

Eleonora Gianti

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

351
citations

840776

11
h-index

1058476

14
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17
all docs

17
docs citations

17
times ranked

614
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Flipping in the Pore: Discovery of Dual Inhibitors That Bind in Different Orientations to the Wild-Type versus the Amantadine-Resistant S31N Mutant of the Influenza A Virus M2 Proton Channel. <i>Journal of the American Chemical Society</i> , 2014, 136, 17987-17995. | 13.7 | 78 |
| 2 | Does Proton Conduction in the Voltage-Gated H ⁺ Channel hHv1 Involve Grotthuss-Like Hopping via Acidic Residues?. <i>Journal of Physical Chemistry B</i> , 2017, 121, 3340-3351. | 2.6 | 34 |
| 3 | Hydrogen-Bonded Water Molecules in the M2 Channel of the Influenza A Virus Guide the Binding Preferences of Ammonium-Based Inhibitors. <i>Journal of Physical Chemistry B</i> , 2015, 119, 1173-1183. | 2.6 | 33 |
| 4 | On the role of water density fluctuations in the inhibition of a proton channel. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E8359-E8368. | 7.1 | 33 |
| 5 | Computational analysis of EBNA1 "druggability" suggests novel insights for Epstein-Barr virus inhibitor design. <i>Journal of Computer-Aided Molecular Design</i> , 2016, 30, 285-303. | 2.9 | 27 |
| 6 | Sites Contributing to TRPA1 Activation by the Anesthetic Propofol Identified by Photoaffinity Labeling. <i>Biophysical Journal</i> , 2017, 113, 2168-2172. | 0.5 | 26 |
| 7 | Identification and Selection of "Privileged Fragments" Suitable for Primary Screening. <i>Journal of Chemical Information and Modeling</i> , 2008, 48, 2129-2139. | 5.4 | 25 |
| 8 | Suppression of Zika Virus Infection in the Brain by the Antiretroviral Drug Rilpivirine. <i>Molecular Therapy</i> , 2019, 27, 2067-2079. | 8.2 | 20 |
| 9 | TRPA1 modulation by piperidine carboxamides suggests an evolutionarily conserved binding site and gating mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 26008-26019. | 7.1 | 18 |
| 10 | Fragment-based Shape Signatures: a new tool for virtual screening and drug discovery. <i>Journal of Computer-Aided Molecular Design</i> , 2013, 27, 1009-1036. | 2.9 | 16 |
| 11 | Dual regulation of TRPV1 channels by phosphatidylinositol via functionally distinct binding sites. <i>Journal of Biological Chemistry</i> , 2021, 296, 100573. | 3.4 | 16 |
| 12 | An SH2 domain model of STAT5 in complex with phospho-peptides define "STAT5 Binding Signatures". <i>Journal of Computer-Aided Molecular Design</i> , 2015, 29, 451-470. | 2.9 | 11 |
| 13 | Modeling Androgen Receptor Flexibility: A Binding Mode Hypothesis of CYP17 Inhibitors/Antiandrogens for Prostate Cancer Therapy. <i>Journal of Chemical Information and Modeling</i> , 2012, 52, 2670-2683. | 5.4 | 9 |
| 14 | Computational Approaches to Studying Voltage-Gated Ion Channel Modulation by General Anesthetics. <i>Methods in Enzymology</i> , 2018, 602, 25-59. | 1.0 | 3 |
| 15 | Structure-activity relationships and drug design. , 2021, , 129-153. | | 2 |