Richard Sherva

List of Publications by Year
in descending order

56 papers	3,966 citations	159525 30 h-index	182361 51 g-index
58 all docs	58 docs citations	58 times ranked	7422 citing authors

#	Article	IF	Citations
1	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. Nature Neuroscience, 2018, 21, 1656-1669.	7.1	490
2	Common variants in WFS1 confer risk of type 2 diabetes. Nature Genetics, 2007, 39, 951-953.	9.4	333
3	Multiple Independent Loci at Chromosome 15q25.1 Affect Smoking Quantity: a Meta-Analysis and Comparison with Lung Cancer and COPD. PLoS Genetics, 2010, 6, e1001053.	1.5	332
4	A large-scale genome-wide association study meta-analysis of cannabis use disorder. Lancet Psychiatry,the, 2020, 7, 1032-1045.	3.7	200
5	Genome-Wide Association Study of Opioid Dependence: Multiple Associations Mapped to Calcium and Potassium Pathways. Biological Psychiatry, 2014, 76, 66-74.	0.7	192
6	Variants in the CD36 gene associate with the metabolic syndrome and high-density lipoprotein cholesterol. Human Molecular Genetics, 2008, 17, 1695-1704.	1.4	164
7	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. JAMA Psychiatry, 2016, 73, 472.	6.0	148
8	Genomeâ€wide association study of the rate of cognitive decline in Alzheimer's disease. Alzheimer's and Dementia, 2014, 10, 45-52.	0.4	147
9	Fetal hemoglobin in sickle cell anemia: genome-wide association studies suggest a regulatory region in the 5â€² olfactory receptor gene cluster. Blood, 2010, 115, 1815-1822.	0.6	146
10	Association of Granulomatosis With Polyangiitis (Wegener's) With HLAâ€“DPB1*04 and SEMA6A Gene Variants: Evidence From Genomeâ€Wide Analysis. Arthritis and Rheumatism, 2013, 65, 2457-2468.	6.7	138
11	Association of a single nucleotide polymorphism in neuronal acetylcholine receptor subunit alpha 5 (CHRNA5) with smoking status and with â€˜pleasurable buzzâ€™ during early experimentation with smoking. Addiction, 2008, 103, 1544-1552.	1.7	129
12	Common CD36 SNPs reduce protein expression and may contribute to a protective atherogenic profile. Human Molecular Genetics, 2011, 20, 193-201.	1.4	126
13	A 3-bp deletion in the HBS1L-MYB intergenic region on chromosome 6q23 is associated with HbF expression. Blood, 2011, 117, 4935-4945.	0.6	116
14	Variation in Nicotinic Acetylcholine Receptor Genes is Associated with Multiple Substance Dependence Phenotypes. Neuropsychopharmacology, 2010, 35, 1921-1931.	2.8	103
15	Two novel loci, COBL and SLC10A2, for Alzheimer's disease in African Americans. Alzheimer's and Dementia, 2017, 13, 119-129.	0.4	87
16	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. Nature Communications, 2020, 11, 5562.	5.8	80
17	Genome-Wide Association Study of Nicotine Dependence in American Populations: Identification of Novel Risk Loci in Both African-Americans and European-Americans. Biological Psychiatry, 2015, 77, 493-503.	0.7	78
18	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. JAMA Psychiatry, 2017, 74, 1234.	6.0	74

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19	Genome-wide Association Study Identifies a Regulatory Variant of RGMA Associated With Opioid Dependence in European Americans. Biological Psychiatry, 2018, 84, 762-770.	0.7	64
20	Genomewide Association Study for Maximum Number of Alcoholic Drinks in European Americans and African Americans. Alcoholism: Clinical and Experimental Research, 2015, 39, 1137-1147.	1.4	58
21	The genetics of alcohol dependence: Twin and SNPâ€based heritability, and genomeâ€wide association study based on AUDIT scores. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 739-748.	1.1	56
22	Variations in opioid receptor genes in neonatal abstinence syndrome. Drug and Alcohol Dependence, 2015, 155, 253-259.	1.6	55
23	Associations and Interactions Between SNPs in the Alcohol Metabolizing Genes and Alcoholism Phenotypes in European Americans. Alcoholism: Clinical and Experimental Research, 2009, 33, 848-857.	1.4	46
24	Identification of Novel Candidate Genes for Alzheimer's Disease by Autozygosity Mapping using Genome Wide SNP Data. Journal of Alzheimer's Disease, 2011, 23, 349-359.	1.2	46
25	Oxytocin receptor gene polymorphisms, attachment, and PTSD: Results from the National Health and Resilience in Veterans Study. Journal of Psychiatric Research, 2017, 94, 139-147.	1.5	46
26	Calibrating Longitudinal Cognition in Alzheimer's Disease Across Diverse Test Batteries and Datasets. Neuroepidemiology, 2014, 43, 194-205.	1.1	43
27	Genomewide Association Study of Alcohol Dependence Identifies Risk Loci Altering Ethanolâ€Response Behaviors in Model Organisms. Alcoholism: Clinical and Experimental Research, 2017, 41, 911-928.	1.4	43
28	Association of maternal and infant variants inPNOCandCOMTgenes with neonatal abstinence syndrome severity. American Journal on Addictions, 2017, 26, 42-49.	1.3	39
29	Population-Specific Risk of Type 2 Diabetes Conferred by HNF4A P2 Promoter Variants: A Lesson for Replication Studies. Diabetes, 2008, 57, 3161-3165.	0.3	37
30	Metaâ€analysis of genetic polymorphisms in granulomatosis with polyangiitis (Wegener's) reveals shared susceptibility loci with rheumatoid arthritis. Arthritis and Rheumatism, 2012, 64, 3463-3471.	6.7	33
31	Genomeâ€wide association study of rate of cognitive decline in Alzheimer's disease patients identifies novel genes and pathways. Alzheimer's and Dementia, 2020, 16, 1134-1145.	0.4	28
32	Shared genetic risk between eating disorderâ€and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	1.4	28
33	Power and Pitfalls of the Genome-Wide Association Study Approach to Identify Genes for Alzheimerâ€™s Disease. Current Psychiatry Reports, 2011, 13, 138-146.	2.1	27
34	Pharmacogenetic Effect of the Stromelysin (MMP3) Polymorphism on Stroke Risk in Relation to Antihypertensive Treatment. Stroke, 2011, 42, 330-335.	1.0	26
35	A Whole Genome Scan for Pulse Pressure/Stroke Volume Ratio in African Americans: The HyperGEN Study. American Journal of Hypertension, 2007, 20, 398-402.	1.0	25
36	Genetic modifiers of Hb E/Î²0 thalassemia identified by a two-stage genome-wide association study. BMC Medical Genetics, 2010, 11, 51.	2.1	25

