

Miguel Machuqueiro

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

74
papers

1,844
citations

22
h-index

40
g-index

102
ext. papers

2,288
ext. citations

6.2
avg, IF

5.06
L-index

#	Paper	IF	Citations
74	Network biology and artificial intelligence drive the understanding of the multidrug resistance phenotype in cancer.. <i>Drug Resistance Updates</i> , 2022 , 60, 100811	23.2	3
73	Key Factors for Activated Carbon Adsorption of Pharmaceutical Compounds from Wastewaters: A Multivariate Modelling Approach. <i>Water (Switzerland)</i> , 2022 , 14, 166	3	3
72	Mutant p53 reactivator SLMP53-2 hinders ultraviolet B radiation-induced skin carcinogenesis. <i>Pharmacological Research</i> , 2021 , 106026	10.2	0
71	In Silico End-to-End Protein-Ligand Interaction Characterization Pipeline: The Case of SARS-CoV-2. <i>ACS Synthetic Biology</i> , 2021 , 10, 3209-3235	5.7	0
70	Designing new antitubercular isoniazid derivatives with improved reactivity and membrane trafficking abilities. <i>Biomedicine and Pharmacotherapy</i> , 2021 , 144, 112362	7.5	5
69	Computational Analysis of the Interactions between the S100B Extracellular Chaperone and Its Amyloid β Peptide Client. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	1
68	Improved Protocol to Tackle the pH Effects on Membrane-Inserting Peptides. <i>Journal of Chemical Theory and Computation</i> , 2021 , 17, 3830-3840	6.4	1
67	pKPDB: a Protein Data Bank extension database of pKa and pI theoretical values. <i>Bioinformatics</i> , 2021 ,	7.2	1
66	Cu-binding to S100B triggers polymerization of disulfide cross-linked tetramers with enhanced chaperone activity against amyloid- β aggregation. <i>Chemical Communications</i> , 2021 , 57, 379-382	5.8	2
65	Identification of Pan-Assay Interference compoundS (PAINS) Using an MD-Based Protocol. <i>Methods in Molecular Biology</i> , 2021 , 2315, 263-271	1.4	1
64	pK Calculations in Membrane Proteins from Molecular Dynamics Simulations. <i>Methods in Molecular Biology</i> , 2021 , 2315, 185-195	1.4	1
63	Predicting stable binding modes from simulated dimers of the D76N mutant of 2-microglobulin. <i>Computational and Structural Biotechnology Journal</i> , 2021 , 19, 5160-5169	6.8	0
62	The Lysosomotropic Activity of Hydrophobic Weak Base Drugs is Mediated via Their Intercalation into the Lysosomal Membrane. <i>Cells</i> , 2020 , 9,	7.9	20
61	Improvement of conventional anti-cancer drugs as new tools against multidrug resistant tumors. <i>Drug Resistance Updates</i> , 2020 , 50, 100682	23.2	72
60	Glucosylpolyphenols as Inhibitors of A β -induced Fyn Kinase Activation and Tau Phosphorylation: Synthesis, Membrane Permeability, and Exploratory Target Assessment within the Scope of Type 2 Diabetes and Alzheimer's Disease. <i>Journal of Medicinal Chemistry</i> , 2020 , 63, 11663-11690	8.3	3
59	PypKa: A Flexible Python Module for Poisson-Boltzmann-Based p Calculations. <i>Journal of Chemical Information and Modeling</i> , 2020 , 60, 4442-4448	6.1	13
58	Improved GROMOS 54A7 Charge Sets for Phosphorylated Tyr, Ser, and Thr to Deal with pH-Dependent Binding Phenomena. <i>Journal of Chemical Theory and Computation</i> , 2020 , 16, 6368-6376	6.4	6

