

Laura Zanetti-Polzi

List of Publications by Year in descending order

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43
papers

848
citations

430754

18
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501076

28
g-index

48
all docs

48
docs citations

48
times ranked

1052
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | A general statistical mechanical model for fluid system thermodynamics: Application to sub- and super-critical water. <i>Journal of Chemical Physics</i> , 2022, 156, 044506. | 1.2 | 3 |
| 2 | Low- and high-density forms of liquid water revealed by a new medium-range order descriptor. <i>Journal of Molecular Liquids</i> , 2022, 355, 118922. | 2.3 | 11 |
| 3 | Photophysical and Computational Insights into Ag(I) Complexation of Porphyrinic Covalent Cages Equipped with Triazoles-Incorporating Linkers. <i>Journal of Physical Chemistry B</i> , 2022, 126, 3450-3459. | 1.2 | 0 |
| 4 | Evidence for a high pK_a of an aspartic acid residue in the active site of CALB by a fully atomistic multiscale approach. <i>Journal of Biomolecular Structure and Dynamics</i> , 2022, , 1-8. | 2.0 | 2 |
| 5 | A computational insight into the relationship between side chain IR line shapes and local environment in fibril-like structures. <i>Journal of Chemical Physics</i> , 2021, 154, 084105. | 1.2 | 1 |
| 6 | Tuning Proton Transfer Thermodynamics in SARS-CoV-2 Main Protease: Implications for Catalysis and Inhibitor Design. <i>Journal of Physical Chemistry Letters</i> , 2021, 12, 4195-4202. | 2.1 | 19 |
| 7 | Segregation on the nanoscale coupled to liquid water polyamorphism in supercooled aqueous ionic-liquid solution. <i>Journal of Chemical Physics</i> , 2021, 155, 104502. | 1.2 | 3 |
| 8 | Inhibitor binding influences the protonation states of histidines in SARS-CoV-2 main protease. <i>Chemical Science</i> , 2021, 12, 1513-1527. | 3.7 | 47 |
| 9 | Allosteric Control of Naphthalene Diimide Encapsulation and Electron Transfer in Porphyrin Containers: Photophysical Studies and Molecular Dynamics Simulation. <i>Chemistry - A European Journal</i> , 2020, 26, 17514-17524. | 1.7 | 7 |
| 10 | Supercomputer-Based Ensemble Docking Drug Discovery Pipeline with Application to Covid-19. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 5832-5852. | 2.5 | 134 |
| 11 | Fully Atomistic Multiscale Approach for pK_a Prediction. <i>Journal of Physical Chemistry B</i> , 2020, 124, 4712-4722. | 1.2 | 10 |
| 12 | Modeling amino-acid side chain infrared spectra: the case of carboxylic residues. <i>Physical Chemistry Chemical Physics</i> , 2020, 22, 3008-3016. | 1.3 | 10 |
| 13 | The Interaction between Amyloid Prefibrillar Oligomers of Salmon Calcitonin and a Lipid-Raft Model: Molecular Mechanisms Leading to Membrane Damage, Ca^{2+} -Influx and Neurotoxicity. <i>Biomolecules</i> , 2020, 10, 58. | 1.8 | 3 |
| 14 | Cooperative protein-solvent tuning of proton transfer energetics: carbonic anhydrase as a case study. <i>Physical Chemistry Chemical Physics</i> , 2020, 22, 19975-19981. | 1.3 | 3 |
| 15 | Hydration Shell of Antifreeze Proteins: Unveiling the Role of Non-Ice-Binding Surfaces. <i>Journal of Physical Chemistry B</i> , 2019, 123, 6474-6480. | 1.2 | 20 |
| 16 | Interpretation of Experimental Soret Bands of Porphyrins in Flexible Covalent Cages and in Their Related Ag(I) Fixed Complexes. <i>Journal of Physical Chemistry C</i> , 2019, 123, 13094-13103. | 1.5 | 17 |
| 17 | Evidence of a Thermodynamic Ramp for Hole Hopping to Protect a Redox Enzyme from Oxidative Damage. <i>Journal of Physical Chemistry Letters</i> , 2019, 10, 1450-1456. | 2.1 | 10 |
| 18 | Tip-Enhanced Infrared Difference-Nanospectroscopy of the Proton Pump Activity of Bacteriorhodopsin in Single Purple Membrane Patches. <i>Nano Letters</i> , 2019, 19, 3104-3114. | 4.5 | 36 |

