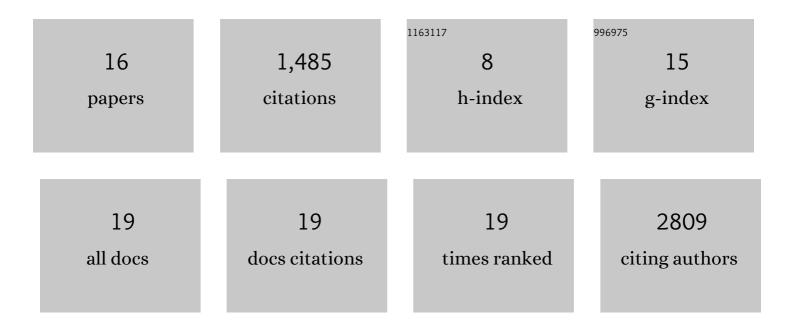
## A P Jason De Koning

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/215410/publications.pdf Version: 2024-02-01



A P JASON DE KONING

#	Article	IF	CITATIONS
1	Transcriptome analysis from muscle biopsy tissues in late-onset myopathies identifies potential biomarkers correlating to muscle pathology. Neuromuscular Disorders, 2022, , .	0.6	О
2	Associations Among Parental Caregiving Quality, Cannabinoid Receptor 1 Expression-Based Polygenic Scores, and Infant-Parent Attachment: Evidence for Differential Genetic Susceptibility?. Frontiers in Neuroscience, 2021, 15, 704392.	2.8	3
3	Parenting Interacts With Plasticity Genes in Predicting Behavioral Outcomes in Preschoolers. Canadian Journal of Nursing Research, 2020, 52, 290-307.	1.5	6
4	Long noncoding RNAs associated with phenotypic severity in multiple sclerosis. Multiple Sclerosis and Related Disorders, 2019, 36, 101407.	2.0	14
5	Molecular Adaptations for Sensing and Securing Prey and Insight into Amniote Genome Diversity from the Garter Snake Genome. Genome Biology and Evolution, 2018, 10, 2110-2129.	2.5	72
6	Wright–Fisher exact solver (WFES): scalable analysis of population genetic models without simulation or diffusion theory. Bioinformatics, 2017, 33, 1416-1417.	4.1	9
7	EzMap: a simple pipeline for reproducible analysis of the human virome. Bioinformatics, 2017, 33, 2573-2574.	4.1	8
8	Allele Age Under Non-Classical Assumptions is Clarified by an Exact Computational Markov Chain Approach. Scientific Reports, 2017, 7, 11869.	3.3	5
9	The Burmese python genome reveals the molecular basis for extreme adaptation in snakes. Proceedings of the United States of America, 2013, 110, 20645-20650.	7.1	260
10	Phylogenetics, likelihood, evolution and complexity. Bioinformatics, 2012, 28, 2989-2990.	4.1	10
11	Sequencing the genome of the Burmese python (Python molurus bivittatus) as a model for studying extreme adaptations in snakes. Genome Biology, 2011, 12, 406.	9.6	58
12	Discovery of Highly Divergent Repeat Landscapes in Snake Genomes Using High-Throughput Sequencing. Genome Biology and Evolution, 2011, 3, 641-653.	2.5	87
13	A multi-organ transcriptome resource for the Burmese Python (Python molurus bivittatus). BMC Research Notes, 2011, 4, 310.	1.4	18
14	Repetitive Elements May Comprise Over Two-Thirds of the Human Genome. PLoS Genetics, 2011, 7, e1002384.	3.5	907
15	Dynamic Nucleotide Mutation Gradients and Control Region Usage in Squamate Reptile Mitochondrial Genomes. Cytogenetic and Genome Research, 2009, 127, 112-127.	1.1	22
16	An ancient adaptive episode of convergent molecular evolution confounds phylogenetic inference. Nature Precedings, 2008, , .	0.1	1