

# Cleydson Breno Rodrigues dos Santos

## List of Publications by Year in descending order

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80  
papers

1,433  
citations

331670

21  
h-index

377865

34  
g-index

81  
all docs

81  
docs citations

81  
times ranked

1326  
citing authors

#	ARTICLE	IF	CITATIONS
1	Alzheimer's Disease: A Review from the Pathophysiology to Diagnosis, New Perspectives for Pharmacological Treatment. <i>Current Medicinal Chemistry</i> , 2018, 25, 3141-3159.	2.4	193
2	In Silico Study to Identify New Antituberculosis Molecules from Natural Sources by Hierarchical Virtual Screening and Molecular Dynamics Simulations. <i>Pharmaceuticals</i> , 2019, 12, 36.	3.8	55
3	Novel chalcones derivatives with potential antineoplastic activity investigated by docking and molecular dynamics simulations. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 2204-2216.	3.5	53
4	Identification of New Rofecoxib-Based Cyclooxygenase-2 Inhibitors: A Bioinformatics Approach. <i>Pharmaceuticals</i> , 2020, 13, 209.	3.8	52
5	Potential inhibitors of the enzyme acetylcholinesterase and juvenile hormone with insecticidal activity: study of the binding mode via docking and molecular dynamics simulations. <i>Journal of Biomolecular Structure and Dynamics</i> , 2020, 38, 4687-4709.	3.5	51
6	Identification of Potential COX-2 Inhibitors for the Treatment of Inflammatory Diseases Using Molecular Modeling Approaches. <i>Molecules</i> , 2020, 25, 4183.	3.8	45
7	Identification of Novel Chemical Entities for Adenosine Receptor Type 2A Using Molecular Modeling Approaches. <i>Molecules</i> , 2020, 25, 1245.	3.8	45
8	Methoxylated flavonols from <i>Vellozia dasypus</i> Seub ethyl acetate active myeloperoxidase extract: in vitro and in silico assays. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 7574-7583.	3.5	44
9	Chemometric methods in antimalarial drug design from 1,2,4,5-tetraoxanes analogues. <i>SAR and QSAR in Environmental Research</i> , 2020, 31, 677-695.	2.2	43
10	Identification of Potential Inhibitors from Pyriproxyfen with Insecticidal Activity by Virtual Screening. <i>Pharmaceuticals</i> , 2019, 12, 20.	3.8	42
11	A SAR and QSAR Study of New Artemisinin Compounds with Antimalarial Activity. <i>Molecules</i> , 2014, 19, 367-399.	3.8	38
12	An In Silico Study of the Antioxidant Ability for Two Caffeine Analogs Using Molecular Docking and Quantum Chemical Methods. <i>Molecules</i> , 2018, 23, 2801.	3.8	38
13	Pharmacophore-based virtual screening and molecular docking to identify promising dual inhibitors of human acetylcholinesterase and butyrylcholinesterase. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 6021-6030.	3.5	35
14	An Antioxidant Potential, Quantum-Chemical and Molecular Docking Study of the Major Chemical Constituents Present in the Leaves of <i>Curatella americana</i> Linn. <i>Pharmaceuticals</i> , 2018, 11, 72.	3.8	33
15	Identification of Novel Protein Kinase Receptor Type 2 Inhibitors Using Pharmacophore and Structure-Based Virtual Screening. <i>Molecules</i> , 2018, 23, 453.	3.8	30
16	Pharmacophore and structure-based drug design, molecular dynamics and admet/tox studies to design novel potential pad4 inhibitors. <i>Journal of Biomolecular Structure and Dynamics</i> , 2019, 37, 966-981.	3.5	30
17	Molecular modeling approaches of selective adenosine receptor type 2A agonists as potential anti-inflammatory drugs. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 1-13.	3.5	30
18	Virtual Screening and Statistical Analysis in the Design of New Caffeine Analogues Molecules with Potential Epithelial Anticancer Activity. <i>Current Pharmaceutical Design</i> , 2018, 24, 576-594.	1.9	28

