

Icaro P Caruso

List of Publications by Year in descending order

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55
papers

550
citations

686830

13
h-index

752256

20
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59
all docs

59
docs citations

59
times ranked

833
citing authors

#	ARTICLE	IF	CITATIONS
1	Hesperetin targets the hydrophobic pocket of the nucleoprotein/phosphoprotein binding site of human respiratory syncytial virus. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, 40, 2156-2168.	2.0	6
2	Insights into the specificity for the interaction of the promiscuous SARS-CoV-2 nucleocapsid protein N-terminal domain with deoxyribonucleic acids. <i>International Journal of Biological Macromolecules</i> , 2022, 203, 466-480.	3.6	16
3	Characterization of secondary structure and thermal stability by biophysical methods of the D-alanyl,D-alanine ligase B protein from <i>Escherichia coli</i> . <i>Protein and Peptide Letters</i> , 2022, 29, .	0.4	0
4	Searching for drug leads targeted to the hydrophobic cleft of dengue virus capsid protein. <i>Journal of Enzyme Inhibition and Medicinal Chemistry</i> , 2022, 37, 287-298.	2.5	1
5	Deciphering the Path of S-nitrosation of Human Thioredoxin: Evidence of an Internal NO Transfer and Implication for the Cellular Responses to NO. <i>Antioxidants</i> , 2022, 11, 1236.	2.2	2
6	Interaction between iron ion and dipole carbon monoxide inside spherical cavities. <i>European Physical Journal D</i> , 2021, 75, 1.	0.6	0
7	¹ H, ¹⁵ N and ¹³ C resonance assignments of the N-terminal domain of the nucleocapsid protein from the endemic human coronavirus HKU1. <i>Biomolecular NMR Assignments</i> , 2021, 15, 153-157.	0.4	2
8	Unique structural features of flaviviruses' capsid proteins: new insights on structure-function relationship. <i>Current Opinion in Virology</i> , 2021, 47, 106-112.	2.6	5
9	Structural and biophysical characterization of the major proteins from the seminal plasma of Dorper rams. <i>Theriogenology</i> , 2021, 166, 64-70.	0.9	0
10	Large-Scale Recombinant Production of the SARS-CoV-2 Proteome for High-Throughput and Structural Biology Applications. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 653148.	1.6	29
11	Thermodynamic profile and molecular modeling of the interaction between Grb2 dimer and flavonoids Rutin and Morin. <i>Journal of Molecular Structure</i> , 2021, 1234, 130164.	1.8	1
12	Dynamics of the SARS-CoV-2 nucleoprotein N-terminal domain triggers RNA duplex destabilization. <i>Biophysical Journal</i> , 2021, 120, 2814-2827.	0.2	12
13	Protein Surface Interactions' Theoretical and Experimental Studies. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 706002.	1.6	9
14	¹ H, ¹⁵ N, and ¹³ C resonance assignments of the SH3-like tandem domain of human KIN protein. <i>Biomolecular NMR Assignments</i> , 2021, 15, 449-453.	0.4	0
15	What Is the Effect of Physical Exercise on Nafld/Nash. <i>American Journal of Sports Science</i> , 2021, 9, 1.	0.2	0
16	Annexin A1 Mimetic Peptide and Piperlongumine: Anti-Inflammatory Profiles in Endotoxin-Induced Uveitis. <i>Cells</i> , 2021, 10, 3170.	1.8	9
17	Quercetin pentaacetate inhibits in vitro human respiratory syncytial virus adhesion. <i>Virus Research</i> , 2020, 276, 197805.	1.1	40
18	4-Methylesculetin, a natural coumarin with intestinal anti-inflammatory activity, elicits a glutathione antioxidant response by different mechanisms. <i>Chemico-Biological Interactions</i> , 2020, 315, 108876.	1.7	19

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19	Fluorescence spectroscopic and dynamics simulation studies on isoorientin binding with human serum albumin. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2020, 228, 117738.	2.0	8
20	Biophysical and Dynamic Characterization of Fine-Tuned Binding of the Human Respiratory Syncytial Virus M2-1 Core Domain to Long RNAs. <i>Journal of Virology</i> , 2020, 94, .	1.5	3
21	The dynamics of free and phosphopeptide-bound Grb2-SH2 reveals two dynamically independent subdomains and an encounter complex with fuzzy interactions. <i>Scientific Reports</i> , 2020, 10, 13040.	1.6	11
22	Biophysical Characterization of the Binding of hRSV M2-1 Protein to RNA and Solasodine. <i>Biophysical Journal</i> , 2020, 118, 515a.	0.2	0
23	Study protocol to investigate biomolecular muscle profile as predictors of long-term urinary incontinence in women with gestational diabetes mellitus. <i>BMC Pregnancy and Childbirth</i> , 2020, 20, 117.	0.9	14
24	The Application of NMR Metabolomics in the Study of the Wound Healing Process. <i>Biophysical Journal</i> , 2020, 118, 135a-136a.	0.2	0
25	Characterization of the Thermal and Chemical Denaturation of the Matrix Protein from hRSV. <i>Biophysical Journal</i> , 2020, 118, 358a-359a.	0.2	2
26	Insights into Interactions of Flavanones with Target Human Respiratory Syncytial Virus M2-1 Protein from STD-NMR, Fluorescence Spectroscopy, and Computational Simulations. <i>International Journal of Molecular Sciences</i> , 2020, 21, 2241.	1.8	15
27	Biophysical characterization and molecular phylogeny of human KIN protein. <i>European Biophysics Journal</i> , 2019, 48, 645-657.	1.2	5
28	An investigation into the interaction between pipartine (piperlongumine) and human serum albumin. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2019, 220, 117084.	2.0	11
29	NMR assignment of free ¹ H, ¹⁵ N and ¹³ C-Grb2-SH2 domain. <i>Biomolecular NMR Assignments</i> , 2019, 13, 295-298.	0.4	5
30	Biophysical and flavonoid-binding studies of the G protein ectodomain of group A human respiratory syncytial virus. <i>Heliyon</i> , 2019, 5, e01394.	1.4	10
31	Grb2 dimer interacts with Coumarin through SH2 domains: A combined experimental and molecular modeling study. <i>Heliyon</i> , 2019, 5, e02869.	1.4	13
32	Binding investigation between M2-1protein from hRSV and acetylated quercetin derivatives: ¹ H NMR, fluorescence spectroscopy, and molecular docking. <i>International Journal of Biological Macromolecules</i> , 2018, 111, 33-38.	3.6	7
33	Spectral and computational features of the binding between riparins and human serum albumin. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2018, 190, 81-88.	2.0	16
34	Structure and interaction of <i>Corynebacterium pseudotuberculosis</i> cold shock protein A with Y-box single-stranded DNA fragment. <i>FEBS Journal</i> , 2018, 285, 372-390.	2.2	6
35	Investigation on the interaction between nanocarriers and functional compounds of flavonoids. <i>Acta Horticulturae</i> , 2018, , 93-96.	0.1	0
36	Biophysical characterization of the interaction between M2-1 protein of hRSV and quercetin. <i>International Journal of Biological Macromolecules</i> , 2017, 95, 63-71.	3.6	8

