Peristera Paschou

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7077109/publications.pdf

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45 papers	2,178 citations	257101 24 h-index	243296 44 g-index
48 all docs	48 docs citations	48 times ranked	3918 citing authors

#	Article	IF	Citations
1	Reconstructing SNP allele and genotype frequencies from GWAS summary statistics. Scientific Reports, 2022, 12, 8242.	1.6	0
2	Synaptic processes and immune-related pathways implicated in Tourette syndrome. Translational Psychiatry, 2021, 11, 56.	2.4	31
3	Integrating Linguistics, Social Structure, and Geography to Model Genetic Diversity within India. Molecular Biology and Evolution, 2021, 38, 1809-1819.	3.5	7
4	Candidate Genes and Pathways Associated with Gilles de la Tourette Syndromeâ€"Where Are We?. Genes, 2021, 12, 1321.	1.0	9
5	Investigating Shared Genetic Basis Across Tourette Syndrome and Comorbid Neurodevelopmental Disorders Along the Impulsivity-Compulsivity Spectrum. Biological Psychiatry, 2021, 90, 317-327.	0.7	49
6	Association between CUBN gene variants, type 2 diabetes and vitamin D concentrations in an elderly Greek population. Journal of Steroid Biochemistry and Molecular Biology, 2020, 198, 105549.	1.2	7
7	Genetic variation in CARD8, a gene coding for an NLRP3 inflammasome-associated protein, alters the genetic risk for diabetic nephropathy in the context of type 2 diabetes mellitus. Diabetes and Vascular Disease Research, 2020, 17, 147916412097089.	0.9	6
8	Association of Genetic Variation in the 3'UTR of LHX6, IMMP2L, and AADAC With Tourette Syndrome. Frontiers in Neurology, 2020, 11, 803.	1.1	13
9	Population structure of modern-day Italians reveals patterns of ancient and archaic ancestries in Southern Europe. Science Advances, 2019, 5, eaaw3492.	4.7	53
10	Genetic history of the population of Crete. Annals of Human Genetics, 2019, 83, 373-388.	0.3	2
11	Interrogating the Genetic Determinants of Touretteâ€™s Syndrome and Other Tic Disorders Through Genome-Wide Association Studies. American Journal of Psychiatry, 2019, 176, 217-227.	4.0	242
12	TeraPCA: a fast and scalable software package to study genetic variation in tera-scale genotypes. Bioinformatics, 2019, 35, 3679-3683.	1.8	31
13	Association of ALOX12 gene polymorphism with all-cause and cardiovascular mortality in diabetic nephropathy. International Urology and Nephrology, 2018, 50, 321-329.	0.6	20
14	De Novo Sequence and Copy Number Variants Are Strongly Associated with Tourette Disorder and Implicate Cell Polarity in Pathogenesis. Cell Reports, 2018, 24, 3441-3454.e12.	2.9	91
15	Gilles de la Tourette syndrome. Nature Reviews Disease Primers, 2017, 3, 16097.	18.1	257
16	Genetics of the peloponnesean populations and the theory of extinction of the medieval peloponnesean Greeks. European Journal of Human Genetics, 2017, 25, 637-645.	1.4	22
17	Rare Copy Number Variants in NRXN1 and CNTN6 Increase Risk for Tourette Syndrome. Neuron, 2017, 94, 1101-1111.e7.	3.8	137
18	17q21.31 duplication causes prominent tau-related dementia with increased MAPT expression. Molecular Psychiatry, 2017, 22, 1119-1125.	4.1	57

