

# Cesar A LÃ³pez

## List of Publications by Year in descending order

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Version: 2024-02-01

39  
papers

3,881  
citations

257357

24  
h-index

302012

39  
g-index

43  
all docs

43  
docs citations

43  
times ranked

6200  
citing authors

| #  | ARTICLE                                                                                                                                                                                                             | IF  | CITATIONS |
|----|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 1  | Machine learning-driven multiscale modeling reveals lipid-dependent dynamics of RAS signaling proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .       | 3.3 | 44        |
| 2  | Exploring CRD mobility during RAS/RAF engagement at the membrane. <i>Biophysical Journal</i> , 2022, 121, 3630-3650.                                                                                                | 0.2 | 9         |
| 3  | Predictive Rules of Efflux Inhibition and Avoidance in <i>Pseudomonas aeruginosa</i> . <i>MBio</i> , 2021, 12, .                                                                                                    | 1.8 | 28        |
| 4  | Molecular origins of reduced activity and binding commitment of processive cellulases and associated carbohydrate-binding proteins to cellulose III. <i>Journal of Biological Chemistry</i> , 2021, 296, 100431.    | 1.6 | 20        |
| 5  | Development of Martini 2.2 parameters for N-glycans: a case study of the HIV-1 Env glycoprotein dynamics. <i>Glycobiology</i> , 2021, 31, 787-799.                                                                  | 1.3 | 7         |
| 6  | Unveiling the Dynamics of KRAS4b on Lipid Model Membranes. <i>Journal of Membrane Biology</i> , 2021, 254, 201-216.                                                                                                 | 1.0 | 6         |
| 7  | Molecular characterization of the outer membrane of <i>Pseudomonas aeruginosa</i> . <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2020, 1862, 183151.                                                       | 1.4 | 28        |
| 8  | Visualization of the HIV-1 Env glycan shield across scales. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 28014-28025.                                        | 3.3 | 57        |
| 9  | Uncovering a membrane-distal conformation of KRAS available to recruit RAF to the plasma membrane. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 24258-24268. | 3.3 | 34        |
| 10 | Machine Learning Algorithm Identifies an Antibiotic Vocabulary for Permeating Gram-Negative Bacteria. <i>Journal of Chemical Information and Modeling</i> , 2020, 60, 2838-2847.                                    | 2.5 | 21        |
| 11 | Anionic Lipids Impact RAS-Binding Site Accessibility and Membrane Binding Affinity of CRAF RBD-CRD. <i>Biophysical Journal</i> , 2020, 119, 525-538.                                                                | 0.2 | 13        |
| 12 | Two distinct anionic phospholipid-dependent events involved in SecA-mediated protein translocation. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2019, 1861, 183035.                                       | 1.4 | 16        |
| 13 | Oligomeric state of the ZIKV E protein defines protective immune responses. <i>Nature Communications</i> , 2019, 10, 4606.                                                                                          | 5.8 | 22        |
| 14 | Efficient transplacental IgG transfer in women infected with Zika virus during pregnancy. <i>PLoS Neglected Tropical Diseases</i> , 2019, 13, e0007648.                                                             | 1.3 | 22        |
| 15 | Unsupervised Machine Learning for Analysis of Phase Separation in Ternary Lipid Mixture. <i>Journal of Chemical Theory and Computation</i> , 2019, 15, 6343-6357.                                                   | 2.3 | 18        |
| 16 | Biophysical Characterization of a Nanodisc with and without BAX: An Integrative Study Using Molecular Dynamics Simulations and Cryo-EM. <i>Structure</i> , 2019, 27, 988-999.e4.                                    | 1.6 | 19        |
| 17 | Sequence- and structure-based computational analyses of Gram-negative tripartite efflux pumps in the context of bacterial membranes. <i>Research in Microbiology</i> , 2018, 169, 414-424.                          | 1.0 | 6         |
| 18 | Development of Envelope Protein Antigens To Serologically Differentiate Zika Virus Infection from Dengue Virus Infection. <i>Journal of Clinical Microbiology</i> , 2018, 56, .                                     | 1.8 | 53        |

