Jungsoon Lee

List of Publications by Year in descending order

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	687363	940533
687	13	16
citations	h-index	g-index
19	19	1101
docs citations	times ranked	citing authors
	citations 19	687 13 citations h-index 19 19

#	Article	IF	CITATIONS
1	Yeast-expressed recombinant SARS-CoV-2 receptor binding domain RBD203-N1 as a COVID-19 protein vaccine candidate. Protein Expression and Purification, 2022, 190, 106003.	1.3	21
2	Receptor-binding domain recombinant protein on alum-CpG induces broad protection against SARS-CoV-2 variants of concern. Vaccine, 2022, 40, 3655-3663.	3.8	21
3	SARSâ€'CoV-2 RBD219-N1C1: A yeast-expressed SARS-CoV-2 recombinant receptor-binding domain candidate vaccine stimulates virus neutralizing antibodies and T-cell immunity in mice. Human Vaccines and Immunotherapeutics, 2021, 17, 2356-2366.	3.3	64
4	Process development and scale-up optimization of the SARS-CoV-2 receptor binding domain–based vaccine candidate, RBD219-N1C1. Applied Microbiology and Biotechnology, 2021, 105, 4153-4165.	3.6	37
5	Genetic modification to design a stable yeast-expressed recombinant SARS-CoV-2 receptor binding domain as a COVID-19 vaccine candidate. Biochimica Et Biophysica Acta - General Subjects, 2021, 1865, 129893.	2.4	49
6	A yeast-expressed RBD-based SARS-CoV-2 vaccine formulated with 3M-052-alum adjuvant promotes protective efficacy in non-human primates. Science Immunology, 2021, 6, .	11.9	53
7	The mitochondrial HSP90 paralog TRAP1 forms an OXPHOS-regulated tetramer and is involved in mitochondrial metabolic homeostasis. BMC Biology, 2020, 18, 10.	3.8	56
8	Crystal Structure of the YcjX Stress Protein Reveals a Ras-Like GTP-Binding Protein. Journal of Molecular Biology, 2019, 431, 3179-3190.	4.2	3
9	Cryo-EM Structures of the Hsp104 Protein Disaggregase Captured in the ATP Conformation. Cell Reports, 2019, 26, 29-36.e3.	6.4	36
10	Mapping interactions with the chaperone network reveals factors that protect against tau aggregation. Nature Structural and Molecular Biology, 2018, 25, 384-393.	8.2	119
11	Overlapping and Specific Functions of the Hsp104 N Domain Define Its Role in Protein Disaggregation. Scientific Reports, 2017, 7, 11184.	3.3	15
12	Structural determinants for protein unfolding and translocation by the Hsp104 protein disaggregase. Bioscience Reports, 2017, 37, .	2.4	15
13	2.4â€Ã resolution crystal structure of human TRAP1 _{NM} , the Hsp90 paralog in the mitochondrial matrix. Acta Crystallographica Section D: Structural Biology, 2016, 72, 904-911.	2.3	16
14	Mitochondrial Hsp90 is a ligand-activated molecular chaperone coupling ATP binding to dimer closure through a coiled-coil intermediate. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 2952-2957.	7.1	42
15	Molecular chaperones: guardians of the proteome in normal and disease states. F1000Research, 2015, 4, 1448.	1.6	39
16	Heat shock protein (Hsp) 70 is an activator of the Hsp104 motor. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 8513-8518.	7.1	101