

# Denis A Baird

## List of Publications by Citations

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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	The MR-Base platform supports systematic causal inference across the human phenome. <i>ELife</i> , <b>2018</b> , 7,	8.9	1190
16	Recent Developments in Mendelian Randomization Studies. <i>Current Epidemiology Reports</i> , <b>2017</b> , 4, 330-345	34.5	218
15	Phenome-wide Mendelian randomization mapping the influence of the plasma proteome on complex diseases. <i>Nature Genetics</i> , <b>2020</b> , 52, 1122-1131	36.3	75
14	Genome-wide association study implicates novel loci and reveals candidate effector genes for longitudinal pediatric bone accrual. <i>Genome Biology</i> , <b>2021</b> , 22, 1	18.3	58
13	Identification of Novel Loci Associated With Hip Shape: A Meta-Analysis of Genomewide Association Studies. <i>Journal of Bone and Mineral Research</i> , <b>2019</b> , 34, 241-251	6.3	32
12	Phenome-wide Mendelian randomization mapping the influence of the plasma proteome on complex diseases		21
11	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> , <b>2018</b> , 70, 1984-1993	9.5	19
10	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> , <b>2021</b> , 53, 1636-1648	36.3	19
9	Author response: The MR-Base platform supports systematic causal inference across the human phenome <b>2018</b> ,		17
8	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> , <b>2019</b> , 34, 1824-1836	6.3	11
7	Identifying drug targets for neurological and psychiatric disease via genetics and the brain transcriptome. <i>PLoS Genetics</i> , <b>2021</b> , 17, e1009224	6	10
6	Multi-omics study revealing putative drug targets of COVID-19 severity and other viral infection diseases		7
5	Identifying Highly Penetrant Disease Causal Mutations Using Next Generation Sequencing: Guide to Whole Process. <i>BioMed Research International</i> , <b>2015</b> , 2015, 923491	3	6
4	Using Y-Chromosomal Haplogroups in Genetic Association Studies and Suggested Implications. <i>Genes</i> , <b>2018</b> , 9,	4.2	4
3	Brain expression quantitative trait locus and network analysis reveals downstream effects and putative drivers for brain-related diseases		4
2	Genome-wide study of DNA methylation shows alterations in metabolic, inflammatory, and cholesterol pathways in ALS.. <i>Science Translational Medicine</i> , <b>2022</b> , 14, eabj0264	17.5	4
1	The influence of adult hip shape genetic variants on adolescent hip shape: Findings from a population-based DXA study. <i>Bone</i> , <b>2021</b> , 143, 115792	4.7	3