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19	Computational design of new protein kinase 2 inhibitors for the treatment of inflammatory diseases using QSAR, pharmacophore-structure-based virtual screening, and molecular dynamics. <i>Journal of Molecular Modeling</i> , 2018, 24, 225.	1.8	26
20	A QSAR, Pharmacokinetic and Toxicological Study of New Artemisinin Compounds with Anticancer Activity. <i>Molecules</i> , 2014, 19, 10670-10697.	3.8	24
21	Natural Products-Based Drug Design against SARS-CoV-2 Mpro 3CLpro. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11739.	4.1	24
22	Oil from the fruits of <i>Pterodon emarginatus</i> Vog.: A traditional anti-inflammatory. Study combining in vivo and in silico. <i>Journal of Ethnopharmacology</i> , 2018, 222, 107-120.	4.1	23
23	In Silico Evaluation of Ibuprofen and Two Benzoylpropionic Acid Derivatives with Potential Anti-Inflammatory Activity. <i>Molecules</i> , 2019, 24, 1476.	3.8	23
24	<i>Acmella oleracea</i> (L) R. K. Jansen Reproductive Toxicity in Zebrafish: An <i>In Vivo</i> and <i>In Silico</i> Assessment. <i>Evidence-based Complementary and Alternative Medicine</i> , 2019, 2019, 1-19.	1.2	21
25	Molecular Modeling: Origin, Fundamental Concepts and Applications Using Structure-Activity Relationship and Quantitative Structure-Activity Relationship. <i>Reviews in Theoretical Science</i> , 2014, 2, 91-115.	0.5	21
26	Computational Analysis of Physicochemical, Pharmacokinetic and Toxicological Properties of Deoxyhypusine Synthase Inhibitors with Antimalarial Activity. <i>Computational Molecular Bioscience</i> , 2014, 04, 47-57.	0.4	19
27	Computational Investigation of Antifungal Compounds Using Molecular Modeling and Prediction of ADME/Tox Properties. <i>Journal of Computational and Theoretical Nanoscience</i> , 2015, 12, 3682-3691.	0.4	18
28	Molecular modeling and statistical analysis in the design of derivatives of human dipeptidyl peptidase IV. <i>Journal of Biomolecular Structure and Dynamics</i> , 2018, 36, 318-334.	3.5	18
29	Hierarchical Virtual Screening of Potential Insecticides Inhibitors of Acetylcholinesterase and Juvenile Hormone from Temephos. <i>Pharmaceuticals</i> , 2019, 12, 61.	3.8	18
30	Toward of Safer Phenylbutazone Derivatives by Exploration of Toxicity Mechanism. <i>Molecules</i> , 2019, 24, 143.	3.8	17
31	Application of Hartree-Fock Method for Modeling of Bioactive Molecules Using SAR and QSPR. <i>Computational Molecular Bioscience</i> , 2014, 04, 1-24.	0.4	17
32	Identification of New Inhibitors with Potential Antitumor Activity from Polypeptide Structures via Hierarchical Virtual Screening. <i>Molecules</i> , 2019, 24, 2943.	3.8	16
33	Studies of NMR, molecular docking, and molecular dynamics simulation of new promising inhibitors of cruzaine from the parasite <i>Trypanosoma cruzi</i> . <i>Medicinal Chemistry Research</i> , 2019, 28, 246-259.	2.4	15
34	Development of nano-emulsions based on <i>Ayapana triplinervis</i> essential oil for the control of <i>Aedes aegypti</i> larvae. <i>PLoS ONE</i> , 2021, 16, e0254225.	2.5	14
35	Hydroethanolic extract from <i>Endopleura uchi</i> (Huber) Cuatrecasas and its marker bergenin: Toxicological and pharmacokinetic studies in silico and in vivo on zebrafish. <i>Toxicology Reports</i> , 2020, 7, 217-232.	3.3	14
36	Rational Design of Antimalarial Drugs Using Molecular Modeling and Statistical Analysis. <i>Current Pharmaceutical Design</i> , 2015, 21, 4112-4127.	1.9	13