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19	Candidate gene investigation of spinal degenerative osteoarthritis in Greek population. Spine Journal, 2017, 17, 1881-1888.	0.6	6
20	Variant Ranker: a web-tool to rank genomic data according to functional significance. BMC Bioinformatics, 2017, 18, 341.	1.2	21
21	Meta-Analysis of Tourette Syndrome and Attention Deficit Hyperactivity Disorder Provides Support for a Shared Genetic Basis. Frontiers in Neuroscience, 2016, 10, 340.	1.4	26
22	The Genetic Etiology of Tourette Syndrome: Large-Scale Collaborative Efforts on the Precipice of Discovery. Frontiers in Neuroscience, 2016, 10, 351.	1.4	45
23	Targeted Re-Sequencing Approach of Candidate Genes Implicates Rare Potentially Functional Variants in Tourette Syndrome Etiology. Frontiers in Neuroscience, 2016, 10, 428.	1.4	29
24	Investigation of SNP rs2060546 Immediately Upstream to NTN4 in a Danish Gilles de la Tourette Syndrome Cohort. Frontiers in Neuroscience, 2016, 10, 531.	1.4	5
25	TS-EUROTRAIN: A European-Wide Investigation and Training Network on the Etiology and Pathophysiology of Gilles de la Tourette Syndrome. Frontiers in Neuroscience, 2016, 10, 384.	1.4	21
26	The Genetics of Gilles de la Tourette Syndrome: a Common Aetiological Basis with Comorbid Disorders?. Current Behavioral Neuroscience Reports, 2016, 3, 218-231.	0.6	7
27	Familial early-onset dementia with complex neuropathologic phenotype and genomic background. Neurobiology of Aging, 2016, 42, 199-204.	1.5	16
28	Comment: Dissecting the genetic architecture of Tourette syndrome into subphenotypes. Neurology, 2016, 87, 503-503.	1.5	1
29	Association of AADAC Deletion and Gilles de la Tourette Syndrome in a Large European Cohort. Biological Psychiatry, 2016, 79, 383-391.	0.7	41
30	Epigenome-Wide Association Study of Tic Disorders. Twin Research and Human Genetics, 2015, 18, 699-709.	0.3	31
31	Reduced genetic variation and strong genetic population structure in the freshwater killifishValencia letourneuxi(Valenciidae) based on nuclear and mitochondrial markers. Biological Journal of the Linnean Society, 2014, 111, 334-349.	0.7	14
32	Intragenic deletions affecting two alternative transcripts of the IMMP2L gene in patients with Tourette syndrome. European Journal of Human Genetics, 2014, 22, 1283-1289.	1.4	69
33	Genetic association signal near <scp>NTN</scp>4 in <scp>T</scp>ourette syndrome. Annals of Neurology, 2014, 76, 310-315.	2.8	53
34	Maritime route of colonization of Europe. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 9211-9216.	3.3	71
35	The genetic basis of Gilles de la Tourette Syndrome. Neuroscience and Biobehavioral Reviews, 2013, 37, 1026-1039.	2.9	105
36	A European population in Minoan Bronze Age Crete. Nature Communications, 2013, 4, 1861.	5.8	31

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37	Genetic Susceptibility and Neurotransmitters in Tourette Syndrome. International Review of Neurobiology, 2013, 112, 155-177.	0.9	48
38	A Brief History of the European Society for the Study of Tourette Syndrome. Behavioural Neurology, 2013, 27, 3-5.	1.1	16
39	Evaluation of the LIM homeobox genes LHX6 and LHX8 as candidates for Tourette syndrome. Genes, Brain and Behavior, 2012, 11, 444-451.	1.1	23
40	Genetic variation in the visfatin (PBEF1/NAMPT) gene and type 2 diabetes in the Greek population. Cytokine, 2010, 51, 25-27.	1.4	13
41	Ancestry informative markers for fine-scale individual assignment to worldwide populations. Journal of Medical Genetics, 2010, 47, 835-847.	1.5	65
42	Tracing Sub-Structure in the European American Population with PCA-Informative Markers. PLoS Genetics, 2008, 4, e1000114.	1.5	60
43	PCA-Correlated SNPs for Structure Identification in Worldwide Human Populations. PLoS Genetics, 2007, 3, e160.	1.5	224
44	Intra- and interpopulation genotype reconstruction from tagging SNPs. Genome Research, 2006, 17, 96-107.	2.4	35
45	Indications of Linkage and Association of Gilles de la Tourette Syndrome in Two Independent Family Samples: 17q25 Is a Putative Susceptibility Region. American Journal of Human Genetics, 2004, 75, 545-560.	2.6	62