Julie G Hussin

List of Publications by Year in descending order

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471061 525886 1,215 27 17 27 citations h-index g-index papers 46 46 46 2630 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Re-engineering the zinc fingers of PRDM9 reverses hybrid sterility in mice. Nature, 2016, 530, 171-176.	13.7	194
2	Mutations in Centrosomal Protein CEP152 in Primary Microcephaly Families Linked to MCPH4. American Journal of Human Genetics, 2010, 87, 40-51.	2.6	174
3	Rare Copy Number Variants Contribute to Congenital Left-Sided Heart Disease. PLoS Genetics, 2012, 8, e1002903.	1.5	119
4	A Population Genetic Approach to Mapping Neurological Disorder Genes Using Deep Resequencing. PLoS Genetics, 2011, 7, e1001318.	1.5	89
5	Recombination affects accumulation of damaging and disease-associated mutations in human populations. Nature Genetics, 2015, 47, 400-404.	9.4	84
6	Whole-Exome Sequencing Reveals a Rapid Change in the Frequency of Rare Functional Variants in a Founding Population of Humans. PLoS Genetics, 2013, 9, e1003815.	1.5	70
7	Plasmodium falciparum Genetic Diversity Maintained and Amplified Over 5 Years of a Low Transmission Endemic in the Peruvian Amazon. Molecular Biology and Evolution, 2011, 28, 1973-1986.	3.5	50
8	Age-Dependent Recombination Rates in Human Pedigrees. PLoS Genetics, 2011, 7, e1002251.	1.5	48
9	Rare allelic forms of <i>PRDM9</i> associated with childhood leukemogenesis. Genome Research, 2013, 23, 419-430.	2.4	45
10	SARS–CoV-2 Receptor ACE2 Gene Is Associated with Hypertension and Severity of COVID 19: Interaction with Sex, Obesity, and Smoking. American Journal of Hypertension, 2021, 34, 367-376.	1.0	42
11	Multiscale PHATE identifies multimodal signatures of COVID-19. Nature Biotechnology, 2022, 40, 681-691.	9.4	39
12	The impact of recombination on human mutation load and disease. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160465.	1.8	31
13	Multicohort analysis of the maternal age effect on recombination. Nature Communications, 2015, 6, 7846.	5.8	29
14	Insights into Platypus Population Structure and History from Whole-Genome Sequencing. Molecular Biology and Evolution, 2018, 35, 1238-1252.	3.5	27
15	Polygenic risk scores predict diabetes complications and their response to intensive blood pressure and glucose control. Diabetologia, 2021, 64, 2012-2025.	2.9	24
16	The mutational landscape of SARS-CoV-2 variants diversifies TÂcell targets in an HLA-supertype-dependent manner. Cell Systems, 2022, 13, 143-157.e3.	2.9	22
17	Next-generation sequencing approaches for genetic mapping of complex diseases. Journal of Neuroimmunology, 2012, 248, 10-22.	1.1	19
18	Haplotype allelic classes for detecting ongoing positive selection. BMC Bioinformatics, 2010, 11, 65.	1.2	18

#	Article	IF	CITATIONS
19	A Family-Based Probabilistic Method for Capturing De Novo Mutations from High-Throughput Short-Read Sequencing Data. Statistical Applications in Genetics and Molecular Biology, 2012, 11, .	0.2	16
20	Implementing Machine Learning in Interventional Cardiology: The Benefits Are Worth the Trouble. Frontiers in Cardiovascular Medicine, 2021, 8, 711401.	1.1	12
21	A sex-specific evolutionary interaction between ADCY9 and CETP. ELife, 2021, 10, .	2.8	8
22	Genetics of symptom remission in outpatients with COVID-19. Scientific Reports, 2021, 11, 10847.	1.6	7
23	Population Genomics Approaches for Genetic Characterization of SARS-CoV-2 Lineages. Frontiers in Medicine, 2022, 9, 826746.	1.2	7
24	Long-Chain Acylcarnitines and Monounsaturated Fatty Acids Discriminate Heart Failure Patients According to Pulmonary Hypertension Status. Metabolites, 2021, 11, 196.	1.3	5
25	deepSimDEF: deep neural embeddings of gene products and gene ontology terms for functional analysis of genes. Bioinformatics, 2022, 38, 3051-3061.	1.8	4
26	Differential modulation of polyunsaturated fatty acids in patients with myocardial infarction treated with ticagrelor or clopidogrel. Cell Reports Medicine, 2021, 2, 100299.	3.3	2
27	Multiscale PHATE Exploration of SARS-CoV-2 Data Reveals Multimodal Signatures of Disease. SSRN Electronic Journal, 0, , .	0.4	1