John Connell

List of Publications by Year
in descending order

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	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. Nature Genetics, 2020, 52, 1314-1332.	21.4	91
2	Gene-educational attainment interactions in a multi-ancestry genome-wide meta-analysis identify novel blood pressure loci. Molecular Psychiatry, 2020, 26, 2111-2125.	7.9	17
3	Multi-ancestry study of blood lipid levels identifies four loci interacting with physical activity. Nature Communications, 2019, 10, 376.	12.8	64
4	Genetic analysis of over 1 million people identifies 535 new loci associated with blood pressure traits. Nature Genetics, 2018, 50, 1412-1425.	21.4	924
5	Genome-wide association analysis identifies novel blood pressure loci and offers biological insights into cardiovascular risk. Nature Genetics, 2017, 49, 403-415.	21.4	492
6	Novel Blood Pressure Locus and Gene Discovery Using Genome-Wide Association Study and Expression Data Sets From Blood and the Kidney. Hypertension, 2017, 70, .	2.7	123
7	ACTH and Polymorphisms at Steroidogenic Loci as Determinants of Aldosterone Secretion and Blood Pressure. International Journal of Molecular Sciences, 2017, 18, 579.	4.1	5
8	Meta-analysis of Dense Genecentric Association Studies Reveals Common and Uncommon Variants Associated with Height. American Journal of Human Genetics, 2011, 88, 6-18.	6.2	122
9	Targeting 160 Candidate Genes for Blood Pressure Regulation with a Genome-Wide Genotyping Array. PLoS ONE, 2009, 4, e6034.	2.5	98
10	Association of the thyroid stimulating hormone receptor gene (TSHR) with Graves' disease. Human Molecular Genetics, 2009, 18, 1704-1713.	2.9	122
11	Meta-Analysis of 28,141 Individuals Identifies Common Variants within Five New Loci That Influence Uric Acid Concentrations. PLoS Genetics, 2009, 5, e1000504.	3.5	572
12	Genome-wide scan identifies CDH13 as a novel susceptibility locus contributing to blood pressure determination in two European populations. Human Molecular Genetics, 2009, 18, 2288-2296.	2.9	170
13	Association scan of 14,500 nonsynonymous SNPs in four diseases identifies autoimmunity variants. Nature Genetics, 2007, 39, 1329-1337.	21.4	1,298
14	Two-dimensional genome-scan identifies novel epistatic loci for essential hypertension. Human Molecular Genetics, 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. Hypertension, 2006, 48, 105-111.	2.7	22
16	Genetic Variation at the Locus Encompassing 11-Î² Hydroxylase and Aldosterone Synthase Accounts for Heritability in Cortisol Precursor (11-Deoxycortisol) Urinary Metabolite Excretion. Journal of Clinical Endocrinology and Metabolism, 2005, 90, 1072-1077.	3.6	52
17	Genome-wide mapping of human loci for essential hypertension. Lancet, The, 2003, 361, 2118-2123.	13.7	247