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37	Bacterial and Arachnid Sphingomyelinases D: Comparison of Biophysical and Pathological Activities. <i>Journal of Cellular Biochemistry</i> , 2017, 118, 2053-2063.	1.2	6
38	Potential Implications for Designing Drugs Against the Brown Spider Venom Phospholipase A ₂ . <i>Journal of Cellular Biochemistry</i> , 2017, 118, 726-738.	1.2	26
39	Exploring the Binding Mechanism of Flavonoid Quercetin to Phospholipase A ₂ : Fluorescence Spectroscopy and Computational Approach. <i>European Journal of Experimental Biology</i> , 2017, 07, .	0.3	7
40	Cold Shock Protein A from <i>Corynebacterium pseudotuberculosis</i> : Role of Electrostatic Forces in the Stability of the Secondary Structure. <i>Protein and Peptide Letters</i> , 2017, 24, 358-367.	0.4	15
41	Active site mapping of <i>Loxosceles</i> phospholipases D: Biochemical and biological features. <i>Biochimica Et Biophysica Acta - Molecular and Cell Biology of Lipids</i> , 2016, 1861, 970-979.	1.2	29
42	An integrated approach with experimental and computational tools outlining the cooperative binding between 2-phenylchromone and human serum albumin. <i>Food Chemistry</i> , 2016, 196, 935-942.	4.2	15
43	Experimental evidence and molecular modeling of the interaction between hRSV-NS1 and quercetin. <i>International Journal of Biological Macromolecules</i> , 2016, 85, 40-47.	3.6	11
44	Functional expression, monodispersity and conformational changes in the SBMV virus viral VPg on binding TFE. <i>International Journal of Biological Macromolecules</i> , 2016, 83, 178-184.	3.6	0
45	Investigation of the Interaction between Coumarins and Its Derivatives with Human Serum Albumin: STD-NMR, Fluorescence and Docking Molecular Studies. <i>International Journal of Sciences</i> , 2016, 2, 1-11.	0.2	1
46	Expression, purification and characterization of cold shock protein A of <i>Corynebacterium pseudotuberculosis</i> . <i>Protein Expression and Purification</i> , 2015, 112, 15-20.	0.6	13
47	Crystal structure of mature 2S albumin from <i>Moringa oleifera</i> seeds. <i>Biochemical and Biophysical Research Communications</i> , 2015, 468, 365-371.	1.0	43
48	Crystal structure of <i>Staphylococcus aureus</i> exfoliative toxin D-like protein: Structural basis for the high specificity of exfoliative toxins. <i>Biochemical and Biophysical Research Communications</i> , 2015, 467, 171-177.	1.0	13
49	Chemical and thermal influence of the [4Fe-4S] ²⁺ cluster of A/G-specific adenine glycosylase from <i>Corynebacterium pseudotuberculosis</i> . <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2015, 1850, 393-400.	1.1	6
50	Preparation and Characterization of Monomodal Grapevine Virus A Capsid Protein. <i>Protein and Peptide Letters</i> , 2015, 22, 712-718.	0.4	0
51	Identification of Folding Intermediates of Streblin, The Most Stable Serine Protease: Biophysical Analysis. <i>Applied Biochemistry and Biotechnology</i> , 2014, 172, 658-671.	1.4	7
52	Binding of antioxidant flavone isovitexin to human serum albumin investigated by experimental and computational assays. <i>Journal of Pharmaceutical and Biomedical Analysis</i> , 2014, 98, 100-106.	1.4	25
53	Exploring the binding mechanism of Guajaverin to human serum albumin: Fluorescence spectroscopy and computational approach. <i>Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy</i> , 2012, 97, 449-455.	2.0	33
54	Effect of modified clays on the structure and functional properties of biofilms produced with zein. <i>Food Science and Technology</i> , 2012, 32, 314-322.	0.8	8

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55	Molecular Docking Studies of Curcumin Analogues against SARS-CoV-2 Spike Protein. Journal of the Brazilian Chemical Society, 0, , .	0.6	1