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37	Cannabinoid Type 1 Receptor (CB1) Ligands with Therapeutic Potential for Withdrawal Syndrome in Chemical Dependents of <i>Cannabis sativa</i> . <i>ChemMedChem</i> , 2017, 12, 1408-1416.	3.2	12
38	New PPAR $\alpha$ Optimal Activator Rationally Designed by Computational Methods. <i>Journal of the Brazilian Chemical Society</i> , 2016, , .	0.6	11
39	Identification of Potential Antiviral Inhibitors from Hydroxychloroquine and 1,2,4,5-Tetraoxanes Analogues and Investigation of the Mechanism of Action in SARS-CoV-2. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1781.	4.1	11
40	Evaluation of Quantum Chemical Methods and Basis Sets Applied in the Molecular Modeling of Artemisinin. <i>Computational Molecular Bioscience</i> , 2013, 03, 66-79.	0.4	10
41	A Computational Approach Applied to the Study of Potential Allosteric Inhibitors Protease NS2B/NS3 from Dengue Virus. <i>Molecules</i> , 2022, 27, 4118.	3.8	10
42	Antimalarial Artemisinins Derivatives Study: Molecular Modeling and Multivariate Analysis (PCA, HCA,) Tj ETQq0 0 0 rgBT /Overlock 10 T	0.4	9
43	A Viability Study for the Production of Biofilms and <i>In Silico</i> Predictions of Major Compounds in Kefir. <i>Journal of Computational and Theoretical Nanoscience</i> , 2017, 14, 2915-2926.	0.4	9
44	Revisiting the Proposition of Binding Pockets and Bioactive Poses for GSK-3 $\beta$ Allosteric Modulators Addressed to Neurodegenerative Diseases. <i>International Journal of Molecular Sciences</i> , 2021, 22, 8252.	4.1	9
45	Hierarchical Virtual Screening Based on Rocaglamide Derivatives to Discover New Potential Anti-Skin Cancer Agents. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, .	3.5	9
46	Identification of novel <i>Aedes aegypti</i> odorant-binding protein 1 modulators by ligand and structure-based approaches and bioassays. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 117-129.	3.5	8
47	ANTIOPHIDIAN ACTIVITY OF <i>BROSIMUM GUIANENSE</i> (AUBL) HUBER. <i>American Journal of Pharmacology and Toxicology</i> , 2014, 9, 148-156.	0.7	7
48	Identification of novel $\beta$ -tubulin modulators with antiproliferative activity directed to cancer therapy using ligand and structure-based virtual screening. <i>International Journal of Biological Macromolecules</i> , 2020, 165, 3040-3050.	7.5	7
49	The Role of Celecoxib as a Potential Inhibitor in the Treatment of Inflammatory Diseases - A Review. <i>Current Medicinal Chemistry</i> , 2022, 29, 3028-3049.	2.4	7
50	Potential colchicine binding site inhibitors unraveled by virtual screening, molecular dynamics and MM/PBSA. <i>Computers in Biology and Medicine</i> , 2021, 137, 104817.	7.0	7
51	Alkylated Sesamol Derivatives as Potent Antioxidants. <i>Molecules</i> , 2020, 25, 3300.	3.8	6
52	Chemical profiling of <i>Curatella americana</i> Linn leaves by UPLC-HRMS and its wound healing activity in mice. <i>PLoS ONE</i> , 2020, 15, e0225514.	2.5	6
53	Validation of Computational Methods Applied in Molecular Modeling of Artemisinin with Antimalarial Activity. <i>Journal of Computational and Theoretical Nanoscience</i> , 2014, 11, 553-561.	0.4	5
54	Development of Monoamine Oxidase B Inhibitors with Antiparkinson Activity. <i>Current Physical Chemistry</i> , 2016, 6, 40-52.	0.2	5