57	Novel "ruthenium cyclopentadienyl"-peptide conjugate complexes against human FGFR(+) breast cancer. <i>Dalton Transactions</i> , 2020 , 49, 5974-5987	4.3	5
56	The Early Phase of β m Aggregation: An Integrative Computational Study Framed on the D76N Mutant and the β 6 Variant. <i>Biomolecules</i> , 2019 , 9,	5.9	6
55	The multi-factorial nature of clinical multidrug resistance in cancer. <i>Drug Resistance Updates</i> , 2019 , 46, 100645	23.2	155
54	Evaluation of EGCG Loading Capacity in DMPC Membranes. <i>Langmuir</i> , 2019 , 35, 6771-6781	4	3
53	Molybdenum(ii) complexes with p-substituted BIAN ligands: synthesis, characterization, biological activity and computational study. <i>Dalton Transactions</i> , 2019 , 48, 8449-8463	4.3	7
52	A pH Replica Exchange Scheme in the Stochastic Titration Constant-pH MD Method. <i>Journal of Chemical Theory and Computation</i> , 2019 , 15, 3108-3116	6.4	7
51	Bioactivities of (Gentianaceae) Decoctions: Antioxidant Activity, Enzyme Inhibition and Docking Studies. <i>Molecules</i> , 2019 , 24,	4.8	15
50	Dynamical Rearrangement of Human Epidermal Growth Factor Receptor 2 upon Antibody Binding: Effects on the Dimerization. <i>Biomolecules</i> , 2019 , 9,	5.9	1
49	Bioactivities of decoctions from <i>Plectranthus</i> species related to their traditional use on the treatment of digestive problems and alcohol intoxication. <i>Journal of Ethnopharmacology</i> , 2018 , 220, 147-154	5	8
48	Role of Counterions in Constant-pH Molecular Dynamics Simulations of PAMAM Dendrimers. <i>ACS Omega</i> , 2018 , 3, 2001-2009	3.9	11
47	Apple tree branches derived activated carbons for the removal of β blocker atenolol. <i>Chemical Engineering Journal</i> , 2018 , 345, 669-678	14.7	32
46	Proactive response to tackle the threat of emerging drugs: Synthesis and toxicity evaluation of new cathinones. <i>Forensic Science International</i> , 2018 , 290, 146-156	2.6	22
45	Membrane-Induced pK Shifts in wt-pHLIP and Its L16H Variant. <i>Journal of Chemical Theory and Computation</i> , 2018 , 14, 3289-3297	6.4	18
44	Differential targeting of membrane lipid domains by caffeic acid and its ester derivatives. <i>Free Radical Biology and Medicine</i> , 2018 , 115, 232-245	7.8	31
43	Sugar-based bactericides targeting phosphatidylethanolamine-enriched membranes. <i>Nature Communications</i> , 2018 , 9, 4857	17.4	15
42	The Impact of Using Single Atomistic Long-Range Cutoff Schemes with the GROMOS 54A7 Force Field. <i>Journal of Chemical Theory and Computation</i> , 2018 , 14, 5823-5833	6.4	16
41	Biomolecular Simulations of Halogen Bonds with a GROMOS Force Field. <i>Journal of Chemical Theory and Computation</i> , 2018 , 14, 5383-5392	6.4	14
40	The Catalase Activity of Catalase-Peroxidases Is Modulated by Changes in the pK of the Distal Histidine. <i>Biochemistry</i> , 2017 , 56, 2271-2281	3.2	9

39	Targeting Type 2 Diabetes with C-Glucosyl Dihydrochalcones as Selective Sodium Glucose Co-Transporter 2 (SGLT2) Inhibitors: Synthesis and Biological Evaluation. <i>Journal of Medicinal Chemistry</i> , 2017 , 60, 568-579	8.3	35
38	Insights on the Mechanism of Action of INH-C as an Antitubercular Prodrug. <i>Molecular Pharmaceutics</i> , 2017 , 14, 4597-4605	5.6	10
37	Isorhamnetin derivatives and piscidic acid for hypercholesterolemia: cholesterol permeability, HMG-CoA reductase inhibition, and docking studies. <i>Archives of Pharmacal Research</i> , 2017 , 40, 1278-1286	6.1	29
36	Antiacetylcholinesterase activity and docking studies with chlorogenic acid, cynarin and arzanol from <i>Helichrysum stoechas</i> (Lamiaceae). <i>Medicinal Chemistry Research</i> , 2017 , 26, 2942-2950	2.2	14
35	A tale of two tails: The importance of unstructured termini in the aggregation pathway of β -microglobulin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2017 , 85, 2045-2057	4.2	10
34	Exploring the Structural Properties of Positively Charged Peptide Dendrimers. <i>Journal of Physical Chemistry B</i> , 2016 , 120, 11323-11330	3.4	15
33	pK(a) Values of Titrable Amino Acids at the Water/Membrane Interface. <i>Journal of Chemical Theory and Computation</i> , 2016 , 12, 930-4	6.4	42
32	Hydrogen peroxide regulates cell adhesion through the redox sensor RPSA. <i>Free Radical Biology and Medicine</i> , 2016 , 90, 145-57	7.8	12
31	Structuring Peptide Dendrimers through pH Modulation and Substrate Binding. <i>Journal of Physical Chemistry B</i> , 2016 , 120, 10138-10152	3.4	7
30	The role of electrostatics in TrxR electron transfer mechanism: A computational approach. <i>Proteins: Structure, Function and Bioinformatics</i> , 2016 , 84, 1836-1843	4.2	6
29	Constant-pH MD Simulations of an Oleic Acid Bilayer. <i>Journal of Chemical Theory and Computation</i> , 2015 , 11, 2367-76	6.4	22
28	Constant-pH Molecular Dynamics Study of Kyotorphin in an Explicit Bilayer. <i>Biophysical Journal</i> , 2015 , 108, 2282-90	2.9	11
27	Constant-pH MD Simulations of DMPA/DMPC Lipid Bilayers. <i>Journal of Chemical Theory and Computation</i> , 2015 , 11, 5973-9	6.4	23
26	Molecular details of INH-C10 binding to wt KatG and Its S315T mutant. <i>Molecular Pharmaceutics</i> , 2015 , 12, 898-909	5.6	10
25	Protonation of DMPC in a Bilayer Environment Using a Linear Response Approximation. <i>Journal of Chemical Theory and Computation</i> , 2014 , 10, 2176-84	6.4	22
24	Influence of activated carbons porous structure on iopamidol adsorption. <i>Carbon</i> , 2014 , 77, 607-615	10.4	21
23	Why a diaminopyrrolic tripodal receptor binds mannosides in acetonitrile but not in water?. <i>Beilstein Journal of Organic Chemistry</i> , 2014 , 10, 1513-23	2.5	6
22	A simulated intermediate state for folding and aggregation provides insights into β 6 β -microglobulin amyloidogenic behavior. <i>PLoS Computational Biology</i> , 2014 , 10, e1003606	5	28

