

Lluís Raich

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

1,107
citations

471509

17
h-index

610901

24
g-index

26
all docs

26
docs citations

26
times ranked

1946
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular mechanism of inhibiting the SARS-CoV-2 cell entry facilitator TMPRSS2 with camostat and nafamostat. <i>Chemical Science</i> , 2021, 12, 983-992.	7.4	66
2	Discovery of a hidden transient state in all bromodomain families. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	24
3	Synergistic inhibition of SARS-CoV-2 cell entry by otamixaban and covalent protease inhibitors: pre-clinical assessment of pharmacological and molecular properties. <i>Chemical Science</i> , 2021, 12, 12600-12609.	7.4	11
4	Camostat mesylate inhibits SARS-CoV-2 activation by TMPRSS2-related proteases and its metabolite GBPA exerts antiviral activity. <i>EBioMedicine</i> , 2021, 65, 103255.	6.1	256
5	Alpha 1 Antitrypsin is an Inhibitor of the SARS-CoV-2â€œPriming Protease TMPRSS2. <i>Pathogens and Immunity</i> , 2021, 6, 55-74.	3.1	73
6	A Single Point Mutation Converts GH84 <i>GlcNAc</i> Hydrolases into Phosphorylases: Experimental and Theoretical Evidence. <i>Journal of the American Chemical Society</i> , 2020, 142, 2120-2124.	13.7	25
7	An Epoxide Intermediate in Glycosidase Catalysis. <i>ACS Central Science</i> , 2020, 6, 760-770.	11.3	34
8	Î±-Gal-cyclophellitol cyclosulfamidate is a Michaelis complex analog that stabilizes therapeutic lysosomal Î±-galactosidase A in Fabry disease. <i>Chemical Science</i> , 2019, 10, 9233-9243.	7.4	11
9	Modeling catalytic reaction mechanisms in glycoside hydrolases. <i>Current Opinion in Chemical Biology</i> , 2019, 53, 183-191.	6.1	26
10	Dynamic and Functional Profiling of Xylan-Degrading Enzymes in <i>Aspergillus</i> Secretomes Using Activity-Based Probes. <i>ACS Central Science</i> , 2019, 5, 1067-1078.	11.3	34
11	The Molecular Mechanism of Substrate Recognition and Catalysis of the Membrane Acyltransferase PatA from <i>Mycobacteria</i> . <i>ACS Chemical Biology</i> , 2018, 13, 131-140.	3.4	10
12	Palladium-mediated enzyme activation suggests multiphase initiation of glycogenesis. <i>Nature</i> , 2018, 563, 235-240.	27.8	42
13	Molecular-Scale Ligand Effects in Small Goldâ€œThiolate Nanoclusters. <i>Journal of the American Chemical Society</i> , 2018, 140, 15430-15436.	13.7	90
14	Structural and Mechanistic Insights into the Catalytic-Domain-Mediated Short-Range Glycosylation Preferences of GalNAc-T4. <i>ACS Central Science</i> , 2018, 4, 1274-1290.	11.3	35
15	The molecular mechanism of the ligand exchange reaction of an antibody against a glutathione-coated gold cluster. <i>Nanoscale</i> , 2017, 9, 3121-3127.	5.6	15
16	Conformational Analysis of the Mannosidase Inhibitor Kifunensine: A Quantum Mechanical and Structural Approach. <i>ChemBioChem</i> , 2017, 18, 1496-1501.	2.6	12
17	Carba-cyclophellitols Are Neutral Retaining-Glucosidase Inhibitors. <i>Journal of the American Chemical Society</i> , 2017, 139, 6534-6537.	13.7	24
18	Contribution of Shape and Charge to the Inhibition of a Family GH99 <i>endo</i> -Î±-1,2-Mannanase. <i>Journal of the American Chemical Society</i> , 2017, 139, 1089-1097.	13.7	17

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19	Precise Probing of Residue Roles by Post-Translational $\hat{1}^2, \hat{1}^3$ -C,N Aza-Michael Mutagenesis in Enzyme Active Sites. ACS Central Science, 2017, 3, 1168-1173.	11.3	30
20	1,6-Cyclophellitol Cyclosulfates: A New Class of Irreversible Glycosidase Inhibitor. ACS Central Science, 2017, 3, 784-793.	11.3	43
21	Selective Derivatization of <i>N</i> -Terminal Cysteines Using Cyclopentenediones. Organic Letters, 2016, 18, 4836-4839.	4.6	10
22	A $\hat{1}^2$ -Mannanase with a Lysozyme-like Fold and a Novel Molecular Catalytic Mechanism. ACS Central Science, 2016, 2, 896-903.	11.3	39
23	A Trapped Covalent Intermediate of a Glycoside Hydrolase on the Pathway to Transglycosylation. Insights from Experiments and Quantum Mechanics/Molecular Mechanics Simulations. Journal of the American Chemical Society, 2016, 138, 3325-3332.	13.7	47
24	The complete conformational free energy landscape of $\hat{1}^2$ -xylose reveals a two-fold catalytic itinerary for $\hat{1}^2$ -xylanases. Chemical Science, 2015, 6, 1167-1177.	7.4	44