Burcu F Darst

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5365606/publications.pdf

Version: 2024-02-01

47 papers

2,167 citations

331670 21 h-index 243625 44 g-index

56 all docs

56 docs citations 56 times ranked 4563 citing authors

#	Article	IF	CITATIONS
1	The gut microbiota-derived metabolite trimethylamine N-oxide is elevated in Alzheimer's disease. Alzheimer's Research and Therapy, 2018, 10, 124.	6.2	273
2	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nature Genetics, 2021, 53, 65-75.	21.4	264
3	Using recursive feature elimination in random forest to account for correlated variables in high dimensional data. BMC Genetics, 2018, 19, 65.	2.7	222
4	Multimodal imaging of the self-regulating developing brain. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 19620-19625.	7.1	192
5	Long-term influence of normal variation in neonatal characteristics on human brain development. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20089-20094.	7.1	158
6	Impact of direct-to-consumer genomic testing at long term follow-up. Journal of Medical Genetics, 2013, 50, 393-400.	3.2	125
7	Longitudinal plasma metabolomics of aging and sex. Aging, 2019, 11, 1262-1282.	3.1	115
8	Pathway-Specific Polygenic Risk Scores as Predictors of Amyloid-β Deposition and Cognitive Function in a Sample at Increased Risk for Alzheimer's Disease. Journal of Alzheimer's Disease, 2016, 55, 473-484.	2.6	93
9	Direct-to-consumer personalized genomic testing. Human Molecular Genetics, 2011, 20, R132-R141.	2.9	84
10	<i>BDNF</i> Val66Met predicts cognitive decline in the Wisconsin Registry for Alzheimer's Prevention. Neurology, 2017, 88, 2098-2106.	1.1	52
11	Integrated analysis of genomics, longitudinal metabolomics, and Alzheimer's risk factors among 1,111 cohort participants. Genetic Epidemiology, 2019, 43, 657-674.	1.3	41
12	<i>KLOTHO</i> heterozygosity attenuates <i>APOE4</i> -related amyloid burden in preclinical AD. Neurology, 2019, 92, e1878-e1889.	1.1	40
13	Germline Sequencing DNA Repair Genes in 5545 Men With Aggressive and Nonaggressive Prostate Cancer. Journal of the National Cancer Institute, 2021, 113, 616-625.	6.3	40
14	Combined Effect of a Polygenic Risk Score and Rare Genetic Variants on Prostate Cancer Risk. European Urology, 2021, 80, 134-138.	1.9	39
15	Cardiorespiratory fitness alters the influence of a polygenic risk score on biomarkers of AD. Neurology, 2017, 88, 1650-1658.	1.1	35
16	A genome sequencing program for novel undiagnosed diseases. Genetics in Medicine, 2015, 17, 995-1001.	2.4	32
17	A Germline Variant at 8q24 Contributes to Familial Clustering of Prostate Cancer in Men of African Ancestry. European Urology, 2020, 78, 316-320.	1.9	32
18	Macrovascular and microvascular cerebral blood flow in adults at risk for Alzheimer's disease. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2017, 7, 48-55.	2.4	31

