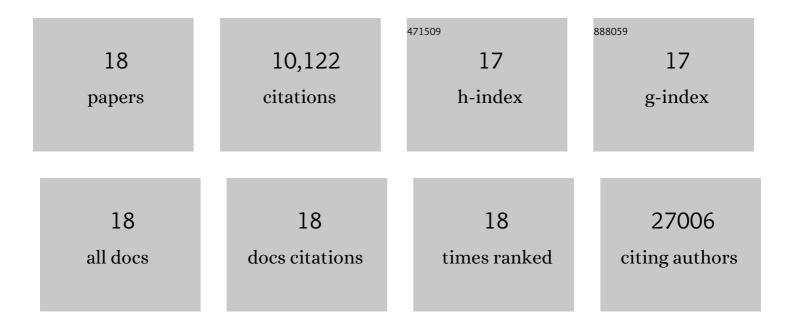
Graham R S Ritchie

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5485860/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Whole-Genome Sequencing Coupled to Imputation Discovers Genetic Signals for Anthropometric Traits. American Journal of Human Genetics, 2017, 100, 865-884.	6.2	131
2	The Ensembl Variant Effect Predictor. Genome Biology, 2016, 17, 122.	8.8	5,181
3	The Ensembl REST API: Ensembl Data for Any Language. Bioinformatics, 2015, 31, 143-145.	4.1	161
4	The African Genome Variation Project shapes medical genetics in Africa. Nature, 2015, 517, 327-332.	27.8	473
5	Functional Annotation of Rare Genetic Variants. , 2015, , 57-70.		2
6	Genetic characterization of Greek population isolates reveals strong genetic drift at missense and trait-associated variants. Nature Communications, 2014, 5, 5345.	12.8	60
7	Functional annotation of noncoding sequence variants. Nature Methods, 2014, 11, 294-296.	19.0	493
8	Computational approaches to identify functional genetic variants in cancer genomes. Nature Methods, 2013, 10, 723-729.	19.0	161
9	Integrative Annotation of Variants from 1092 Humans: Application to Cancer Genomics. Science, 2013, 342, 1235587.	12.6	341
10	A rare functional cardioprotective APOC3 variant has risen in frequency in distinct population isolates. Nature Communications, 2013, 4, 2872.	12.8	77
11	A Combined Functional Annotation Score for Non-Synonymous Variants. Human Heredity, 2012, 73, 47-51.	0.8	90
12	Ensembl 2012. Nucleic Acids Research, 2012, 40, D84-D90.	14.5	840
13	Ensembl 2013. Nucleic Acids Research, 2012, 41, D48-D55.	14.5	856
14	Genome-wide meta-analysis of common variant differences between men and women. Human Molecular Genetics, 2012, 21, 4805-4815.	2.9	33
15	Ensembl 2011. Nucleic Acids Research, 2011, 39, D800-D806.	14.5	630
16	Modernizing Reference Genome Assemblies. PLoS Biology, 2011, 9, e1001091.	5.6	458
17	Signalling signalhood and the emergence of communication. Cognition, 2009, 113, 226-233.	2.2	115
18	Song learning as an indicator mechanism: Modelling the developmental stress hypothesis. Journal of Theoretical Biology, 2008, 251, 570-583.	1.7	20

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