

# Andrew Bjonnes

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10580580/publications.pdf>

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14  
papers

3,352  
citations

687363

13  
h-index

1058476

14  
g-index

14  
all docs

14  
docs citations

14  
times ranked

8301  
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. <i>Journal of the American College of Cardiology</i> , 2019, 73, 58-66.	2.8	147
2	Transethnic Evaluation Identifies Low-Frequency Loci Associated With 25-Hydroxyvitamin D Concentrations. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2018, 103, 1380-1392.	3.6	33
3	An integrated clinical program and crowdsourcing strategy for genomic sequencing and Mendelian disease gene discovery. <i>Npj Genomic Medicine</i> , 2018, 3, 21.	3.8	24
4	Genetic predisposition to elevated levels of C-reactive protein is associated with a decreased risk for preeclampsia. <i>Hypertension in Pregnancy</i> , 2017, 36, 30-35.	1.1	4
5	No Association of Coronary Artery Disease with X-Chromosomal Variants in Comprehensive International Meta-Analysis. <i>Scientific Reports</i> , 2016, 6, 35278.	3.3	25
6	Genome-wide association study of 25(OH) Vitamin D concentrations in Punjabi Sikhs: Results of the Asian Indian diabetic heart study. <i>Journal of Steroid Biochemistry and Molecular Biology</i> , 2016, 158, 149-156.	2.5	43
7	Genetic Risk Score for Essential Hypertension and Risk of Preeclampsia. <i>American Journal of Hypertension</i> , 2016, 29, 17-24.	2.0	19
8	Genetic Predisposition to Dyslipidemia and Risk of Preeclampsia. <i>American Journal of Hypertension</i> , 2015, 28, 915-923.	2.0	19
9	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , 2015, 523, 459-462.	27.8	173
10	Causal mechanisms and balancing selection inferred from genetic associations with polycystic ovary syndrome. <i>Nature Communications</i> , 2015, 6, 8464.	12.8	304
11	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. <i>Nature Genetics</i> , 2015, 47, 1282-1293.	21.4	294
12	A comprehensive 1000 Genomes-based genome-wide association meta-analysis of coronary artery disease. <i>Nature Genetics</i> , 2015, 47, 1121-1130.	21.4	2,054
13	Genome-Wide Association Study Identifies Variants in Casein Kinase II ( <i>CSNK2A2</i> ) to be Associated With Leukocyte Telomere Length in a Punjabi Sikh Diabetic Cohort. <i>Circulation: Cardiovascular Genetics</i> , 2014, 7, 287-295.	5.1	46
14	Genome-Wide Association Study Identifies a Novel Locus Contributing to Type 2 Diabetes Susceptibility in Sikhs of Punjabi Origin From India. <i>Diabetes</i> , 2013, 62, 1746-1755.	0.6	167