546-550.	1.0	4
95	Highly predictive hologram QSAR models of nitrile-containing cruzain inhibitors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 3232-3249.	2.0	4
96	N-Sulfonyl dipeptide nitriles as inhibitors of human cathepsin S: In silico design, synthesis and biochemical characterization. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2020, 30, 127420.	1.0	4
97	Química Medicinal: 25 anos de planejamento racional de fármacos. <i>Química Nova</i> , 0, 25, 39-44.	0.3	4
98	A patent review on cathepsin K inhibitors to treat osteoporosis (2011 – 2021). <i>Expert Opinion on Therapeutic Patents</i> , 2022, 32, 561-573.	2.4	4
99	A Combined Study Using Ligand-Based Design, Synthesis, and Pharmacological Evaluation of Analogues of the Acetaminophen <i>ortho</i> Regioisomer with Potent Analgesic Activity. <i>Chemical Biology and Drug Design</i> , 2012, 80, 99-105.	1.5	3
100	Dipeptidyl nitrile derivatives have cytostatic effects against <i>Leishmania</i> spp. promastigotes. <i>Experimental Parasitology</i> , 2019, 200, 84-91.	0.5	3
101	Dipeptidyl nitrile derivatives suppress the <i>Trypanosoma cruzi</i> in vitro infection. <i>Experimental Parasitology</i> , 2020, 219, 108032.	0.5	3
102	A química medicinal na próxima década. <i>Química Nova</i> , 2000, 23, 134-137.	0.3	2
103	The role of ion pairing in the chromatographic study of propranolol analogues. <i>Chromatographia</i> , 2001, 53, 11-16.	0.7	2
104	Estudos de relações estrutura-atividade quantitativas (QSAR) de bis-benzamidas com atividade antifúngica. <i>Química Nova</i> , 2010, 33, 1482-1489.	0.3	2
105	Synthesis and matched molecular pair analysis of covalent reversible inhibitors of the cysteine protease CPB. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2020, 30, 127439.	1.0	2
106	Estudos de QSAR baseados em dados de atividade biológica obtidos por microcalorimetria: III interação de m-alcofifenis e p-hidroxibenzoatos de alquila com <i>Escherichia coli</i> . <i>Química Nova</i> , 1997, 20, 125-131.	0.3	2
107	A QSAR Study on <i>Pneumocystis carinii</i> Topoisomerases of bis-Benzimidazoles. <i>QSAR and Combinatorial Science</i> , 2000, 19, 173-175.	1.4	1
108	Interação de compostos organossulfurados derivados do alho com o citocromo-c: um estudo eletroquímico. <i>Química Nova</i> , 2002, 25, 5-9.	0.3	1

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109	Um novo modelo para os sítios de interação dos antagonistas H2, baseado na caracterização química dos sítios primário e secundário por QSAR- 3D. Química Nova, 2003, 26, 499-506.	0.3	1
110	Preparation and Chemical and Biological Studies of the Novel Complex $[\frac{1}{4}\text{-}1,4,5\text{-triphenyl-}1,3,4\text{-triazole-}2\text{-thiolate-bis}[\text{dichlorotriethylphosphineplatinum(II)}]_a$. Transition Metal Chemistry, 2005, 30, 733-737.	0.7	1
111	Chemometric Characterization of Chromatographic Retention Parameters of Mesoionic 1,3,4-thiadiazolium β -Aminides by Molecular Interaction Fields. Journal of Liquid Chromatography and Related Technologies, 2006, 29, 307-327.	0.5	1
112	Antifungal activity of tria β - and tetra β -thioureido amino derivatives against different <i>Candida</i> species. Mycoses, 2011, 54, e389-93.	1.8	1
113	Investigação eletroquímica e calorimétrica da interação de novos agentes antitumorais biscati β nicos com DNA. Química Nova, 2012, 35, 1318-1324.	0.3	1
114	Molecular design aided by random forests and synthesis of potent trypanocidal agents as cruzain inhibitors for Chagas disease treatment. Chemical Biology and Drug Design, 2020, 96, 948-960.	1.5	1
115	Estrogen Receptor: Structural Differences and Potential Implications on Selectivity Examined by the GRID/CPCA Approach. Letters in Drug Design and Discovery, 2008, 5, 182-192.	0.4	1
116	Molecular Design, Synthesis and Evaluation of 2,3-Diarylquinoxalines as Estrogen Receptor Ligands. Medicinal Chemistry, 2015, 11, 736-746.	0.7	1
117	The Effect of Dipeptidyl Nitrile Derivatives on Pancreatic Ductal Adenocarcinoma Cells In Vitro. Current Chemical Biology, 2021, 15, 278-286.	0.2	1
118	A CoMFA study on antileishmaniasis bisamidines. , 2000, , 314-315.		0
119	The history, evolution and importance of lipophilicity in medicinal chemistry: from Hippocrates and Galeno to Paracelsus and the contributions of Overton and Hansch. Revista Virtual De Química, 2009, 1, .	0.1	0
120	Structure Based Design of Potential Inhibitors of Steroid Sulfatase. Advances in Intelligent and Soft Computing, 2010, , 151-156.	0.2	0
121	Structure-based design of anti-trypanosomal drugs. Acta Crystallographica Section A: Foundations and Advances, 2011, 67, C294-C294.	0.3	0
122	Discovery of 2-aminopyridine Derivatives with Antichagasic and Antileishmanial Activity Using Phenotypic Assays. Letters in Drug Design and Discovery, 2020, 17, 867-872.	0.4	0