Qian Peng

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3670415/publications.pdf

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	1478505	1372567
160	6	10
citations	h-index	g-index
1.1		5.60
11	11	563
docs citations	times ranked	citing authors
	citations 11	160 6 citations h-index 11 11

#	Article	IF	CITATIONS
1	Combinatorial interactions of genetic variants in human cardiomyopathy. Nature Biomedical Engineering, 2019, 3, 147-157.	22.5	37
2	Large-scale genomics unveil polygenic architecture of human cortical surface area. Nature Communications, 2015, 6, 7549.	12.8	30
3	Genetic loci for alcohol-related life events and substance-induced affective symptoms: indexing the $\hat{a}\in \infty$ dark side $\hat{a}\in \infty$ addiction. Translational Psychiatry, 2019, 9, 71.	4.8	19
4	Association and ancestry analysis of sequence variants in ADH and ALDH using alcoholâ€related phenotypes in a Native American community sample. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2014, 165, 673-683.	1.7	17
5	Conservation of Distinct Genetically-Mediated Human Cortical Pattern. PLoS Genetics, 2016, 12, e1006143.	3.5	15
6	Two single nucleotide polymorphisms in IL13 and IL13RA1 from individuals with idiopathic Parkinson's disease increase cellular susceptibility to oxidative stress. Brain, Behavior, and Immunity, 2020, 88, 920-924.	4.1	11
7	Fine mapping and subphenotyping implicates <i>ADRA1B</i> gene variants in psoriasis susceptibility in a Chinese population. Epigenomics, 2019, 11, 455-467.	2.1	10
8	Associations Between Genomic Variants in Alcohol Dehydrogenase Genes and Alcohol Symptomatology in American Indians and European Americans: Distinctions and Convergence. Alcoholism: Clinical and Experimental Research, 2017, 41, 1695-1704.	2.4	7
9	Common genetic substrates of alcohol and substance use disorder severity revealed by pleiotropy detection against GWAS catalog in two populations. Addiction Biology, 2021, 26, e12877.	2.6	7
10	Whole genome sequence association and ancestryâ€informed polygenic profile of EEG alpha in a Native American population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2017, 174, 435-450.	1.7	5
11	Long tracks of homozygosity predict the severity of alcohol use disorders in an American Indian population. Molecular Psychiatry, 2021, 26, 2200-2211.	7.9	2