

# John Connell

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12141574/publications.pdf>

Version: 2024-02-01

17  
papers

4,501  
citations

516710

16  
h-index

839539

18  
g-index

20  
all docs

20  
docs citations

20  
times ranked

8679  
citing authors

#	ARTICLE	IF	CITATIONS
1	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. <i>Nature Genetics</i> , 2007, 39, 1329-1337.	21.4	1,298
2	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. <i>Nature Genetics</i> , 2018, 50, 1412-1425.	21.4	924
3	Meta-Analysis of 28,141 Individuals Identifies Common Variants within Five New Loci That Influence Uric Acid Concentrations. <i>PLoS Genetics</i> , 2009, 5, e1000504.	3.5	572
4	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. <i>Nature Genetics</i> , 2017, 49, 403-415.	21.4	492
5	Genome-wide mapping of human loci for essential hypertension. <i>Lancet, The</i> , 2003, 361, 2118-2123.	13.7	247
6	Genome-wide scan identifies CDH13 as a novel susceptibility locus contributing to blood pressure determination in two European populations. <i>Human Molecular Genetics</i> , 2009, 18, 2288-2296.	2.9	170
7	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. <i>Hypertension</i> , 2017, 70, .	2.7	123
8	Association of the thyroid stimulating hormone receptor gene (TSHR) with Graves' disease. <i>Human Molecular Genetics</i> , 2009, 18, 1704-1713.	2.9	122
9	Meta-analysis of Dense Genecentric Association Studies Reveals Common and Uncommon Variants Associated with Height. <i>American Journal of Human Genetics</i> , 2011, 88, 6-18.	6.2	122
10	Targeting 160 Candidate Genes for Blood Pressure Regulation with a Genome-Wide Genotyping Array. <i>PLoS ONE</i> , 2009, 4, e6034.	2.5	98
11	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. <i>Nature Genetics</i> , 2020, 52, 1314-1332.	21.4	91
12	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. <i>Nature Communications</i> , 2019, 10, 376.	12.8	64
13	Genetic Variation at the Locus Encompassing 11- $\beta$ Hydroxylase and Aldosterone Synthase Accounts for Heritability in Cortisol Precursor (11-Deoxycortisol) Urinary Metabolite Excretion. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2005, 90, 1072-1077.	3.6	52
14	Two-dimensional genome-scan identifies novel epistatic loci for essential hypertension. <i>Human Molecular Genetics</i> , 2006, 15, 1365-1374.	2.9	50
15	Increased Support for Linkage of a Novel Locus on Chromosome 5q13 for Essential Hypertension in the British Genetics of Hypertension Study. <i>Hypertension</i> , 2006, 48, 105-111.	2.7	22
16	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. <i>Molecular Psychiatry</i> , 2020, 26, 2111-2125.	7.9	17
17	ACTH and Polymorphisms at Steroidogenic Loci as Determinants of Aldosterone Secretion and Blood Pressure. <i>International Journal of Molecular Sciences</i> , 2017, 18, 579.	4.1	5