Farid Rajabli

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/10711930/publications.pdf

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38	471	1040056	839539
papers	citations	9	18
		h-index	g-index

44	44	44	1375
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Identifying differential regulatory control of APOE É›4 on African versus European haplotypes as potential therapeutic targets. Alzheimer's and Dementia, 2022, 18, 1930-1942.	0.8	12
2	A locus at 19q13.31 significantly reduces the ApoE Îµ4 risk for Alzheimerâ€™s Disease in African Ancestry. PLoS Genetics, 2022, 18, e1009977.	3.5	19
3	Dissecting the role of Amerindian genetic ancestry and the ApoE Îµ4 allele on Alzheimer disease in an admixed Peruvian population. Neurobiology of Aging, 2021, 101, 298.e11-298.e15.	3.1	11
4	Increased APOE Îµ4 expression is associated with the difference in Alzheimer's disease risk from diverse ancestral backgrounds. Alzheimer's and Dementia, 2021, 17, 1179-1188.	0.8	33
5	Linkage of Alzheimer disease families with Puerto Rican ancestry identifies a chromosome 9 locus. Neurobiology of Aging, 2021, 104, 115.e1-115.e7.	3.1	4
6	Assessment of ADâ€related plasma biomarkers in diverse ancestral populations. Alzheimer's and Dementia, 2021, 17, .	0.8	0
7	Does higher educational attainment influence functional capabilities among African Americans with Alzheimerâ€™s disease?. Alzheimer's and Dementia, 2021, 17, .	0.8	0
8	Ancestryâ€specific intronic variants on the APOE É›4 haplotype influence enhancer activity and interaction with APOE promoter. Alzheimer's and Dementia, 2021, 17, e055266.	0.8	0
9	APOEâ€stratified genomeâ€wide association analysis identifies novel Alzheimer disease candidate risk loci for African Americans. Alzheimer's and Dementia, 2021, 17, e056383.	0.8	2
10	ATAC-seq on iPSC derived astrocytes to assess chromatin accessibility differences between African and European local ancestry.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056086.	0.8	0
11	African locus reduces the effect of ApoE Îµ4 allele in Alzheimer's disease.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056210.	0.8	0
12	Admixture mapping identifies novel regions influencing Alzheimer disease in African Americans.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e056443.	0.8	0
13	Whole exome sequencing study identifies novel rare and common Alzheimerâ€™s-Associated variants involved in immune response and transcriptional regulation. Molecular Psychiatry, 2020, 25, 1859-1875.	7.9	191
14	Novel Variants in LRRK2 and GBA Identified in Latino Parkinson Disease Cohort Enriched for Caribbean Origin. Frontiers in Neurology, 2020, 11, 573733.	2.4	6
15	Comparative transâ€ethnic metaâ€analysis of whole exome sequencing variation for Alzheimerâ€™s disease (AD) in 18,402 individuals of the Alzheimerâ€™s Disease Sequencing Project (ADSP). Alzheimer's and Dementia, 2020, 16, e041583.	0.8	0
16	Exploring the role of Amerindian genetic ancestry and ApoEÎµ4 gene on Alzheimer disease in the Peruvian population. Alzheimer's and Dementia, 2020, 16, e045012.	0.8	0
17	Increased APOEâ€e4 expression is associated with reactive A1 astrocytes and may confer the difference in Alzheimer disease risk from different ancestral backgrounds. Alzheimer's and Dementia, 2020, 16, e045415.	0.8	0
18	Transcriptomic characterization of a Puerto Rican Alzheimer disease cohort implicates convergent immuneâ€related pathways. Alzheimer's and Dementia, 2020, 16, e045890.	0.8	0

