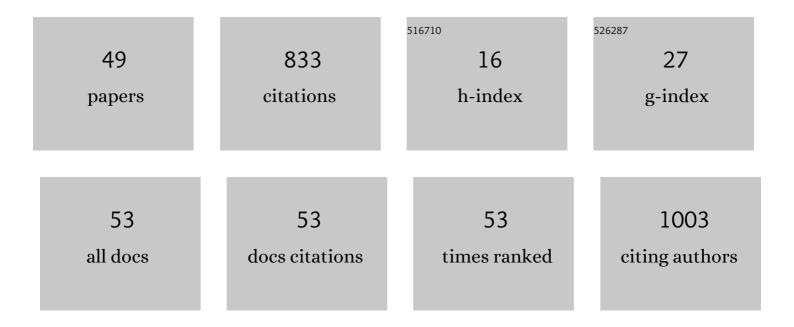
## Ronaldo J Oliveira

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Cholinesterase inhibitors assessment of aporphine alkaloids from Annona crassiflora and molecular docking studies. Bioorganic Chemistry, 2022, 120, 105593.	4.1	6
2	Quantifying biomolecular diffusion with a "spherical cow―model. American Journal of Physics, 2022, 90, 225-238.	0.7	1
3	Gold(III) heteroleptic complexes with SNS-thiosemicarbazonate ligands as cytotoxic agents: Experimental and computational insights into the mechanism of action. Polyhedron, 2022, 219, 115767.	2.2	Ο
4	Biotin-painted proteins have thermodynamic stability switched by kinetic folding routes. Journal of Chemical Physics, 2022, 156, .	3.0	6
5	Shedding Light on the Inhibitory Mechanisms of SARS-CoV-1/CoV-2 Spike Proteins by ACE2-Designed Peptides. Journal of Chemical Information and Modeling, 2021, 61, 1226-1243.	5.4	24
6	The Dynamics of Subunit Rotation in a Eukaryotic Ribosome. Biophysica, 2021, 1, 204-221.	1.4	10
7	Nitroisatin dithiocarbazate: Synthesis, structural characterization, DFT, and docking studies. European Journal of Chemistry, 2021, 12, 235-241.	0.6	1
8	Copper(II) complexes based on thiosemicarbazone ligand: Preparation, crystal structure, Hirshfeld surface, energy framework, antiMycobacterium activity, in silico and molecular docking studies. Journal of Inorganic Biochemistry, 2021, 223, 111543.	3.5	11
9	Ninhydrin as a novel DNA hybridization indicator applied to a highly reusable electrochemical genosensor for Candida auris. Talanta, 2021, 235, 122694.	5.5	6
10	Small Neutral Crowding Solute Effects on Protein Folding Thermodynamic Stability and Kinetics. Journal of Physical Chemistry B, 2021, 125, 11673-11686.	2.6	5
11	The Role of Electrostatics and Folding Kinetics on the Thermostability of Homologous Cold Shock Proteins. Journal of Chemical Information and Modeling, 2020, 60, 546-561.	5.4	10
12	Extension-Dependent Drift Velocity and Diffusion (DrDiff) Directly Reconstructs the Folding Free Energy Landscape of Atomic Force Microscopy Experiments. Journal of Physical Chemistry Letters, 2020, 11, 800-807.	4.6	3
13	Fragmentation Study, Dual Anti-Bactericidal and Anti-Viral Effects and Molecular Docking of Cobalt(III) Complexes. International Journal of Molecular Sciences, 2020, 21, 8355.	4.1	10
14	Preparation, structural characterization, voltammetry and Hirshfeld surface analysis of homoleptic iron(III) thiosemicarbazone complexes. Transition Metal Chemistry, 2020, 45, 511-521.	1.4	3
15	Development of gold(III) thiosemicarbazonate complex–loaded PLGA nanoparticles: characterization and sustained release studies. Journal of Nanoparticle Research, 2020, 22, 1.	1.9	6
16	Intramolecular C(sp <sup>2</sup> )–C (sp <sup>2</sup> ) bond formation between phenanthroline and β-diketone thiosemicarbazones in Pt <sup>II</sup> complexes: crystal structures and computational studies. Dalton Transactions, 2020, 49, 9564-9567.	3.3	1
17	DNA binding, cleavage, apoptosis and cytotoxicity studies of three heteroleptic nickel complexes bearing β-diketones. Inorganica Chimica Acta, 2020, 511, 119824.	2.4	20
18	Novel tetranuclear Pd <sup>II</sup> and Pt <sup>II</sup> anticancer complexes derived from pyrene thiosemicarbazones. Dalton Transactions, 2020, 49, 9595-9604.	3.3	25