21	Treatment of Ionic Strength in Biomolecular Simulations of Charged Lipid Bilayers. <i>Journal of Chemical Theory and Computation</i> , 2014 , 10, 5483-92	6.4	17
20	Structural effects of pH and deacylation on surfactant protein C in an organic solvent mixture: a constant-pH MD study. <i>Journal of Chemical Information and Modeling</i> , 2013 , 53, 2979-89	6.1	19
19	Unraveling the Conformational Determinants of Peptide Dendrimers Using Molecular Dynamics Simulations. <i>Macromolecules</i> , 2013 , 46, 9427-9436	5.5	7
18	Charge parametrization of the DvH-c3 heme group: validation using constant-(pH,E) molecular dynamics simulations. <i>Journal of Physical Chemistry B</i> , 2013 , 117, 70-82	3.4	17
17	Tuning the Bioactivity of Tensioactive Deoxy Glycosides to Structure: Antibacterial Activity Versus Selective Cholinesterase Inhibition Rationalized by Molecular Docking. <i>European Journal of Organic Chemistry</i> , 2013 , 2013, 1448-1459	3.2	6
16	Conformational study of GSH and GSSG using constant-pH molecular dynamics simulations. <i>Journal of Physical Chemistry B</i> , 2013 , 117, 7507-17	3.4	24
15	Reversibility of prion misfolding: insights from constant-pH molecular dynamics simulations. <i>Journal of Physical Chemistry B</i> , 2012 , 116, 8812-21	3.4	33
14	Indenyl ring slippage in crown thioether complexes [IndMo(CO) ₂ L] ⁺ and C-S activation of trithiacyclononane: experimental and theoretical studies. <i>Dalton Transactions</i> , 2011 , 40, 10513-25	4.3	17
13	Is the prediction of pKa values by constant-pH molecular dynamics being hindered by inherited problems?. <i>Proteins: Structure, Function and Bioinformatics</i> , 2011 , 79, 3437-47	4.2	50
12	Unfolding the conformational behavior of peptide dendrimers: insights from molecular dynamics simulations. <i>Journal of the American Chemical Society</i> , 2011 , 133, 5042-52	16.4	30
11	Membrane-induced conformational changes of kyotorphin revealed by molecular dynamics simulations. <i>Journal of Physical Chemistry B</i> , 2010 , 114, 11659-67	3.4	19
10	Constant-pH molecular dynamics simulations reveal a β -rich form of the human prion protein. <i>Journal of Physical Chemistry B</i> , 2010 , 114, 12692-700	3.4	77
9	Molecular dynamics at constant pH and reduction potential: application to cytochrome c(3). <i>Journal of the American Chemical Society</i> , 2009 , 131, 12586-94	16.4	51
8	Acidic range titration of HEWL using a constant-pH molecular dynamics method. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008 , 72, 289-98	4.2	66
7	The pH-dependent conformational states of kyotorphin: a constant-pH molecular dynamics study. <i>Biophysical Journal</i> , 2007 , 92, 1836-45	2.9	44
6	Constant-pH molecular dynamics with ionic strength effects: protonation-conformation coupling in decalysine. <i>Journal of Physical Chemistry B</i> , 2006 , 110, 2927-33	3.4	93
5	On the use of different dielectric constants for computing individual and pairwise terms in poisson-boltzmann studies of protein ionization equilibrium. <i>Journal of Physical Chemistry B</i> , 2005 , 109, 14691-706	3.4	75
4	A Selective Direct Aldol Reaction in Aqueous Media Catalyzed by ZincProline. <i>European Journal of Organic Chemistry</i> , 2005 , 2005, 5268-5276	3.2	90

- 3 Zinc-proline catalyzed pathway for the formation of sugars. *Chemical Communications*, **2004**, 1540-1 5.8 88
- 2 Zinc mediated methyl transfer from trimethyl phosphate to chelating and non-chelating alkyl thiols. Model for Zn-dependent methyltransferases. *Journal of Inorganic Biochemistry*, **2003**, 94, 193-6 4.2 18
- 1 Zn-Proline catalyzed direct aldol reaction in aqueous media. *Chemical Communications*, **2003**, 1090-1091 5.8 165