

Nicholas B Blackburn

List of Publications by Citations

Source: <https://exaly.com/author-pdf/7072862/nicholas-b-blackburn-publications-by-citations.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

25
papers

272
citations

11
h-index

15
g-index

27
ext. papers

377
ext. citations

6
avg, IF

2.71
L-index

#	Paper	IF	Citations
25	Rediscovering the value of families for psychiatric genetics research. <i>Molecular Psychiatry</i> , 2019 , 24, 523-535	5.1	30
24	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	4.8	25
23	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, 420-430	14.5	24
22	Dose response of the 16p11.2 distal copy number variant on intracranial volume and basal ganglia. <i>Molecular Psychiatry</i> , 2020 , 25, 584-602	15.1	24
21	Abdominal Obesity and Brain Atrophy in Type 2 Diabetes Mellitus. <i>PLoS ONE</i> , 2015 , 10, e0142589	3.7	23
20	Association of CREBRF variants with obesity and diabetes in Pacific Islanders from Guam and Saipan. <i>Diabetologia</i> , 2019 , 62, 1647-1652	10.3	15
19	Common genetic variation within miR-146a predicts disease onset and relapse in multiple sclerosis. <i>Neurological Sciences</i> , 2018 , 39, 297-304	3.5	15
18	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	3.7	14
17	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	8.4	14
16	Rare, Potentially Pathogenic Variants in ZNF469 Are Not Enriched in Keratoconus in a Large Australian Cohort of European Descent 2017 , 58, 6248-6256		12
15	Molecular characterization of a marine turtle tumor epizootic, profiling external, internal and postsurgical regrowth tumors. <i>Communications Biology</i> , 2021 , 4, 152	6.7	12
14	Evaluating a CLL susceptibility variant in in families with multiple subtypes of hematological malignancies. <i>Blood</i> , 2017 , 130, 86-88	2.2	10
13	Rare variant significantly alters de novo ceramide synthesis pathway. <i>Journal of Lipid Research</i> , 2019 , 60, 1630-1639	6.3	7
12	A retrospective examination of mean relative telomere length in the Tasmanian Familial Hematological Malignancies Study. <i>Oncology Reports</i> , 2015 , 33, 25-32	3.5	7
11	Genome-wide linkage scan for loci influencing plasma triglyceride levels. <i>BMC Proceedings</i> , 2018 , 12, 52	2.3	7
10	1q21.1 distal copy number variants are associated with cerebral and cognitive alterations in humans. <i>Translational Psychiatry</i> , 2021 , 11, 182	8.6	6
9	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	4.9	5

8	Reliability of genomic predictions of complex human phenotypes. <i>BMC Proceedings</i> , 2018 , 12, 51	2.3	5
7	Heritability and genetic associations of triglyceride and HDL-C levels using pedigree-based and empirical kinships. <i>BMC Proceedings</i> , 2018 , 12, 34	2.3	4
6	Glycated Serum Protein Genetics and Pleiotropy with Cardiometabolic Risk Factors. <i>Journal of Diabetes Research</i> , 2019 , 2019, 2310235	3.9	3
5	Transcriptomic profiling of fibropapillomatosis in green sea turtles (<i>Chelonia mydas</i>) from South Texas		3
4	Mesenchymal plasticity of devil facial tumour cells during in vivo vaccine and immunotherapy trials. <i>Immunology and Cell Biology</i> , 2021 , 99, 711-723	5	3
3	Genotype phasing in pedigrees using whole-genome sequence data. <i>European Journal of Human Genetics</i> , 2020 , 28, 790-803	5.3	2
2	Identifying the Lipidomic Effects of a Rare Loss-of-Function Deletion in. <i>Circulation Genomic and Precision Medicine</i> , 2021 , 14, e003232	5.2	1
1	Massively parallel sequencing in hereditary prostate cancer families reveals a rare risk variant in the DNA repair gene, RAD51C. <i>European Journal of Cancer</i> , 2021 , 159, 52-55	7.5	0