

Estimating the proportion of variation in susceptibility common SNPs

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Runs of Homozygosity Implicate Autozygosity as a Schizophrenia Risk Factor. <i>PLoS Genetics</i> , 2012, 8, e1002656.	1.5	109
2	Estimating the proportion of variation in susceptibility to multiple sclerosis captured by common SNPs. <i>Scientific Reports</i> , 2012, 2, 770.	1.6	16
3	Prenatal nutrition, epigenetics and schizophrenia risk: can we test causal effects?. <i>Epigenomics</i> , 2012, 4, 303-315.	1.0	76
5	A commentary on the gender-specific association of TSNAX/DISC1 locus for schizophrenia and bipolar affective disorder in South Indian population. <i>Journal of Human Genetics</i> , 2012, 57, 475-476.	1.1	0
6	Estimation of pleiotropy between complex diseases using single-nucleotide polymorphism-derived genomic relationships and restricted maximum likelihood. <i>Bioinformatics</i> , 2012, 28, 2540-2542.	1.8	564
7	The gene in its natural habitat: The importance of gene-trait interactions. <i>Development and Psychopathology</i> , 2012, 24, 1307-1318.	1.4	26
8	Genome-wide approaches in pharmacogenomics: heritability estimation and pharmacoethnicity as primary challenges. <i>Pharmacogenomics</i> , 2012, 13, 1101-1104.	0.6	11
9	The genetic architecture of economic and political preferences. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 8026-8031.	3.3	225
10	Implications of Genetic Findings for Understanding Schizophrenia. <i>Schizophrenia Bulletin</i> , 2012, 38, 904-907.	2.3	48
11	A mixed-model approach for genome-wide association studies of correlated traits in structured populations. <i>Nature Genetics</i> , 2012, 44, 1066-1071.	9.4	380
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14	Improved Heritability Estimation from Genome-wide SNPs. <i>American Journal of Human Genetics</i> , 2012, 91, 1011-1021.	2.6	656
15	Target practice: HDAC inhibitors for schizophrenia. <i>Nature Neuroscience</i> , 2012, 15, 1180-1181.	7.1	15
16	Puzzling over schizophrenia: Schizophrenia as a pathway disease. <i>Nature Medicine</i> , 2012, 18, 210-211.	15.2	80
17	Confluence of genes, environment, development, and behavior in a post Genome-Wide Association Study world. <i>Development and Psychopathology</i> , 2012, 24, 1195-1214.	1.4	43
18	Common genetic variants, acting additively, are a major source of risk for autism. <i>Molecular Autism</i> , 2012, 3, 9.	2.6	357
19	Using summary data from the Danish National Registers to estimate heritabilities for schizophrenia, bipolar disorder, and major depressive disorder. <i>Frontiers in Genetics</i> , 2012, 3, 118.	1.1	176

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21	Genetic architectures of psychiatric disorders: the emerging picture and its implications. <i>Nature Reviews Genetics</i> , 2012, 13, 537-551.	7.7	1,025
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31	Genetics of psychiatric disorders in the GWAS era: an update on schizophrenia. <i>European Archives of Psychiatry and Clinical Neuroscience</i> , 2013, 263, 147-154.	1.8	49
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