

Vasyl Pihur

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

3,853
citations

623734

14
h-index

940533

16
g-index

17
all docs

17
docs citations

17
times ranked

9058
citing authors

#	ARTICLE	IF	CITATIONS
1	Causal Effect of Plasminogen Activator Inhibitor Type 1 on Coronary Heart Disease. Journal of the American Heart Association, 2017, 6, .	3.7	89
2	The genetics of blood pressure regulation and its target organs from association studies in 342,415 individuals. Nature Genetics, 2016, 48, 1171-1184.	21.4	362
3	Loss of β -catenin function in severe autism. Nature, 2015, 520, 51-56.	27.8	145
4	Effects of Long-Term Averaging of Quantitative Blood Pressure Traits on the Detection of Genetic Associations. American Journal of Human Genetics, 2014, 95, 49-65.	6.2	73
5	An Enhancer Polymorphism at the Cardiomyocyte Intercalated Disc Protein NOS1AP Locus Is a Major Regulator of the QT Interval. American Journal of Human Genetics, 2014, 94, 854-869.	6.2	72
6	MmPalateMiRNA, an R package compendium illustrating analysis of miRNA microarray data. Source Code for Biology and Medicine, 2013, 8, 1.	1.7	36
7	Effects of Rare and Common Blood Pressure Gene Variants on Essential Hypertension. Circulation Research, 2013, 112, 318-326.	4.5	24
8	Genetic variants in novel pathways influence blood pressure and cardiovascular disease risk. Nature, 2011, 478, 103-109.	27.8	1,855
9	TGF β -1 and Wnt-3a interact to induce unique gene expression profiles in murine embryonic palate mesenchymal cells. Reproductive Toxicology, 2011, 31, 128-133.	2.9	1
10	Association of genetic variation with systolic and diastolic blood pressure among African Americans: the Candidate Gene Association Resource study. Human Molecular Genetics, 2011, 20, 2273-2284.	2.9	168
11	Genome-wide association study identifies six new loci influencing pulse pressure and mean arterial pressure. Nature Genetics, 2011, 43, 1005-1011.	21.4	403
12	Meta analysis of Chronic Fatigue Syndrome through integration of clinical, gene expression, SNP and proteomic data. Bioinformatics, 2011, 6, 120-124.	0.5	8
13	Developmental microRNA expression profiling of murine embryonic orofacial tissue. Birth Defects Research Part A: Clinical and Molecular Teratology, 2010, 88, 511-534.	1.6	44
14	Reconstruction of genetic association networks from microarray data: a partial least squares approach. Bioinformatics, 2008, 24, 561-568.	4.1	60
15	clValid : An R Package for Cluster Validation. Journal of Statistical Software, 2008, 25, .	3.7	472
16	Epidemiologic Associations of Carpal Tunnel Syndrome and Sleep Position: Is there a Case for Causation?. Hand, 2007, 2, 127-134.	1.2	41
17	Combined Effects of Variant N-acetyltransferase 1 (NAT1) and Multiple Drug Resistance 1 (MDR1) Genes in Prostate Cancer Risk. FASEB Journal, 2007, 21, A416.	0.5	0