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37	Genomeâ€wide association metaâ€analysis of age at first cannabis use. Addiction, 2018, 113, 2073-2086.	1.7	24
38	Post-GWAS analysis of six substance use traits improves the identification and functional interpretation of genetic risk loci. Drug and Alcohol Dependence, 2020, 206, 107703.	1.6	19
39	Genome-Wide Meta-Analyses of FTND and TTFC Phenotypes. Nicotine and Tobacco Research, 2020, 22, 900-909.	1.4	17
40	Nf1 Regulates Alcohol Dependence-Associated Excessive Drinking and Gamma-Aminobutyric Acid Release in the Central Amygdala in Mice and Is Associated with Alcohol Dependence in Humans. Biological Psychiatry, 2015, 77, 870-879.	0.7	14
41	ACSL6 Is Associated with the Number of Cigarettes Smoked and Its Expression Is Altered by Chronic Nicotine Exposure. PLoS ONE, 2011, 6, e28790.	1.1	11
42	The executive prominent/memory prominent spectrum in Alzheimer's disease is highly heritable. Neurobiology of Aging, 2016, 41, 115-121.	1.5	11
43	Evidence for a quantitative trait locus affecting low levels of apolipoprotein B and low density lipoprotein on chromosome 10 in Caucasian families. Journal of Lipid Research, 2007, 48, 2632-2639.	2.0	9
44	A Whole-Genome Scan for Stroke or Myocardial Infarction in Family Blood Pressure Program Families. Stroke, 2008, 39, 1115-1120.	1.0	9
45	S100A10 identified in a genome-wide gene Ã— cannabis dependence interaction analysis of risky sexual behaviours. Journal of Psychiatry and Neuroscience, 2017, 42, 252-261.	1.4	9
46	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. Exploration of Medicine, 2021, 2, 60-73.	1.5	6
47	Genome-wide association study of stimulant dependence. Translational Psychiatry, 2021, 11, 363.	2.4	4
48	Using linkage and association to identify and model genetic effects: summary of GAW15 Group 4. Genetic Epidemiology, 2007, 31, S34-S42.	0.6	3
49	Genome-Wide Studies in Sickle Cell Anemia Show Associations Between SNPs in the Olfactory Receptor Gene Cluster and Fetal Hemoglobin Concentration.. Blood, 2009, 114, 821-821.	0.6	2
50	No evidence for multiple loci affecting rheumatoid arthritis risk on chromosome 6p21. BMC Proceedings, 2007, 1, S42.	1.8	1
51	POLYGENIC RISK BURDEN AND CANNABIS USE COMORBIDITY IN PATIENTS WITH SCHIZOPHRENIA AND BIPOLAR DISORDER. European Neuropsychopharmacology, 2019, 29, S951.	0.3	1
52	P2-016: Identification of genetic variants associated with Alzheimer's disease: Progression rate. , 2015, 11, P487-P487.		0
53	EXPLORING THE ROLE OF GENETIC REGULATION OF GENE EXPRESSION IN SUBSTANCE USE AND DEPENDENCE. European Neuropsychopharmacology, 2019, 29, S803-S804.	0.3	0
54	GENOME-WIDE ASSOCIATION STUDY OF COMORBID ALCOHOL DEPENDENCE AND MAJOR DEPRESSION. European Neuropsychopharmacology, 2019, 29, S971.	0.3	0

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55	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. Exploration of Medicine, 0, , .	1.5	0
56	P90. A Genetically Informed Examination of Posttraumatic Stress Disorder and Traumatic Brain Injuryâ€™s Impact on Dementia Risk in US Veterans. Biological Psychiatry, 2022, 91, S123-S124.	0.7	0