

Denis A Baird

List of Publications by Year in Descending Order

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

17
papers

1,699
citations

10
h-index

23
g-index

23
ext. papers

3,297
ext. citations

11.7
avg, IF

4.19
L-index

#	Paper	IF	Citations
17	Genome-wide study of DNA methylation shows alterations in metabolic, inflammatory, and cholesterol pathways in ALS.. <i>Science Translational Medicine</i> , 2022 , 14, eabj0264	17.5	4
16	Common and rare variant association analyses in amyotrophic lateral sclerosis identify 15 risk loci with distinct genetic architectures and neuron-specific biology. <i>Nature Genetics</i> , 2021 , 53, 1636-1648	36.3	19
15	Identifying drug targets for neurological and psychiatric disease via genetics and the brain transcriptome. <i>PLoS Genetics</i> , 2021 , 17, e1009224	6	10
14	Genome-wide association study implicates novel loci and reveals candidate effector genes for longitudinal pediatric bone accrual. <i>Genome Biology</i> , 2021 , 22, 1	18.3	58
13	The influence of adult hip shape genetic variants on adolescent hip shape: Findings from a population-based DXA study. <i>Bone</i> , 2021 , 143, 115792	4.7	3
12	Phenome-wide Mendelian randomization mapping the influence of the plasma proteome on complex diseases. <i>Nature Genetics</i> , 2020 , 52, 1122-1131	36.3	75
11	Mendelian Randomization Analysis Reveals a Causal Influence of Circulating Sclerostin Levels on Bone Mineral Density and Fractures. <i>Journal of Bone and Mineral Research</i> , 2019 , 34, 1824-1836	6.3	11
10	Identification of Novel Loci Associated With Hip Shape: A Meta-Analysis of Genomewide Association Studies. <i>Journal of Bone and Mineral Research</i> , 2019 , 34, 241-251	6.3	32
9	Using Y-Chromosomal Haplogroups in Genetic Association Studies and Suggested Implications. <i>Genes</i> , 2018 , 9,	4.2	4
8	Investigation of the Relationship Between Susceptibility Loci for Hip Osteoarthritis and Dual X-Ray Absorptiometry-Derived Hip Shape in a Population-Based Cohort of Perimenopausal Women. <i>Arthritis and Rheumatology</i> , 2018 , 70, 1984-1993	9.5	19
7	Author response: The MR-Base platform supports systematic causal inference across the human phenome 2018 ,		17
6	The MR-Base platform supports systematic causal inference across the human phenome. <i>ELife</i> , 2018 , 7,	8.9	1190
5	Recent Developments in Mendelian Randomization Studies. <i>Current Epidemiology Reports</i> , 2017 , 4, 330-345	3.45	218
4	Identifying Highly Penetrant Disease Causal Mutations Using Next Generation Sequencing: Guide to Whole Process. <i>BioMed Research International</i> , 2015 , 2015, 923491	3	6
3	Multi-omics study revealing putative drug targets of COVID-19 severity and other viral infection diseases		7
2	Phenome-wide Mendelian randomization mapping the influence of the plasma proteome on complex diseases		21
1	Brain expression quantitative trait locus and network analysis reveals downstream effects and putative drivers for brain-related diseases		4

