

Richard Sherva

List of Publications by Year in descending order

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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	P90. A Genetically Informed Examination of Posttraumatic Stress Disorder and Traumatic Brain Injury's Impact on Dementia Risk in US Veterans. <i>Biological Psychiatry</i> , 2022, 91, S123-S124.	0.7	0
2	Shared genetic risk between eating disorder and substance use related phenotypes: Evidence from genome-wide association studies. <i>Addiction Biology</i> , 2021, 26, e12880.	1.4	28
3	Genome-wide association study of phenotypes measuring progression from first cocaine or opioid use to dependence reveals novel risk genes. <i>Exploration of Medicine</i> , 2021, 2, 60-73.	1.5	6
4	Genome-wide association study of stimulant dependence. <i>Translational Psychiatry</i> , 2021, 11, 363.	2.4	4
5	Genome-Wide Meta-Analyses of FTND and TTFC Phenotypes. <i>Nicotine and Tobacco Research</i> , 2020, 22, 900-909.	1.4	17
6	Post-GWAS analysis of six substance use traits improves the identification and functional interpretation of genetic risk loci. <i>Drug and Alcohol Dependence</i> , 2020, 206, 107703.	1.6	19
7	A large-scale genome-wide association study meta-analysis of cannabis use disorder. <i>Lancet Psychiatry</i> , 2020, 7, 1032-1045.	3.7	200
8	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. <i>Nature Communications</i> , 2020, 11, 5562.	5.8	80
9	Genome-wide association study of rate of cognitive decline in Alzheimer's disease patients identifies novel genes and pathways. <i>Alzheimer's and Dementia</i> , 2020, 16, 1134-1145.	0.4	28
10	EXPLORING THE ROLE OF GENETIC REGULATION OF GENE EXPRESSION IN SUBSTANCE USE AND DEPENDENCE. <i>European Neuropsychopharmacology</i> , 2019, 29, S803-S804.	0.3	0
11	GENOME-WIDE ASSOCIATION STUDY OF COMORBID ALCOHOL DEPENDENCE AND MAJOR DEPRESSION. <i>European Neuropsychopharmacology</i> , 2019, 29, S971.	0.3	0
12	POLYGENIC RISK BURDEN AND CANNABIS USE COMORBIDITY IN PATIENTS WITH SCHIZOPHRENIA AND BIPOLAR DISORDER. <i>European Neuropsychopharmacology</i> , 2019, 29, S951.	0.3	1
13	Genome-wide Association Study Identifies a Regulatory Variant of RGMA Associated With Opioid Dependence in European Americans. <i>Biological Psychiatry</i> , 2018, 84, 762-770.	0.7	64
14	Transancestral GWAS of alcohol dependence reveals common genetic underpinnings with psychiatric disorders. <i>Nature Neuroscience</i> , 2018, 21, 1656-1669.	7.1	490
15	Genome-wide association meta-analysis of age at first cannabis use. <i>Addiction</i> , 2018, 113, 2073-2086.	1.7	24
16	Association of maternal and infant variants in <i>PNOC</i> and <i>COMT</i> genes with neonatal abstinence syndrome severity. <i>American Journal on Addictions</i> , 2017, 26, 42-49.	1.3	39
17	Genomewide Association Study of Alcohol Dependence Identifies Risk Loci Altering Ethanol Response Behaviors in Model Organisms. <i>Alcoholism: Clinical and Experimental Research</i> , 2017, 41, 911-928.	1.4	43
18	Genetic Risk Variants Associated With Comorbid Alcohol Dependence and Major Depression. <i>JAMA Psychiatry</i> , 2017, 74, 1234.	6.0	74

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19	Oxytocin receptor gene polymorphisms, attachment, and PTSD: Results from the National Health and Resilience in Veterans Study. <i>Journal of Psychiatric Research</i> , 2017, 94, 139-147.	1.5	46
20	Two novel loci, <i>COBL</i> and <i>SLC10A2</i> , for Alzheimer's disease in African Americans. <i>Alzheimer's and Dementia</i> , 2017, 13, 119-129.	0.4	87
21	S100A10 identified in a genome-wide gene × cannabis dependence interaction analysis of risky sexual behaviours. <i>Journal of Psychiatry and Neuroscience</i> , 2017, 42, 252-261.	1.4	9
22	Genome-wide Association Study of Cannabis Dependence Severity, Novel Risk Variants, and Shared Genetic Risks. <i>JAMA Psychiatry</i> , 2016, 73, 472.	6.0	148
23	The executive prominent/memory prominent spectrum in Alzheimer's disease is highly heritable. <i>Neurobiology of Aging</i> , 2016, 41, 115-121.	1.5	11
24	The genetics of alcohol dependence: Twin and SNP-based heritability, and genome-wide association study based on AUDIT scores. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2015, 168, 739-748.	1.1	56
25	Genomewide Association Study for Maximum Number of Alcoholic Drinks in European Americans and African Americans. <i>Alcoholism: Clinical and Experimental Research</i> , 2015, 39, 1137-1147.	1.4	58
26	Variations in opioid receptor genes in neonatal abstinence syndrome. <i>Drug and Alcohol Dependence</i> , 2015, 155, 253-259.	1.6	55
27	P2-016: Identification of genetic variants associated with Alzheimer's disease: Progression rate. , 2015, 11, P487-P487.		0
28	Genome-Wide Association Study of Nicotine Dependence in American Populations: Identification of Novel Risk Loci in Both African-Americans and European-Americans. <i>Biological Psychiatry</i> , 2015, 77, 493-503.	0.7	78
29	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. <i>Biological Psychiatry</i> , 2015, 77, 870-879.	0.7	14
30	Genome-wide association study of the rate of cognitive decline in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2014, 10, 45-52.	0.4	147
31	Genome-Wide Association Study of Opioid Dependence: Multiple Associations Mapped to Calcium and Potassium Pathways. <i>Biological Psychiatry</i> , 2014, 76, 66-74.	0.7	192
32	Calibrating Longitudinal Cognition in Alzheimer's Disease Across Diverse Test Batteries and Datasets. <i>Neuroepidemiology</i> , 2014, 43, 194-205.	1.1	43
33	Association of Granulomatosis With Polyangiitis (Wegener's) With <i>HLA-DPB1*04</i> and <i>SEMA6A</i> Gene Variants: Evidence From Genome-Wide Analysis. <i>Arthritis and Rheumatism</i> , 2013, 65, 2457-2468.	6.7	138
34	Meta-analysis of genetic polymorphisms in granulomatosis with polyangiitis (Wegener's) reveals shared susceptibility loci with rheumatoid arthritis. <i>Arthritis and Rheumatism</i> , 2012, 64, 3463-3471.	6.7	33
35	Identification of Novel Candidate Genes for Alzheimer's Disease by Autozygosity Mapping using Genome Wide SNP Data. <i>Journal of Alzheimer's Disease</i> , 2011, 23, 349-359.	1.2	46
36	A 3-bp deletion in the HBS1L-MYB intergenic region on chromosome 6q23 is associated with HbF expression. <i>Blood</i> , 2011, 117, 4935-4945.	0.6	116