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19	Organometallic Gold(III) Complex [Au(Hdamp)(L1 <sup>4</sup> )] <sup>+</sup> (L1 = <i>SNS</i> -Donating) T Diseases, 2019, 5, 1698-1707.	j ETQq1 1 3.8	l 0.784314 rg <mark>8</mark> 1 16
20	Importance of the β5â <sup>~</sup> β6 Loop for the Structure, Catalytic Efficiency, and Stability of Carbapenem-Hydrolyzing Class D β-Lactamase Subfamily OXA-143. Biochemistry, 2019, 58, 3604-3616.	2.5	4
21	A novel peptide-based sensor platform for detection of anti-Toxoplasma gondii immunoglobulins. Journal of Pharmaceutical and Biomedical Analysis, 2019, 175, 112778.	2.8	12
22	Heterobimetallic nickel(II) and palladium(II) complexes derived from S-benzyl-N- (ferrocenyl)methylenedithiocarbazate: Trypanocidal activity and interaction with Trypanosoma cruzi Old Yellow Enzyme (TcOYE). European Journal of Medicinal Chemistry, 2019, 180, 213-223.	5.5	20
23	Drift-diffusion (DrDiff) framework determines kinetics and thermodynamics of two-state folding trajectory and tunes diffusion models. Journal of Chemical Physics, 2019, 151, 114106.	3.0	14
24	A Dynamic Hydrophobic Core and Surface Salt Bridges Thermostabilize a Designed Three-Helix Bundle. Biophysical Journal, 2019, 116, 621-632.	0.5	16
25	Identification of Bioactive Compounds and Analysis of Inhibitory Potential of the Digestive Enzymes from Syzygium sp. Extracts. Journal of Chemistry, 2019, 2019, 1-10.	1.9	4
26	In vitro anti-Trypanosoma cruzi activity of ternary copper(II) complexes and in vivo evaluation of the most promising complex. Biomedicine and Pharmacotherapy, 2019, 109, 157-166.	5.6	23
27	Developing a Mathematical Model for the Controlled Release Over Time of Sulfentrazone Herbicide from Biodegradable Polymer. Materials Research, 2019, 22, .	1.3	3
28	Photophysical and DFT Studies of Cationic Ag(I) Complexes with Thiosemicarbazides Derived from <i>p</i> â€Toluenesulfohydrazide. ChemistrySelect, 2018, 3, 2108-2114.	1.5	2
29	Effects of pH and aggregation in the human prion conversion into scrapie form: a study using molecular dynamics with excited normal modes. European Biophysics Journal, 2018, 47, 583-590.	2.2	10
30	Supersymmetric quantum mechanics method for the Fokker–Planck equation with applications to protein folding dynamics. Physica A: Statistical Mechanics and Its Applications, 2018, 493, 286-300.	2.6	17
31	Stochastic diffusion framework determines the free-energy landscape and rate from single-molecule trajectory. Journal of Chemical Physics, 2018, 149, 234107.	3.0	11
32	Isolation, leishmanicidal evaluation and molecular docking simulations of piperidine alkaloids from Senna spectabilis. Bioorganic and Medicinal Chemistry, 2018, 26, 5816-5823.	3.0	24
33	Introdução ao problema de enovelamento de proteÃnas: uma abordagem utilizando modelos computacionais simplificados. Revista Brasileira De Ensino De Fisica, 2018, 40, .	0.2	2
34	Cu(I) complexes with thiosemicarbazides derived from p-toluenesulfohydrazide: Structural, luminescence and biological studies. Polyhedron, 2018, 155, 170-179.	2.2	14
35	Pt II , Pd II and Au III complexes with a thiosemicarbazone derived from diacethylmonooxime: Structural analysis, trypanocidal activity, cytotoxicity and first insight into the antiparasitic mechanism of action. European Journal of Medicinal Chemistry, 2017, 141, 615-631.	5.5	37
36	Gold(III) complexes with ONS-Tridentate thiosemicarbazones: Toward selective trypanocidal drugs. European Journal of Medicinal Chemistry, 2016, 120, 217-226.	5.5	39

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37	VirB7 and VirB9 Interactions Are Required for the Assembly and Antibacterial Activity of a Type IV Secretion System. Structure, 2016, 24, 1707-1718.	3.3	14
38	Quantifying Nonnative Interactions in the Protein-Folding Free-Energy Landscape. Biophysical Journal, 2016, 111, 287-293.	0.5	21
39	Physicochemical composition and ruminal degradability of leucaena ensiled with different levels of buriti fruit peel. Grassland Science, 2016, 62, 160-166.	1.1	4
40	Structure and functional dynamics characterization of the ion channel of the human respiratory syncytial virus (hRSV) small hydrophobic protein (SH) transmembrane domain by combining molecular dynamics with excited normal modes. Journal of Molecular Modeling, 2016, 22, 286.	1.8	13
41	Electropolymerization of hydroxyphenylacetic acid isomers and the development of a bioelectrode for the diagnosis of bacterial meningitis. Journal of Applied Electrochemistry, 2015, 45, 1277-1287.	2.9	9
42	Manganese(II) complexes with thiosemicarbazones as potential anti-Mycobacterium tuberculosis agents. Journal of Inorganic Biochemistry, 2014, 132, 21-29.	3.5	50
43	The characterization of a thermostable and cambialistic superoxide dismutase from <i>Thermus filiformis</i> . Letters in Applied Microbiology, 2013, 57, 40-46.	2.2	23
44	Analyzing the effect of homogeneous frustration in protein folding. Proteins: Structure, Function and Bioinformatics, 2013, 81, 1727-1737.	2.6	32
45	Configuration-Dependent Diffusion Dynamics of Downhill and Two-State Protein Folding. Journal of Physical Chemistry B, 2012, 116, 5152-5159.	2.6	23
46	Topography of funneled landscapes determines the thermodynamics and kinetics of protein folding. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 15763-15768.	7.1	62
47	The Origin of Nonmonotonic Complex Behavior and the Effects of Nonnative Interactions on the Diffusive Properties of Protein Folding. Biophysical Journal, 2010, 99, 600-608.	0.5	33
48	Coordinate and time-dependent diffusion dynamics in protein folding. Methods, 2010, 52, 91-98.	3.8	36
49	Configuration-dependent diffusion can shift the kinetic transition state and barrier height of protein folding. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 14646-14651.	7.1	92