

Lauren Carter

List of Publications by Citations

Source: <https://exaly.com/author-pdf/7103310/lauren-carter-publications-by-citations.pdf>

Version: 2024-04-24

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

39
papers

2,290
citations

16
h-index

47
g-index

47
ext. papers

3,839
ext. citations

26.4
avg, IF

4.9
L-index

#	Paper	IF	Citations
39	Functional SARS-CoV-2-Specific Immune Memory Persists after Mild COVID-19. <i>Cell</i> , 2021 , 184, 169-183.e17	56.2	327
38	Global analysis of protein folding using massively parallel design, synthesis, and testing. <i>Science</i> , 2017 , 357, 168-175	33.3	241
37	Massively parallel de novo protein design for targeted therapeutics. <i>Nature</i> , 2017 , 550, 74-79	50.4	235
36	De novo design of picomolar SARS-CoV-2 miniprotein inhibitors. <i>Science</i> , 2020 , 370, 426-431	33.3	219
35	Elicitation of Potent Neutralizing Antibody Responses by Designed Protein Nanoparticle Vaccines for SARS-CoV-2. <i>Cell</i> , 2020 , 183, 1367-1382.e17	56.2	217
34	Induction of Potent Neutralizing Antibody Responses by a Designed Protein Nanoparticle Vaccine for Respiratory Syncytial Virus. <i>Cell</i> , 2019 , 176, 1420-1431.e17	56.2	190
33	De novo design of potent and selective mimics of IL-2 and IL-15. <i>Nature</i> , 2019 , 565, 186-191	50.4	184
32	De novo design of a fluorescence-activating E-barrel. <i>Nature</i> , 2018 , 561, 485-491	50.4	156
31	Adjuvanting a subunit COVID-19 vaccine to induce protective immunity. <i>Nature</i> , 2021 , 594, 253-258	50.4	92
30	Molecular basis of immune evasion by the Delta and Kappa SARS-CoV-2 variants. <i>Science</i> , 2021 , eabl8506	33.3	65
29	De novo design of a non-local E-sheet protein with high stability and accuracy. <i>Nature Structural and Molecular Biology</i> , 2018 , 25, 1028-1034	17.6	54
28	Quadrivalent influenza nanoparticle vaccines induce broad protection. <i>Nature</i> , 2021 , 592, 623-628	50.4	40
27	A Computationally Designed Hemagglutinin Stem-Binding Protein Provides In Vivo Protection from Influenza Independent of a Host Immune Response. <i>PLoS Pathogens</i> , 2016 , 12, e1005409	7.6	36
26	Elicitation of broadly protective sarbecovirus immunity by receptor-binding domain nanoparticle vaccines. <i>Cell</i> , 2021 , 184, 5432-5447.e16	56.2	34
25	De novo protein design by deep network hallucination. <i>Nature</i> , 2021 ,	50.4	33
24	F-domain valency determines outcome of signaling through the angiotensin pathway 2020 ,		28
23	Engineered SARS-CoV-2 receptor binding domain improves manufacturability in yeast and immunogenicity in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	13

22	Elicitation of broadly protective sarbecovirus immunity by receptor-binding domain nanoparticle vaccines 2021 ,		12
21	Ultrapotent miniproteins targeting the SARS-CoV-2 receptor-binding domain protect against infection and disease. <i>Cell Host and Microbe</i> , 2021 , 29, 1151-1161.e5	23.4	11
20	Elicitation of potent neutralizing antibody responses by designed protein nanoparticle vaccines for SARS-CoV-2 2020 ,		10
19	Engineered SARS-CoV-2 receptor binding domain improves immunogenicity in mice and elicits protective immunity in hamsters 2021 ,		10
18	In silico detection of SARS-CoV-2 specific B-cell epitopes and validation in ELISA for serological diagnosis of COVID-19. <i>Scientific Reports</i> , 2021 , 11, 4290	4.9	9
17	Computational design of a synthetic PD-1 agonist. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	9
16	Functional SARS-CoV-2-specific immune memory persists after mild COVID-19 2020 ,		7
15	Incorporation of sensing modalities into de novo designed fluorescence-activating proteins. <i>Nature Communications</i> , 2021 , 12, 856	17.4	7
14	Structure-based Design of JOC-x, a Conjugatable Tumor Tight Junction Opener to Enhance Cancer Therapy. <i>Scientific Reports</i> , 2019 , 9, 6169	4.9	6
13	Characterizing protein G B1 orientation and its effect on immunoglobulin G antibody binding using XPS, ToF-SIMS, and quartz crystal microbalance with dissipation monitoring. <i>Biointerphases</i> , 2020 , 15, 021002	1.8	6
12	F-domain valency determines outcome of signaling through the angiotensin pathway. <i>EMBO Reports</i> , 2021 , 22, e53471	6.5	4
11	Sampling of Structure and Sequence Space of Small Protein Folds		4
10	SARS-COV-2 spike binding to ACE2 in living cells monitored by TR-FRET. <i>Cell Chemical Biology</i> , 2021 ,	8.2	4
9	Multivalent designed proteins protect against SARS-CoV-2 variants of concern 2021 ,		4
8	High-affinity, neutralizing antibodies to SARS-CoV-2 can be made without T follicular helper cells. <i>Science Immunology</i> , 2022 , 7,	28	3
7	Airway antibodies emerge according to COVID-19 severity and wane rapidly but reappear after SARS-CoV-2 vaccination. <i>JCI Insight</i> , 2021 , 6,	9.9	3
6	Multivalent designed proteins neutralize SARS-CoV-2 variants of concern and confer protection against infection in mice.. <i>Science Translational Medicine</i> , 2022 , 14, eabn1252	17.5	3
5	Qualification of ELISA and neutralization methodologies to measure SARS-CoV-2 humoral immunity using human clinical samples. <i>Journal of Immunological Methods</i> , 2021 , 499, 113160	2.5	2

4	High-affinity, neutralizing antibodies to SARS-CoV-2 can be made without T follicular helper cells.. <i>Science Immunology</i> , 2021 , eabl5652	28	2
3	Ultrapotent miniproteins targeting the receptor-binding domain protect against SARS-CoV-2 infection and disease in mice 2021 ,		1
2	Detection of antibodies neutralizing historical and emerging SARS-CoV-2 strains using a thermodynamically coupled de novo biosensor system 2021 ,		1
1	Qualification of ELISA and neutralization methodologies to measure SARS-CoV-2 humoral immunity using human clinical samples 2021 ,		1