Dawei Li

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6804610/publications.pdf

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23	1,625	623734	642732
papers	citations	14 h-index	23 g-index

24	24	24	2882
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Meta-analysis shows significant association between dopamine system genes and attention deficit hyperactivity disorder (ADHD). Human Molecular Genetics, 2006, 15, 2276-2284.	2.9	519
2	Meta-analysis shows strong positive association of the neuregulin 1 (NRG1) gene with schizophrenia. Human Molecular Genetics, 2006, 15, 1995-2002.	2.9	260
3	Strong Association of the Alcohol Dehydrogenase 1B Gene (ADH1B) with Alcohol Dependence and Alcohol-Induced Medical Diseases. Biological Psychiatry, 2011, 70, 504-512.	1.3	150
4	Strong protective effect of the aldehyde dehydrogenase gene (ALDH2) 504lys (*2) allele against alcoholism and alcohol-induced medical diseases in Asians. Human Genetics, 2012, 131, 725-737.	3.8	132
5	Association of Gamma-Aminobutyric Acid A Receptor Î±2 Gene (GABRA2) with Alcohol Use Disorder. Neuropsychopharmacology, 2014, 39, 907-918.	5.4	93
6	Association study of serotonin 2A receptor (5-HT2A) gene with schizophrenia and suicidal behavior using systematic meta-analysis. Biochemical and Biophysical Research Communications, 2006, 340, 1006-1015.	2.1	86
7	Association of the HTR2A gene with alcohol and heroin abuse. Human Genetics, 2014, 133, 357-365.	3.8	56
8	Associations of the 5â€hydroxytryptamine (serotonin) Receptor 1B gene (HTR1B) with alcohol, cocaine, and heroin abuse. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2013, 162, 169-176.	1.7	48
9	A virome-wide clonal integration analysis platform for discovering cancer viral etiology. Genome Research, 2019, 29, 819-830.	5.5	47
10	Multi-Cultural Association of the Serotonin Transporter Gene (SLC6A4) with Substance Use Disorder. Neuropsychopharmacology, 2013, 38, 1737-1747.	5.4	42
11	Genome-Wide Association Study of Copy Number Variations (CNVs) with Opioid Dependence. Neuropsychopharmacology, 2015, 40, 1016-1026.	5.4	39
12	Comprehensive comparative analysis of methods and software for identifying viral integrations. Briefings in Bioinformatics, 2019, 20, 2088-2097.	6.5	32
13	Searching for human oncoviruses: Histories, challenges, and opportunities. Journal of Cellular Biochemistry, 2018, 119, 4897-4906.	2.6	26
14	ERVcaller: identifying polymorphic endogenous retrovirus and other transposable element insertions using whole-genome sequencing data. Bioinformatics, 2019, 35, 3913-3922.	4.1	25
15	Profile of circulating microRNAs in myalgic encephalomyelitis and their relation to symptom severity, and disease pathophysiology. Scientific Reports, 2020, 10, 19620.	3.3	24
16	Atlas of human diseases influenced by genetic variants with extreme allele frequency differences. Human Genetics, 2017, 136, 39-54.	3.8	15
17	GACT: a Genome build and Allele definition Conversion Tool for SNP imputation and meta-analysis in genetic association studies. BMC Genomics, 2014, 15, 610.	2.8	7
18	Eye color: A potential indicator of alcohol dependence risk in European Americans. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 347-353.	1.7	7

#	Article	IF	Citations
19	Characterization of Hepatitis B Virus Integrations Identified in Hepatocellular Carcinoma Genomes. Viruses, 2021, 13, 245.	3.3	6
20	VIpower: Simulation-based tool for estimating power of viral integration detection via high-throughput sequencing. Genomics, 2020, 112, 207-211.	2.9	4
21	Association of Circulating YKL-40 Levels and CHI3L1 Variants with the Risk of Spinal Deformity Progression in Adolescent Idiopathic Scoliosis. Scientific Reports, 2019, 9, 5712.	3.3	3
22	Sequencing facility and DNA source associated patterns of virus-mappable reads in whole-genome sequencing data. Genomics, 2021, 113, 1189-1198.	2.9	3
23	Further analyses support the association between light eye color and alcohol dependence. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2015, 168, 757-760.	1.7	1