#	Article	IF	Citations
19	Southern European genetic ancestry shows reduced APOE E4 risk for Alzheimer disease in Caribbean Hispanic population. Alzheimer's and Dementia, 2020, 16, e045951.	0.8	0
20	Identification of differential regulation of European versus African local ancestry haplotypes surrounding ApoEÎµ4. Alzheimer's and Dementia, 2020, 16, e046016.	0.8	0
21	Functional analysis of candidate genes identified through whole genome sequencing in Caribbean Hispanic families for lateâ€onset Alzheimer disease. Alzheimer's and Dementia, 2020, 16, e046017.	0.8	1
22	The effect of global ancestry and diabetes on the 3MS score in older Puerto Ricans. Alzheimer's and Dementia, 2020, 16, e046051.	0.8	0
23	Mapping Alzheimer diseaseâ€“associated regions in the African American population. Alzheimer's and Dementia, 2020, 16, e046072.	0.8	0
24	Use of local genetic ancestry to assess TOMM40 -523â€² and risk for Alzheimer disease. Neurology: Genetics, 2020, 6, e404.	1.9	12
25	Immune and Inflammatory Pathways Implicated by Whole Blood Transcriptomic Analysis in a Diverse Ancestry Alzheimerâ€™s Disease Cohort. Journal of Alzheimer's Disease, 2020, 76, 1047-1060.	2.6	6
26	Identification of Main Genetic Causes Responsible for Non-Syndromic Hearing Loss in a Peruvian Population. Genes, 2019, 10, 581.	2.4	5
27	The Puerto Rico Alzheimer Disease Initiative (PRADI): A Multisource Ascertainment Approach. Frontiers in Genetics, 2019, 10, 538.	2.3	10
28	P1â€144: TRANSCRIPTOMIC ANALYSIS OF WHOLE BLOOD IN AFRICAN AMERICAN AND NONâ€HISPANIC WHITE ALZHEIMER DISEASE CASES AND CONTROLS. Alzheimer's and Dementia, 2018, 14, P331.	0.8	0
29	P2â€106: AFRICAN AMERICAN WHOLE EXOME SEQUENCING SUGGESTS RISK CODING VARIANTS IN IDH1 GENE. Alzheimer's and Dementia, 2018, 14, P709.	0.8	0
30	Rare genetic variation implicated in non-Hispanic white families with Alzheimer disease. Neurology: Genetics, 2018, 4, e286.	1.9	27
31	P1â€154: GENOMEâ€WIDE LINKAGE ANALYSES OF AFRICAN AMERICAN FAMILIES SUPPORTS EVIDENCE OF LINKAGE TO CHROMOSOME 12. Alzheimer's and Dementia, 2018, 14, P336.	0.8	0
32	O3â€06â€06: IDENTIFYING A PROTECTIVE VARIANT THAT LOWERS THE RISK FOR DEVELOPING AD IN APOEâ€E4 CARRIERS. Alzheimer's and Dementia, 2018, 14, P1028.	0.8	0
33	Ancestral origin of ApoE Îµ4 Alzheimer disease risk in Puerto Rican and African American populations. PLoS Genetics, 2018, 14, e1007791.	3.5	117
34	[P2â€“124]: THE PUERTO RICAN ALZHEIMER DISEASE INITIATIVE (PRADI): INITIAL CLINICAL FINDINGS. Alzheimer's and Dementia, 2017, 13, P654.	0.8	0
35	Application of kappa statistics in sequential tests for family-based design. Turkish Journal of Electrical Engineering and Computer Sciences, 2016, 24, 1984-1991.	1.4	0
36	Application of Dempster-Schafer Method in Family-Based Association Studies. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2013, 10, 1071-1075.	3.0	0

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37	Power Analysis of C-TDT for Small Sample Size Genome-Wide Association Studies by the Joint Use of Case-Parent Trios and Pairs. Computational and Mathematical Methods in Medicine, 2013, 2013, 1-7.	1.3	1
38	A novel approach for small sample size family-based association studies: sequential tests. European Journal of Human Genetics, 2011, 19, 915-920.	2.8	3