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37	ACSL6 Is Associated with the Number of Cigarettes Smoked and Its Expression Is Altered by Chronic Nicotine Exposure. <i>PLoS ONE</i> , 2011, 6, e28790.	1.1	11
38	Power and Pitfalls of the Genome-Wide Association Study Approach to Identify Genes for Alzheimer's Disease. <i>Current Psychiatry Reports</i> , 2011, 13, 138-146.	2.1	27
39	Common CD36 SNPs reduce protein expression and may contribute to a protective atherogenic profile. <i>Human Molecular Genetics</i> , 2011, 20, 193-201.	1.4	126
40	Pharmacogenetic Effect of the Stromelysin (MMP3) Polymorphism on Stroke Risk in Relation to Antihypertensive Treatment. <i>Stroke</i> , 2011, 42, 330-335.	1.0	26
41	Fetal hemoglobin in sickle cell anemia: genome-wide association studies suggest a regulatory region in the 5q22 olfactory receptor gene cluster. <i>Blood</i> , 2010, 115, 1815-1822.	0.6	146
42	Genetic modifiers of Hb E β 0 thalassemia identified by a two-stage genome-wide association study. <i>BMC Medical Genetics</i> , 2010, 11, 51.	2.1	25
43	Multiple Independent Loci at Chromosome 15q25.1 Affect Smoking Quantity: a Meta-Analysis and Comparison with Lung Cancer and COPD. <i>PLoS Genetics</i> , 2010, 6, e1001053.	1.5	332
44	Variation in Nicotinic Acetylcholine Receptor Genes is Associated with Multiple Substance Dependence Phenotypes. <i>Neuropsychopharmacology</i> , 2010, 35, 1921-1931.	2.8	103
45	Associations and Interactions Between SNPs in the Alcohol Metabolizing Genes and Alcoholism Phenotypes in European Americans. <i>Alcoholism: Clinical and Experimental Research</i> , 2009, 33, 848-857.	1.4	46
46	Genome-Wide Studies in Sickle Cell Anemia Show Associations Between SNPs in the Olfactory Receptor Gene Cluster and Fetal Hemoglobin Concentration.. <i>Blood</i> , 2009, 114, 821-821.	0.6	2
47	Association of a single nucleotide polymorphism in neuronal acetylcholine receptor subunit alpha 5 (CHRNA5) with smoking status and with "pleasurable buzz" during early experimentation with smoking. <i>Addiction</i> , 2008, 103, 1544-1552.	1.7	129
48	A Whole-Genome Scan for Stroke or Myocardial Infarction in Family Blood Pressure Program Families. <i>Stroke</i> , 2008, 39, 1115-1120.	1.0	9
49	Population-Specific Risk of Type 2 Diabetes Conferred by HNF4A P2 Promoter Variants: A Lesson for Replication Studies. <i>Diabetes</i> , 2008, 57, 3161-3165.	0.3	37
50	Variants in the CD36 gene associate with the metabolic syndrome and high-density lipoprotein cholesterol. <i>Human Molecular Genetics</i> , 2008, 17, 1695-1704.	1.4	164
51	Evidence for a quantitative trait locus affecting low levels of apolipoprotein B and low density lipoprotein on chromosome 10 in Caucasian families. <i>Journal of Lipid Research</i> , 2007, 48, 2632-2639.	2.0	9
52	A Whole Genome Scan for Pulse Pressure/Stroke Volume Ratio in African Americans: The HyperGEN Study. <i>American Journal of Hypertension</i> , 2007, 20, 398-402.	1.0	25
53	No evidence for multiple loci affecting rheumatoid arthritis risk on chromosome 6p21. <i>BMC Proceedings</i> , 2007, 1, S42.	1.8	1
54	Using linkage and association to identify and model genetic effects: summary of GAW15 Group 4. <i>Genetic Epidemiology</i> , 2007, 31, S34-S42.	0.6	3

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55	Common variants in WFS1 confer risk of type 2 diabetes. Nature Genetics, 2007, 39, 951-953.	9.4	333
56	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