Jin P Szatkiewicz

List of Publications by Year
in descending order

24 papers	2,903 citations	759233 12 h-index	610901 24 g-index
28 all docs	28 docs citations	28 times ranked	4959 citing authors

#	Article	IF	Citations
1	Mapping genomic loci implicates genes and synaptic biology in schizophrenia. Nature, 2022, 604, 502-508.	27.8	929
2	The impact of educational attainment, intelligence and intellectual disability on schizophrenia: a Swedish population-based register and genetic study. Molecular Psychiatry, 2022, 27, 2439-2447.	7.9	10
3	Shared genetic risk between eating disorderâ€•and substanceâ€useâ€related phenotypes: Evidence from genomeâ€wide association studies. Addiction Biology, 2021, 26, e12880.	2.6	28
4	Neuronal and glial 3D chromatin architecture informs the cellular etiology of brain disorders. Nature Communications, 2021, 12, 3968.	12.8	48
5	Elevated common variant genetic risk for tourette syndrome in a densely-affected pedigree. Molecular Psychiatry, 2021, 26, 7522-7529.	7.9	8
6	Characterization of Single Gene Copy Number Variants in Schizophrenia. Biological Psychiatry, 2020, 87, 736-744.	1.3	10
7	Treatment-resistant psychotic symptoms and early-onset dementia: A case report of the 3q29 deletion syndrome. Schizophrenia Research, 2020, 224, 195-197.	2.0	8
8	Association test using Copy Number Profile Curves (CONCUR) enhances power in rare copy number variant analysis. PLoS Computational Biology, 2020, 16, e1007797.	3.2	6
9	Increased burden of ultra-rare structural variants localizing to boundaries of topologically associated domains in schizophrenia. Nature Communications, 2020, 11, 1842.	12.8	56
10	Treatment-resistant psychotic symptoms and the 15q11.2 BP1â€“BP2 (Burnside-Butler) deletion syndrome: case report and review of the literature. Translational Psychiatry, 2020, 10, 42.	4.8	11
11	Genome-wide association study identifies eight risk loci and implicates metabo-psychiatric origins for anorexia nervosa. Nature Genetics, 2019, 51, 1207-1214.	21.4	641
12	The genomics of major psychiatric disorders in a large pedigree from Northern Sweden. Translational Psychiatry, 2019, 9, 60.	4.8	15
13	Common-variant associations with fragile X syndrome. Molecular Psychiatry, 2019, 24, 338-344.	7.9	8
14	A randomized approach to speed up the analysis of large-scale read-count data in the application of CNV detection. BMC Bioinformatics, 2018, 19, 74.	2.6	1
15	Developmental Delay, Treatment-Resistant Psychosis, and Early-Onset Dementia in a Man With 22q11 Deletion Syndrome and Huntingtonâ€™s Disease. American Journal of Psychiatry, 2018, 175, 400-407.	7.2	9
16	Cover Image, Volume 173A, Number 2, February 2017. American Journal of Medical Genetics, Part A, 2017, 173, i.	1.2	0
17	Implication of LRRC4C and DPP6 in neurodevelopmental disorders. American Journal of Medical Genetics, Part A, 2017, 173, 395-406.	1.2	40
18	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	21.4	838

#	Article	IF	Citations
19	Exploration of large, rare copy number variants associated with psychiatric and neurodevelopmental disorders in individuals with anorexia nervosa. Psychiatric Genetics, 2017, 27, 152-158.	1.1	18
20	One CNV Discordance in NRXN1 Observed Upon Genome-wide Screening in 38 Pairs of Adult Healthy Monozygotic Twins. Twin Research and Human Genetics, 2016, 19, 97-103.	0.6	2
21	A New Method for Detecting Associations with Rare Copy-Number Variants. PLoS Genetics, 2015, 11, e1005403.	3.5	14
22	Allele-specific copy-number discovery from whole-genome and whole-exome sequencing. Nucleic Acids Research, 2015, 43, e90-e90.	14.5	16
23	Clozapine-induced agranulocytosis is associated with rare HLA-DQB1 and HLA-B alleles. Nature Communications, 2014, 5, 4757.	12.8	153
24	Improving detection of copy-number variation by simultaneous bias correction and read-depth segmentation. Nucleic Acids Research, 2013, 41, 1519-1532.	14.5	33