

Gabriele Cerutti

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

3,392
citations

687363
13
h-index

713466
21
g-index

38
all docs

38
docs citations

38
times ranked

7044
citing authors

#	ARTICLE	IF	CITATIONS
1	A monoclonal antibody that neutralizes SARS-CoV-2 variants, SARS-CoV, and other sarbecoviruses. <i>Emerging Microbes and Infections</i> , 2022, 11, 147-157.	6.5	25
2	Cryo-EM structure of the SARS-CoV-2 Omicron spike. <i>Cell Reports</i> , 2022, 38, 110428.	6.4	82
3	Point Mutations at a Key Site Alter the Cytochrome P450 OleP Structural Dynamics. <i>Biomolecules</i> , 2022, 12, 55.	4.0	6
4	An antibody class with a common CDRH3 motif broadly neutralizes sarbecoviruses. <i>Science Translational Medicine</i> , 2022, 14, eabn6859.	12.4	31
5	Modular basis for potent SARS-CoV-2 neutralization by a prevalent VH1-2-derived antibody class. <i>Cell Reports</i> , 2021, 35, 108950.	6.4	54
6	Increased resistance of SARS-CoV-2 variant P.1 to antibody neutralization. <i>Cell Host and Microbe</i> , 2021, 29, 747-751.e4.	11.0	504
7	Potent SARS-CoV-2 neutralizing antibodies directed against spike N-terminal domain target a single supersite. <i>Cell Host and Microbe</i> , 2021, 29, 819-833.e7.	11.0	444
8	Gating movements and ion permeation in HCN4 pacemaker channels. <i>Molecular Cell</i> , 2021, 81, 2929-2943.e6.	9.7	41
9	Structural basis for accommodation of emerging B.1.351 and B.1.1.7 variants by two potent SARS-CoV-2 neutralizing antibodies. <i>Structure</i> , 2021, 29, 655-663.e4.	3.3	52
10	Structural Studies of an Anti-SARS-CoV-2 Antibody Cocktail. <i>Microscopy and Microanalysis</i> , 2021, 27, 2844-2846.	0.4	0
11	Crystal structure and functional characterization of an oligosaccharide dehydrogenase from <i>Pycnoporus cinnabarinus</i> provides insights into fungal breakdown of lignocellulose. <i>Biotechnology for Biofuels</i> , 2021, 14, 161.	6.2	7
12	Paired heavy- and light-chain signatures contribute to potent SARS-CoV-2 neutralization in public antibody responses. <i>Cell Reports</i> , 2021, 37, 109771.	6.4	38
13	Neutralizing antibody 5-7 defines a distinct site of vulnerability in SARS-CoV-2 spike N-terminal domain. <i>Cell Reports</i> , 2021, 37, 109928.	6.4	52
14	Structure-Based Design with Tag-Based Purification and In-Process Biotinylation Enable Streamlined Development of SARS-CoV-2 Spike Molecular Probes. <i>Cell Reports</i> , 2020, 33, 108322.	6.4	59
15	Cryo-EM Structures of SARS-CoV-2 Spike without and with ACE2 Reveal a pH-Dependent Switch to Mediate Endosomal Positioning of Receptor-Binding Domains. <i>Cell Host and Microbe</i> , 2020, 28, 867-879.e5.	11.0	316
16	Biodistribution PET/CT Study of Hemoglobin-DFO-89Zr Complex in Healthy and Lung Tumor-Bearing Mice. <i>International Journal of Molecular Sciences</i> , 2020, 21, 4991.	4.1	1
17	Potent neutralizing antibodies against multiple epitopes on SARS-CoV-2 spike. <i>Nature</i> , 2020, 584, 450-456.	27.8	1,337
18	Dissecting the Cytochrome P450 OleP Substrate Specificity: Evidence for a Preferential Substrate. <i>Biomolecules</i> , 2020, 10, 1411.	4.0	6

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19	Structure-Based Design with Tag-Based Purification and In-Process Biotinylation Enable Streamlined Development of SARS-CoV-2 Spike Molecular Probes. SSRN Electronic Journal, 2020, , 3639618.	0.4	3
20	Proximal and distal control for ligand binding in neuroglobin: role of the CD loop and evidence for His64 gating. Scientific Reports, 2019, 9, 5326.	3.3	10
21	Substrate-induced conformational change in cytochrome P450 OleP. FASEB Journal, 2019, 33, 1787-1800.	0.5	14
22	Subcellular localization of the five members of the human steroid 5 β -reductase family. Biochimie Open, 2017, 4, 99-106.	3.2	11
23	Cryo-EM Structures Delineate a pH-Dependent Switch that Mediates Endosomal Positioning of SARS-CoV-2 Spike Receptor-Binding Domains. SSRN Electronic Journal, 0, , .	0.4	6
24	Paired Heavy and Light Chain Signatures Contribute to Potent SARS-CoV-2 Neutralization in Public Antibody Responses. SSRN Electronic Journal, 0, , .	0.4	1
25	Gating Movements and Ion Permeation in HCN4 Pacemaker Channels. SSRN Electronic Journal, 0, , .	0.4	0
26	Probing the Role of Murine Neuroglobin CDloop-D-Helix Unit in CO Ligand Binding and Structural Dynamics. ACS Chemical Biology, 0, , .	3.4	2