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55	Identification of potential modulator of <i>Anopheles gambiae</i> odorant binding protein 1 by hierarchical virtual screening and molecular dynamics. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 6031-6043.	3.5	5
56	Identification of novel potential cyclooxygenase-2 inhibitors using ligand- and structure-based virtual screening approaches. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 5386-5408.	3.5	5
57	Ethyl Acetate Fraction of <i>Bixa orellana</i> and Its Component Ellagic Acid Exert Antibacterial and Anti-Inflammatory Properties against <i>Mycobacterium abscessus</i> subsp. <i>massiliense</i> . <i>Antibiotics</i> , 2022, 11, 817.	3.7	5
58	Production and characterization of absorbent heat from the bark of residual Brazil nut bark ( <i>Bertholletia Excelsa</i> l.). <i>Chemistry Central Journal</i> , 2015, 9, 36.	2.6	4
59	Assessment of the hypoglycemic effect of Bixin in alloxan-induced diabetic rats: <i>in vivo</i> and <i>in silico</i> studies. <i>Journal of Biomolecular Structure and Dynamics</i> , 2021, 39, 1017-1028.	3.5	4
60	An Overview of the $\alpha_1$ Integrin and the Potential Therapeutic Role of its Antagonists. <i>Current Medicinal Chemistry</i> , 2021, 28, 5884-5895.	2.4	4
61	A comparative theoretical mechanism on simplified flavonoid derivatives and isoxazolone analogous as Michael system inhibitor. <i>Journal of Molecular Modeling</i> , 2021, 27, 26.	1.8	3
62	An <i>in Silico</i> Study of Natural Compounds as Potential MAO-B Inhibitors for the Treatment of Parkinson's Disease. <i>Engineering Materials</i> , 2021, , 591-617.	0.6	3
63	Molecular Modeling of Peptide Derivatives NS3 Protease Inhibitors of the Type 2 Dengue Virus. <i>Current Physical Chemistry</i> , 2016, 6, 28-39.	0.2	2
64	Ligand- and Structure-Based Drug Design of Novel Calcium Channel Blockers. <i>Journal of Computational and Theoretical Nanoscience</i> , 2017, 14, 3489-3502.	0.4	2
65	A MLR and ADME/Tox Study of New Dihydroartemisinin Compounds with Antimalarial Activity. <i>Journal of Computational and Theoretical Nanoscience</i> , 2018, 15, 1785-1794.	0.4	2
66	Chemical Study, Predictions <i>In Silico</i> and Larvicide Activity of the Essential Oil of Root <i>Philodendron deflexum</i> Poepp.. <i>Journal of Computational and Theoretical Nanoscience</i> , 2017, 14, 3330-3337.	0.4	2
67	Evaluation of Computational Method from Crystallographic Structure of Galantamine for Molecular Modeling of Drug Candidates Anti-Alzheimer's Disease. <i>Journal of Computational and Theoretical Nanoscience</i> , 2019, 16, 2673-2686.	0.4	2
68	Theoretical Study of Monoamine Oxidase B Inhibitors as Drug Candidates for Treatment of Parkinson's Disease. <i>Central Nervous System Agents in Medicinal Chemistry</i> , 2020, 20, 128-143.	1.1	2
69	Application of Chemometric Techniques for Validation of Computational Methods Applied in Molecular Modeling of 6-Hydroxyouacapan-7-Lactone with Antiproliferative Activity in Leukemia Cells. <i>Journal of Computational and Theoretical Nanoscience</i> , 2020, 17, 4855-4865.	0.4	2
70	Virtual screening and molecular docking of structures with potential inhibitor of the ebolavirus glycoprotein. <i>Research, Society and Development</i> , 2022, 11, e45311226034.	0.1	2
71	Physicochemical Characterization of Water Quality - Lagoa dos Ándios in Macapá, Brazil. <i>American Chemical Science Journal</i> , 2015, 5, 122-134.	0.2	1
72	<i>In vivo</i> and <i>in vitro</i> evaluation of antiplasmodial activity of <i>Amasonia campestris</i> (Aubl.) Moldenke. <i>African Journal of Pharmacy and Pharmacology</i> , 2017, 11, 377-384.	0.3	1

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73	The role of regioselective hydroxylation on toxicity of diclofenac and related derivatives. <i>Molecular Simulation</i> , 2019, 45, 1454-1458.	2.0	1
74	ADME/Tox Study and Molecular Dynamics Simulations Applied in the Design of New Potential GABA-AT Inhibitors. <i>Engineering Materials</i> , 2021, , 719-738.	0.6	1
75	High-Throughput-Based Virtual Screening via Molecular Docking for Oxidative Stress Mediated by ROS Enzyme. <i>Engineering Materials</i> , 2021, , 489-513.	0.6	1
76	Pharmaceutical and Biological Potential of the Croton palanostigma Isolated Compounds. <i>Journal of Computational and Theoretical Nanoscience</i> , 2019, 16, 1773-1782.	0.4	1
77	Estudo Físico-Químico e Avaliação do Potencial Larvicida do Extrato Etanólico das Cascas do Caule de <i>Licania macrophylla</i> Benth. <i>Biota Amazônia</i> , 2015, 5, 74-78.	0.2	1
78	Molecular Modeling as Motivating Tool in Teaching and Learning in Mechanisms of Diels-Alder Reactions. <i>Revista Virtual De Química</i> , 2016, 8, 2026-2041.	0.4	0
79	Chemical Study, Predictions <i>In Silico</i> and Larvicide Activity of the Essential Oil of Root <i>Philodendron deflexum</i> Poepp. ( <i>Journal of Computational and Theoretical Nanoscience</i> , Vol. 14(7),) Tj ETQq1 1 0.7843140gBT /Over	0.4	0
80	An <i>In Silico</i> and <i>In Vitro</i> Study of the Metal Complex di- $\frac{1}{4}$ -Chloro-Bis[Chlorine (4,7-Dimethyl-1,10-Phenanthroline) Cadmium(ii)] with Antibacterial Potential. <i>Journal of Computational and Theoretical Nanoscience</i> , 2021, 18, 1702-1713.	0.4	0