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|----|--|-----|-----------|
| 19 | Difference mid-IR nanospectroscopy on individual patches of purple membranes: the proton pump activity of bacteriorhodopsin at the nanoscale. , 2019, , . | | 0 |
| 20 | On the nature of solvatochromic effect: The riboflavin absorption spectrum as a case study. Spectrochimica Acta - Part A: Molecular and Biomolecular Spectroscopy, 2018, 192, 451-457. | 2.0 | 8 |
| 21 | A quantitative connection of experimental and simulated folding landscapes by vibrational spectroscopy. Chemical Science, 2018, 9, 9002-9011. | 3.7 | 20 |
| 22 | Extending the perturbed matrix method beyond the dipolar approximation: comparison of different levels of theory. Physical Chemistry Chemical Physics, 2018, 20, 24369-24378. | 1.3 | 49 |
| 23 | Tyrosine absorption spectroscopy: Backbone protonation effects on the side chain electronic properties. Journal of Computational Chemistry, 2018, 39, 1747-1756. | 1.5 | 17 |
| 24 | Theoretical modeling of the absorption spectrum of aqueous riboflavin. Chemical Physics Letters, 2017, 669, 119-124. | 1.2 | 37 |
| 25 | Computational evidence support the hypothesis of neuroglobin also acting as an electron transfer species. Journal of Biological Inorganic Chemistry, 2017, 22, 615-623. | 1.1 | 24 |
| 26 | Computational investigation of the electron transfer complex between neuroglobin and cytochrome c. Supramolecular Chemistry, 2017, 29, 846-852. | 1.5 | 2 |
| 27 | Parallel folding pathways of Fip35 WW domain explained by infrared spectra and their computer simulation. FEBS Letters, 2017, 591, 3265-3275. | 1.3 | 12 |
| 28 | Alternative Electron-Transfer Channels Ensure Ultrafast Deactivation of Light-Induced Excited States in Riboflavin Binding Protein. Journal of Physical Chemistry Letters, 2017, 8, 3321-3327. | 2.1 | 21 |
| 29 | Theoretical-computational characterization of the temperature-dependent folding thermodynamics of α 2-hairpin peptide. Chemical Physics Letters, 2016, 659, 247-251. | 1.2 | 1 |
| 30 | Extending the essential dynamics analysis to investigate molecular properties: application to the redox potential of proteins. Physical Chemistry Chemical Physics, 2016, 18, 18450-18459. | 1.3 | 18 |
| 31 | A dynamical approach to non-adiabatic electron transfers at the bio-inorganic interface. Physical Chemistry Chemical Physics, 2016, 18, 10538-10549. | 1.3 | 11 |
| 32 | A few key residues determine the high redox potential shift in azurin mutants. Organic and Biomolecular Chemistry, 2015, 13, 11003-11013. | 1.5 | 35 |
| 33 | Surface Packing Determines the Redox Potential Shift of Cytochrome c Adsorbed on Gold. Journal of the American Chemical Society, 2014, 136, 12929-12937. | 6.6 | 39 |
| 34 | Simulation of the Amide I Infrared Spectrum in Photoinduced Peptide Folding/Unfolding Transitions. Journal of Physical Chemistry B, 2013, 117, 12383-12390. | 1.2 | 17 |
| 35 | Modeling triplet flavin-indole electron transfer and interradical dipolar interaction: a perturbative approach. Theoretical Chemistry Accounts, 2013, 132, 1. | 0.5 | 14 |
| 36 | A Theoretical Reappraisal of Polylysine in the Investigation of Secondary Structure Sensitivity of Infrared Spectra. Journal of Physical Chemistry B, 2012, 116, 3353-3360. | 1.2 | 19 |

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|----|---|-----|-----------|
| 37 | New Insight into the IR-Spectra/Structure Relationship in Amyloid Fibrils: A Theoretical Study on a Prion Peptide. <i>Journal of the American Chemical Society</i> , 2011, 133, 11414-11417. | 6.6 | 28 |
| 38 | Analysis of Infrared Spectra of \hat{I}^2 -Hairpin Peptides As Derived from Molecular Dynamics Simulations. <i>Journal of Physical Chemistry B</i> , 2011, 115, 11872-11878. | 1.2 | 11 |
| 39 | Structure of the lipodepsipeptide syringomycin E in phospholipids and sodium dodecylsulphate micelle studied by circular dichroism, NMR spectroscopy and molecular dynamics. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2011, 1808, 2102-2110. | 1.4 | 15 |
| 40 | Modeling quantum vibrational excitations in condensed-phase molecular systems. <i>Theoretical Chemistry Accounts</i> , 2011, 129, 31-43. | 0.5 | 26 |
| 41 | On the origin of IR spectral changes upon protein folding. <i>Chemical Physics Letters</i> , 2010, 488, 213-218. | 1.2 | 30 |
| 42 | Structural, thermodynamic, and kinetic properties of Gramicidin analogue GS6 studied by molecular dynamics simulations and statistical mechanics. <i>Biopolymers</i> , 2009, 91, 1154-1160. | 1.2 | 1 |
| 43 | Calcitonin Forms Oligomeric Pore-Like Structures in Lipid Membranes. <i>Biophysical Journal</i> , 2006, 91, 2275-2281. | 0.2 | 52 |