

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

18
papers

10,122
citations

471509

17
h-index

888059

17
g-index

18
all docs

18
docs citations

18
times ranked

27006
citing authors

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