

Elisha D O Roberson

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

4,220
citations

430874

18
h-index

501196

28
g-index

40
all docs

40
docs citations

40
times ranked

7446
citing authors

#	ARTICLE	IF	CITATIONS
1	Frequent Mutation of <i>BAP1</i> in Metastasizing Uveal Melanomas. <i>Science</i> , 2010, 330, 1410-1413.	12.6	1,242
2	Recurrent mutations at codon 625 of the splicing factor SF3B1 in uveal melanoma. <i>Nature Genetics</i> , 2013, 45, 133-135.	21.4	447
3	Early-onset lymphoproliferation and autoimmunity caused by germline STAT3 gain-of-function mutations. <i>Blood</i> , 2015, 125, 591-599.	1.4	436
4	PSORS2 Is Due to Mutations in CARD14. <i>American Journal of Human Genetics</i> , 2012, 90, 784-795.	6.2	365
5	Rare and Common Variants in CARD14, Encoding an Epidermal Regulator of NF-kappaB, in Psoriasis. <i>American Journal of Human Genetics</i> , 2012, 90, 796-808.	6.2	306
6	Retinal Ganglion Cells Downregulate Gene Expression and Lose Their Axons within the Optic Nerve Head in a Mouse Glaucoma Model. <i>Journal of Neuroscience</i> , 2008, 28, 548-561.	3.6	285
7	Mutations in the Gene PRRT2 Cause Paroxysmal Kinesigenic Dyskinesia with Infantile Convulsions. <i>Cell Reports</i> , 2012, 1, 2-12.	6.4	250
8	Psoriasis genetics: breaking the barrier. <i>Trends in Genetics</i> , 2010, 26, 415-423.	6.7	203
9	A Subset of Methylated CpG Sites Differentiate Psoriatic from Normal Skin. <i>Journal of Investigative Dermatology</i> , 2012, 132, 583-592.	0.7	138
10	Inference of Relationships in Population Data Using Identity-by-Descent and Identity-by-State. <i>PLoS Genetics</i> , 2011, 7, e1002287.	3.5	76
11	A mucosal imprint left by prior <i>Escherichia coli</i> bladder infection sensitizes to recurrent disease. <i>Nature Microbiology</i> , 2017, 2, 16196.	13.3	67
12	Rare Variants in the Functional Domains of Complement Factor H Are Associated With Age-Related Macular Degeneration. , 2015, 56, 6873.		60
13	Genomic analysis of partial 21q monosomies with variable phenotypes. <i>European Journal of Human Genetics</i> , 2011, 19, 235-238.	2.8	52
14	Visualization of uniparental inheritance, Mendelian inconsistencies, deletions, and parent of origin effects in single nucleotide polymorphism trio data with SNP trio. <i>Human Mutation</i> , 2007, 28, 1225-1235.	2.5	46
15	Dynamic immunoglobulin responses to gut bacteria during inflammatory bowel disease. <i>Gut Microbes</i> , 2020, 11, 405-420.	9.8	44
16	Protective Low-Frequency Variants for Preeclampsia in the Fms Related Tyrosine Kinase 1 Gene in the Finnish Population. <i>Hypertension</i> , 2017, 70, 365-371.	2.7	37
17	Visualization of Shared Genomic Regions and Meiotic Recombination in High-Density SNP Data. <i>PLoS ONE</i> , 2009, 4, e6711.	2.5	36
18	<i>Nicastrin</i> haploinsufficiency alters expression of type I interferon-stimulated genes: the relationship to familial hidradenitis suppurativa. <i>Clinical and Experimental Dermatology</i> , 2019, 44, e118-e125.	1.3	28

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19	Mucosal infection rewires TNF ϵ signaling dynamics to skew susceptibility to recurrence. <i>ELife</i> , 2019, 8, .	6.0	24
20	STING Gain-of-Function Disrupts Lymph Node Organogenesis and Innate Lymphoid Cell Development in Mice. <i>Cell Reports</i> , 2020, 31, 107771.	6.4	18
21	Transcriptomes of peripheral blood mononuclear cells from juvenile dermatomyositis patients show elevated inflammation even when clinically inactive. <i>Scientific Reports</i> , 2022, 12, 275.	3.3	12
22	C5b-9 and MASP2 deposition in skin and bone marrow microvasculature characterize hematopoietic stem cell transplant-associated thrombotic microangiopathy. <i>Bone Marrow Transplantation</i> , 2022, 57, 1445-1447.	2.4	9
23	Consanguinity in Centre d'Étude du Polymorphisme Humain (CEPH) pedigrees. <i>European Journal of Human Genetics</i> , 2012, 20, 657-667.	2.8	8
24	Motif scraper: a cross-platform, open-source tool for identifying degenerate nucleotide motif matches in FASTA files. <i>Bioinformatics</i> , 2018, 34, 3926-3928.	4.1	7
25	Locations and patterns of meiotic recombination in two-generation pedigrees. <i>BMC Medical Genetics</i> , 2009, 10, 93.	2.1	6
26	Editorial: Genomic Advances in Systemic Sclerosis: It Is Time for Precision. <i>Arthritis and Rheumatology</i> , 2015, 67, 2801-2805.	5.6	5
27	A catalog of CasX genome editing sites in common model organisms. <i>BMC Genomics</i> , 2019, 20, 528.	2.8	4
28	Alterations of the Primary Cilia Gene <i>SPAG17</i> and <i>SOX9</i> Locus Noncoding RNAs Identified by RNA-Seq Analysis in Patients With Systemic Sclerosis. <i>Arthritis and Rheumatology</i> , 2023, 75, 108-119.	5.6	4
29	Identification of high-efficiency 3'GG gRNA motifs in indexed FASTA files with ngg2. <i>PeerJ Computer Science</i> , 2015, 1, e33.	4.5	3
30	Validation of a commercial antibody to detect endogenous human nicastrin by immunoblot. <i>F1000Research</i> , 2019, 8, 1211.	1.6	0
31	Validation of a commercial antibody to detect endogenous human nicastrin by immunoblot. <i>F1000Research</i> , 2019, 8, 1211.	1.6	0