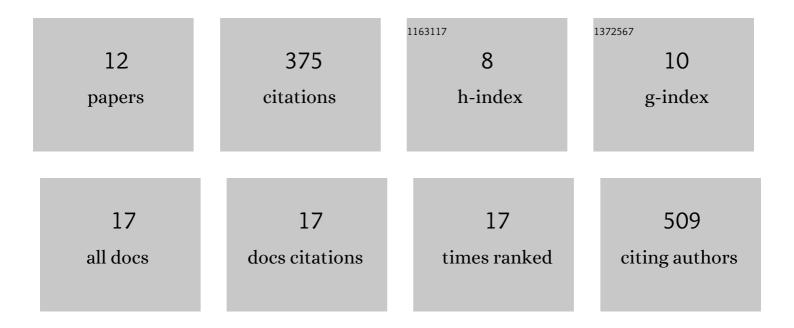
Erik Abner

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5520345/publications.pdf Version: 2024-02-01



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