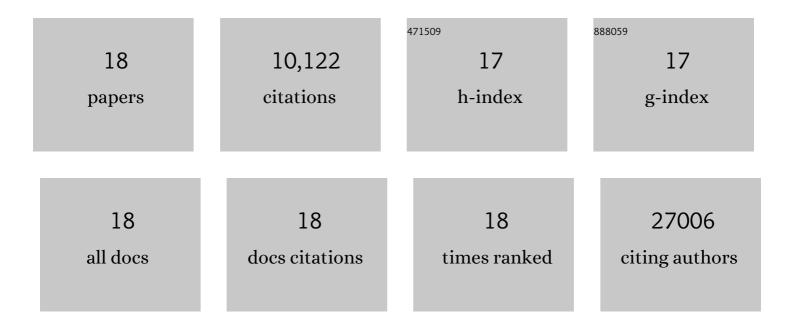
Graham R S Ritchie

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

18 Functional Annotation of Rare Genetic Variants. , 2015, , 57-70.