

Erik Abner

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5520345/publications.pdf>

Version: 2024-02-01

12
papers

375
citations

1163117

8
h-index

1372567

10
g-index

17
all docs

17
docs citations

17
times ranked

509
citing authors

#	ARTICLE	IF	CITATIONS
1	Mendelian Randomization Analysis Identifies Blood Tyrosine Levels as a Biomarker of Non-Alcoholic Fatty Liver Disease. <i>Metabolites</i> , 2022, 12, 440.	2.9	15
2	Comprehensive genome-wide association study of different forms of hernia identifies more than 80 associated loci. <i>Nature Communications</i> , 2022, 13, .	12.8	9
3	Genome-wide association study identifies five risk loci for pernicious anemia. <i>Nature Communications</i> , 2021, 12, 3761.	12.8	27
4	Electronic health record-based genome-wide meta-analysis provides insights on the genetic architecture of non-alcoholic fatty liver disease. <i>Cell Reports Medicine</i> , 2021, 2, 100437.	6.5	56
5	Genetic and InÂVitro Inhibition of PCSK9 and Calcific Aortic Valve Stenosis. <i>JACC Basic To Translational Science</i> , 2020, 5, 649-661.	4.1	45
6	Genome-wide Study Identifies Association between HLA-B*55:01 and Self-Reported Penicillin Allergy. <i>American Journal of Human Genetics</i> , 2020, 107, 612-621.	6.2	34
7	A broad drug arsenal to attack a strenuous latent HIV reservoir. <i>Current Opinion in Virology</i> , 2019, 38, 37-53.	5.4	23
8	HIV shock and kill therapy: In need of revision. <i>Antiviral Research</i> , 2019, 166, 19-34.	4.1	117
9	A New Quinoline BRD4 Inhibitor Targets a Distinct Latent HIV-1 Reservoir for Reactivation from Other Shock-Drugs. <i>Journal of Virology</i> , 2018, 92, .	3.4	38
10	P1-independent replication and local movement of Rice yellow mottle virus in host and non-host plant species. <i>Virology</i> , 2017, 502, 28-32.	2.4	5
11	Electronic Health Record-Based Genome-Wide Meta-Analysis Provides New Insights on the Genetic Architecture of Non-Alcoholic Fatty Liver Disease. <i>SSRN Electronic Journal</i> , 0, , .	0.4	2
12	Genome-wide association study of susceptibility to hospitalised respiratory infections. <i>Wellcome Open Research</i> , 0, 6, 290.	1.8	3