#	Article	IF	CITATIONS
19	Characteristics of Genomic Test Consumers Who Spontaneously Share Results With Their Health Care Provider. Health Communication, 2014, 29, 105-108.	3.1	28
20	Cerebrospinal fluid metabolomics identifies 19 brain-related phenotype associations. Communications Biology, 2021, 4, 63.	4.4	28
21	Dyslexia and language impairment associated genetic markers influence cortical thickness and white matter in typically developing children. Brain Imaging and Behavior, 2016, 10, 272-282.	2.1	27
22	Candidate SNP Associations of Optimism and Resilience in Older Adults: Exploratory Study of 935 Community-Dwelling Adults. American Journal of Geriatric Psychiatry, 2014, 22, 997-1006.e5.	1.2	23
23	Genetic discovery and risk characterization in type 2 diabetes across diverse populations. Human Genetics and Genomics Advances, 2021, 2, 100029.	1.7	23
24	The Four-Kallikrein Panel Is Effective in Identifying Aggressive Prostate Cancer in a Multiethnic Population. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1381-1388.	2.5	22
25	A Rare Germline HOXB13 Variant Contributes to Risk of Prostate Cancer in Men of African Ancestry. European Urology, 2022, 81, 458-462.	1.9	22
26	The effect of rare variants in TREM2 and PLD3 on longitudinal cognitive function in the Wisconsin Registry for Alzheimer's Prevention. Neurobiology of Aging, 2018, 66, 177.e1-177.e5.	3.1	18
27	Multi-ethnic GWAS and fine-mapping of glycaemic traits identify novel loci in the PAGE Study. Diabetologia, 2022, 65, 477-489.	6.3	15
28	Heritability of Cognitive Traits Among Siblings with a Parental History of Alzheimer's Disease. Journal of Alzheimer's Disease, 2015, 45, 1149-1155.	2.6	11
29	Including diverse and admixed populations in genetic epidemiology research. Genetic Epidemiology, 2022, 46, 347-371.	1.3	11
30	Carrier frequency estimation of Zellweger spectrum disorder using ExAC database and bioinformatics tools. Genetics in Medicine, 2019, 21, 1969-1976.	2.4	10
31	Association Between a 22-feature Genomic Classifier and Biopsy Gleason Upgrade During Active Surveillance for Prostate Cancer. European Urology Open Science, 2022, 37, 113-119.	0.4	10
32	Patient perspectives on whole-genome sequencing for undiagnosed diseases. Personalized Medicine, 2017, 14, 17-25.	1.5	8
33	Metabolites Associated with Early Cognitive Changes Implicated in Alzheimer's Disease. Journal of Alzheimer's Disease, 2021, 79, 1041-1054.	2.6	4
34	Data mining and machine learning approaches for the integration of genome-wide association and methylation data: methodology and main conclusions from GAW20. BMC Genetics, 2018, 19, 76.	2.7	2
35	Findings from a Genetic Sequencing Investigation of Men with Familial and Aggressive Prostate Cancer. European Urology, 2021, 79, 362-363.	1.9	2
36	Genome-wide association study of pancreatic fat: The Multiethnic Cohort Adiposity Phenotype Study. PLoS ONE, 2021, 16, e0249615.	2.5	2

3

#	Article	IF	Citations
37	CSF metabolites associate with CSF tau and improve prediction of Alzheimer's disease status. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2021, 13, e12167.	2.4	2
38	Genetic variants predictive of chemotherapy-induced peripheral neuropathy symptoms in gynecologic cancer survivors. Gynecologic Oncology, 2021, 163, 578-582.	1.4	2
39	P4-004: Cardiorespiratory capacity modifies the association between a polygenic risk score and CSF biomarkers in preclinical Alzheimer's disease., 2015, 11, P766-P766.		1
40	Transmission and decorrelation methods for detecting rare variants using sequencing data from related individuals. BMC Proceedings, 2016, 10, 203-207.	1.6	1
41	Prioritization of family member sequencing for the detection of rare variants. BMC Proceedings, 2016, 10, 227-231.	1.6	1
42	Sign-based Shrinkage Based on an Asymmetric LASSO Penalty. Journal of Data Science, 2021, 19, 429-449.	0.9	1
43	P4-193: Effect of TREM2 and PLD3 in an Alzheimer's disease family history cohort: The wisconsin registry for Alzheimer's prevention (WRAP)., 2015, 11, P854-P854.		0
44	IC-P-044: Cardiorespiratory capacity modifies the association between a polygenic risk score and CSF biomarkers in preclinical Alzheimer's disease., 2015, 11, P37-P38.		0
45	[O2–02–01]: PRELIMINARY FINDINGS FROM AN ONGOING LONGITUDINAL METABOLOMEâ€WIDE ASSOCIAT STUDY OF COGNITIVE DECLINE IN HEALTHY ADULTS WITH INCREASED RISK FOR ALZHEIMER's DISEASE. Alzheimer's and Dementia, 2017, 13, P551.	TION 0.8	O
46	O3â€03â€05: INTEGRATIVE NETWORK ANALYSIS IDENTIFIES RELATIONSHIPS BETWEEN METABOLOMICS, GENOMICS, AND RISK FACTORS FOR AD. Alzheimer's and Dementia, 2018, 14, P1016.	0.8	0
47	O3â€06â€03: TISSUEâ€6PECIFIC POLYGENIC RISK SCORES PREDICT AMYLOID BIOMARKERS. Alzheimer's and Dementia, 2018, 14, P1026.	0.8	0