Richard R Stein

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4842374/publications.pdf

Version: 2024-02-01

1163117 1588992 2,434 12 8 citations h-index g-index papers

15 15 15 3736 docs citations times ranked citing authors all docs

8

#	Article	IF	CITATIONS
1	Protein Structure from Experimental Evolution. Cell Systems, 2020, 10, 15-24.e5.	6.2	39
2	Perturbation biology links temporal protein changes to drug responses in a melanoma cell line. PLoS Computational Biology, 2020, 16, e1007909.	3.2	15
3	High-Dimensional Analysis of Immune Cell Composition Predicts Periprosthetic Joint Infections and Dissects Its Pathophysiology. Biomedicines, 2020, 8, 358.	3.2	12
4	Perturbation biology links temporal protein changes to drug responses in a melanoma cell line. , 2020, 16, e1007909.		0
5	Perturbation biology links temporal protein changes to drug responses in a melanoma cell line. , 2020, 16, e1007909.		0
6	Perturbation biology links temporal protein changes to drug responses in a melanoma cell line. , 2020, 16, e1007909.		0
7	Perturbation biology links temporal protein changes to drug responses in a melanoma cell line. , 2020, 16, e1007909.		0
8	Computer-guided design of optimal microbial consortia for immune system modulation. ELife, 2018, 7, .	6.0	65
9	MDSINE: Microbial Dynamical Systems INference Engine for microbiome time-series analyses. Genome Biology, 2016, 17, 121.	8.8	209
10	Precision microbiome reconstitution restores bile acid mediated resistance to Clostridium difficile. Nature, 2015, 517, 205-208.	27.8	1,506
11	Inferring Pairwise Interactions from Biological Data Using Maximum-Entropy Probability Models. PLoS Computational Biology, 2015, 11, e1004182.	3.2	97
12	Ecological Modeling from Time-Series Inference: Insight into Dynamics and Stability of Intestinal Microbiota. PLoS Computational Biology, 2013, 9, e1003388.	3.2	487