Shion A Lim

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10683412/publications.pdf

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		1040056	1281871	
11	725	9	11	
papers	citations	h-index	g-index	
15	15	15	1665	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Engineered ACE2 receptor traps potently neutralize SARS-CoV-2. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 28046-28055.	7.1	219
2	The thermostability and specificity of ancient proteins. Current Opinion in Structural Biology, 2016, 38, 37-43.	5.7	106
3	Engineering luminescent biosensors for point-of-care SARS-CoV-2 antibody detection. Nature Biotechnology, 2021, 39, 928-935.	17.5	106
4	Bi-paratopic and multivalent VH domains block ACE2 binding and neutralize SARS-CoV-2. Nature Chemical Biology, 2021, 17, 113-121.	8.0	78
5	Competitive SARS-CoV-2 Serology Reveals Most Antibodies Targeting the Spike Receptor-Binding Domain Compete for ACE2 Binding. MSphere, 2020, 5, .	2.9	62
6	Evolutionary trend toward kinetic stability in the folding trajectory of RNases H. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 13045-13050.	7.1	37
7	Theranostic Targeting of CUB Domain Containing Protein 1 (CDCP1) in Pancreatic Cancer. Clinical Cancer Research, 2020, 26, 3608-3615.	7.0	24
8	Bispecific VH/Fab antibodies targeting neutralizing and non-neutralizing Spike epitopes demonstrate enhanced potency against SARS-CoV-2. MAbs, 2021, 13, 1893426.	5 . 2	22
9	Targeting a proteolytic neoepitope on CUB domain containing protein 1 (CDCP1) for RAS-driven cancers. Journal of Clinical Investigation, 2022, 132, .	8.2	13
10	The burstâ€phase folding intermediate of ribonuclease H changes conformation over evolutionary history. Biopolymers, 2018, 109, e23086.	2.4	8
11	Exploring the Evolutionary History of Kinetic Stability in the α-Lytic Protease Family. Biochemistry, 2021, 60, 170-181.	2.5	2