

Nicholas B Blackburn

List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

436
citations

759190

12
h-index

794568

19
g-index

27
all docs

27
docs citations

27
times ranked

1444
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of Copy Number Variation of the 15q11.2 BP1-BP2 Region With Cortical and Subcortical Morphology and Cognition. <i>JAMA Psychiatry</i> , 2020, 77, 420.	11.0	54
2	Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. <i>Molecular Psychiatry</i> , 2020, 25, 584-602.	7.9	49
3	Rediscovering the value of families for psychiatric genetics research. <i>Molecular Psychiatry</i> , 2019, 24, 523-535.	7.9	43
4	RNA-seq profiling of a radiation resistant and radiation sensitive prostate cancer cell line highlights opposing regulation of DNA repair and targets for radiosensitization. <i>BMC Cancer</i> , 2014, 14, 808.	2.6	35
5	Association of CREBRF variants with obesity and diabetes in Pacific Islanders from Guam and Saipan. <i>Diabetologia</i> , 2019, 62, 1647-1652.	6.3	26
6	Abdominal Obesity and Brain Atrophy in Type 2 Diabetes Mellitus. <i>PLoS ONE</i> , 2015, 10, e0142589.	2.5	25
7	1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. <i>Translational Psychiatry</i> , 2021, 11, 182.	4.8	24
8	Rare, potentially pathogenic variants in 21 keratoconus candidate genes are not enriched in cases in a large Australian cohort of European descent. <i>PLoS ONE</i> , 2018, 13, e0199178.	2.5	21
9	Molecular characterization of a marine turtle tumor epizootic, profiling external, internal and postsurgical regrowth tumors. <i>Communications Biology</i> , 2021, 4, 152.	4.4	20
10	Comparative Analysis of Immune Checkpoint Molecules and Their Potential Role in the Transmissible Tasmanian Devil Facial Tumor Disease. <i>Frontiers in Immunology</i> , 2017, 8, 513.	4.8	19
11	Common genetic variation within miR-146a predicts disease onset and relapse in multiple sclerosis. <i>Neurological Sciences</i> , 2018, 39, 297-304.	1.9	19
12	Rare DEGS1 variant significantly alters de novo ceramide synthesis pathway. <i>Journal of Lipid Research</i> , 2019, 60, 1630-1639.	4.2	16
13	Rare, Potentially Pathogenic Variants in <i>ZNF469</i> Are Not Enriched in Keratoconus in a Large Australian Cohort of European Descent. , 2017, 58, 6248.		13
14	Evaluating a CLL susceptibility variant in ITGB2 in families with multiple subtypes of hematological malignancies. <i>Blood</i> , 2017, 130, 86-88.	1.4	11
15	A retrospective examination of mean relative telomere length in the Tasmanian Familial Hematological Malignancies Study. <i>Oncology Reports</i> , 2015, 33, 25-32.	2.6	9
16	Impact of the G84E variant on HOXB13 gene and protein expression in formalin-fixed, paraffin-embedded prostate tumours. <i>Scientific Reports</i> , 2017, 7, 17778.	3.3	8
17	Genome-wide linkage scan for loci influencing plasma triglyceride levels. <i>BMC Proceedings</i> , 2018, 12, 52.	1.6	7
18	Reliability of genomic predictions of complex human phenotypes. <i>BMC Proceedings</i> , 2018, 12, 51.	1.6	7

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19	Glycated Serum Protein Genetics and Pleiotropy with Cardiometabolic Risk Factors. <i>Journal of Diabetes Research</i> , 2019, 2019, 1-9.	2.3	6
20	Heritability and genetic associations of triglyceride and HDL-C levels using pedigree-based and empirical kinships. <i>BMC Proceedings</i> , 2018, 12, 34.	1.6	5
21	Mesenchymal plasticity of devil facial tumour cells during in vivo vaccine and immunotherapy trials. <i>Immunology and Cell Biology</i> , 2021, 99, 711-723.	2.3	5
22	Identifying the Lipidomic Effects of a Rare Loss-of-Function Deletion in <i>ANGPTL3</i> . <i>Circulation Genomic and Precision Medicine</i> , 2021, 14, e003232.	3.6	3
23	Genotype phasing in pedigrees using whole-genome sequence data. <i>European Journal of Human Genetics</i> , 2020, 28, 790-803.	2.8	3
24	Massively parallel sequencing in hereditary prostate cancer families reveals a rare risk variant in the DNA repair gene, <i>RAD51C</i> . <i>European Journal of Cancer</i> , 2021, 159, 52-55.	2.8	3