| #  | ARTICLE                                                                                                                                                                                                                            | IF  | CITATIONS |
|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 19 | Capturing Phase Behavior of Ternary Lipid Mixtures with a Refined Martini Coarse-Grained Force Field. <i>Journal of Chemical Theory and Computation</i> , 2018, 14, 6050-6062.                                                     | 2.3 | 63        |
| 20 | Molecular recognition of RAS/RAF complex at the membrane: Role of RAF cysteine-rich domain. <i>Scientific Reports</i> , 2018, 8, 8461.                                                                                             | 1.6 | 71        |
| 21 | Membrane perturbing properties of toxin mycolactone from <i>Mycobacterium ulcerans</i> . <i>PLoS Computational Biology</i> , 2018, 14, e1005972.                                                                                   | 1.5 | 28        |
| 22 | Dynamics of Intact MexAB-OprM Efflux Pump: Focusing on the MexA-OprM Interface. <i>Scientific Reports</i> , 2017, 7, 16521.                                                                                                        | 1.6 | 30        |
| 23 | Lack of Durable Cross-Neutralizing Antibodies Against Zika Virus from Dengue Virus Infection. <i>Emerging Infectious Diseases</i> , 2017, 23, 773-781.                                                                             | 2.0 | 141       |
| 24 | Effect of Glycosylation on an Immunodominant Region in the V1V2 Variable Domain of the HIV-1 Envelope gp120 Protein. <i>PLoS Computational Biology</i> , 2016, 12, e1005094.                                                       | 1.5 | 17        |
| 25 | Broadly targeted CD8 <sup>+</sup> T cell responses restricted by major histocompatibility complex E. <i>Science</i> , 2016, 351, 714-720.                                                                                          | 6.0 | 260       |
| 26 | Membrane-Mediated Regulation of the Intrinsically Disordered CD3 $\mu$ Cytoplasmic Tail of the TCR. <i>Biophysical Journal</i> , 2015, 108, 2481-2491.                                                                             | 0.2 | 21        |
| 27 | MARTINI Coarse-Grained Model for Crystalline Cellulose Microfibers. <i>Journal of Physical Chemistry B</i> , 2015, 119, 465-473.                                                                                                   | 1.2 | 54        |
| 28 | Permeability Barrier of Gram-Negative Cell Envelopes and Approaches To Bypass It. <i>ACS Infectious Diseases</i> , 2015, 1, 512-522.                                                                                               | 1.8 | 442       |
| 29 | Residue Leu940 Has a Crucial Role in the Linkage and Reaction Specificity of the Glucanucrase GTF180 of the Probiotic Bacterium <i>Lactobacillus reuteri</i> 180. <i>Journal of Biological Chemistry</i> , 2014, 289, 32773-32782. | 1.6 | 33        |
| 30 | The power of coarse graining in biomolecular simulations. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2014, 4, 225-248.                                                                              | 6.2 | 423       |
| 31 | Lipid Organization of the Plasma Membrane. <i>Journal of the American Chemical Society</i> , 2014, 136, 14554-14559.                                                                                                               | 6.6 | 734       |
| 32 | Disaccharides Impact the Lateral Organization of Lipid Membranes. <i>Journal of the American Chemical Society</i> , 2014, 136, 16167-16175.                                                                                        | 6.6 | 55        |
| 33 | Martini Force Field Parameters for Glycolipids. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 1694-1708.                                                                                                            | 2.3 | 166       |
| 34 | Computational microscopy of cyclodextrin mediated cholesterol extraction from lipid model membranes. <i>Scientific Reports</i> , 2013, 3, 2071.                                                                                    | 1.6 | 101       |
| 35 | Molecular view on protein sorting into liquid-ordered membrane domains mediated by gangliosides and lipid anchors. <i>Faraday Discussions</i> , 2013, 161, 347-363.                                                                | 1.6 | 76        |
| 36 | Amylose folding under the influence of lipids. <i>Carbohydrate Research</i> , 2012, 364, 1-7.                                                                                                                                      | 1.1 | 72        |

| #  | ARTICLE                                                                                                                                                                                                                                 | IF  | CITATIONS |
|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 37 | Molecular Mechanism of Cyclodextrin Mediated Cholesterol Extraction. PLoS Computational Biology, 2011, 7, e1002020.                                                                                                                     | 1.5 | 165       |
| 38 | Crystal structure of a 117 kDa glucansucrase fragment provides insight into evolution and product specificity of GH70 enzymes. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 21406-21411. | 3.3 | 140       |
| 39 | Martini Coarse-Grained Force Field: Extension to Carbohydrates. Journal of Chemical Theory and Computation, 2009, 5, 3195-3210.                                                                                                         | 2.3 | 363       |