

Shahzad Ahmad

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

4,980
citations

623188

14
h-index

476904

29
g-index

46
all docs

46
docs citations

46
times ranked

7727
citing authors

#	ARTICLE	IF	CITATIONS
1	Matrix metalloproteinase 10 is linked to the risk of progression to dementia of the Alzheimer's type. <i>Brain</i> , 2022, 145, 2507-2517.	3.7	16
2	Circulating Metabolome and White Matter Hyperintensities in Women and Men. <i>Circulation</i> , 2022, 145, 1040-1052.	1.6	17
3	Challenges at the APOE locus: a robust quality control approach for accurate APOE genotyping. <i>Alzheimer's Research and Therapy</i> , 2022, 14, 22.	3.0	5
4	Meta-analysis of genome-wide association studies identifies ancestry-specific associations underlying circulating total tau levels. <i>Communications Biology</i> , 2022, 5, 336.	2.0	6
5	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	9.4	700
6	Immunity and amyloid beta, total tau and neurofilament light chain: Findings from a community-based cohort study. <i>Alzheimer's and Dementia</i> , 2021, 17, 446-456.	0.4	14
7	Circulating metabolites are associated with brain atrophy and white matter hyperintensities. <i>Alzheimer's and Dementia</i> , 2021, 17, 205-214.	0.4	17
8	Common and Rare Variants Genetic Association Analysis of Circulating Neutrophil Extracellular Traps. <i>Frontiers in Immunology</i> , 2021, 12, 615527.	2.2	8
9	Multomics integrative analysis identifies APOE allele-specific blood biomarkers associated to Alzheimer's disease etiopathogenesis. <i>Aging</i> , 2021, 13, 9277-9329.	1.4	15
10	Genome-wide association study of frontotemporal dementia identifies a C9ORF72 haplotype with a median of 12-G4C2 repeats that predisposes to pathological repeat expansions. <i>Translational Psychiatry</i> , 2021, 11, 451.	2.4	6
11	Whole exome sequencing study identifies novel rare and common Alzheimer's-Associated variants involved in immune response and transcriptional regulation. <i>Molecular Psychiatry</i> , 2020, 25, 1859-1875.	4.1	191
12	Association of lysophosphatidic acids with cerebrospinal fluid biomarkers and progression to Alzheimer's disease. <i>Alzheimer's Research and Therapy</i> , 2020, 12, 124.	3.0	12
13	Genetic correlations and genome-wide associations of cortical structure in general population samples of 22,824 adults. <i>Nature Communications</i> , 2020, 11, 4796.	5.8	61
14	Cross-omics studies of the role of apolipoprotein E in Alzheimer's disease and dementia: Searching common pathways in patients, populations and cellular models. <i>Alzheimer's and Dementia</i> , 2020, 16, e040282.	0.4	0
15	Exome sequencing identifies three novel AD-associated genes. <i>Alzheimer's and Dementia</i> , 2020, 16, e041592.	0.4	6
16	The role of the gut microbiome in cognitive function and Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e043197.	0.4	1
17	SORL1 ϵ -variant carriers in ADESADSP: A higher level of variant pathogenicity associates with earlier age at onset of Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e044492.	0.4	1
18	Clostridium shows a higher abundance in less neurovascular and neurodegenerative changes: A microbiome-wide association study. <i>Alzheimer's and Dementia</i> , 2020, 16, e044743.	0.4	4

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19	CDH6 and HACH protein levels in plasma associate with Alzheimer's disease in APOE ϵ 4 carriers. Scientific Reports, 2020, 10, 8233.	1.6	17
20	Prion protein codon 129 polymorphism in mild cognitive impairment and dementia: the Rotterdam Study. Brain Communications, 2020, 2, fcaa030.	1.5	3
21	Association of Altered Liver Enzymes With Alzheimer Disease Diagnosis, Cognition, Neuroimaging Measures, and Cerebrospinal Fluid Biomarkers. JAMA Network Open, 2019, 2, e197978.	2.8	142
22	Genetic predisposition, modifiable-risk-factor profile and long-term dementia risk in the general population. Nature Medicine, 2019, 25, 1364-1369.	15.2	132
23	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates $\text{A}\beta$, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	9.4	1,962
24	O3-05-01: GENETIC PREDISPOSITION, MODIFIABLE RISK FACTOR PROFILE, AND LONG-TERM DEMENTIA RISK IN THE GENERAL POPULATION. Alzheimer's and Dementia, 2019, 15, P889.	0.4	0
25	Altered bile acid profile associates with cognitive impairment in Alzheimer's disease—An emerging role for gut microbiome. Alzheimer's and Dementia, 2019, 15, 76-92.	0.4	396
26	Altered bile acid profile in mild cognitive impairment and Alzheimer's disease: Relationship to neuroimaging and CSF biomarkers. Alzheimer's and Dementia, 2019, 15, 232-244.	0.4	198
27	Increased High-Density Lipoprotein Levels Associated with Age-Related Macular Degeneration. Ophthalmology, 2019, 126, 393-406.	2.5	88
28	Systematic comparative study of computational methods for T-cell receptor sequencing data analysis. Briefings in Bioinformatics, 2019, 20, 222-234.	3.2	10
29	Disentangling the biological pathways involved in early features of Alzheimer's disease in the Rotterdam Study. , 2018, 14, 848-857.		36
30	F3-02-02: CIRCULATING METABOLITES' ASSOCIATION WITH ALZHEIMER'S DISEASE—ASSOCIATED GENETIC VARIANTS. Alzheimer's and Dementia, 2018, 14, P997.	0.4	0
31	P3-134: CIRCULATING METABOLITES ARE ASSOCIATED WITH WHITE MATTER HYPERINTENSITIES. Alzheimer's and Dementia, 2018, 14, P1119.	0.4	0
32	P1-298: CEREBROSPINAL FLUID AND PLASMA LEVELS OF LYSOPHOSPHATIDIC ACIDS (LPAS) ASSOCIATE WITH CEREBROSPINAL FLUID $\text{A}\beta$ AND τ . Alzheimer's and Dementia, 2018, 14, P403.	0.4	0
33	Variants in TTC25 affect autistic trait in patients with autism spectrum disorder and general population. European Journal of Human Genetics, 2017, 25, 982-987.	1.4	5
34	Rare coding variants in PLCG2, ABI3, and TREM2 implicate microglial-mediated innate immunity in Alzheimer's disease. Nature Genetics, 2017, 49, 1373-1384.	9.4	783
35	[P1-139]: PATHWAY-SPECIFIC GENETIC RISK SCORE ASSOCIATED WITH ALZHEIMER'S DISEASE AND WHITE MATTER LESIONS IN COGNITIVELY NORMAL SUBJECTS. Alzheimer's and Dementia, 2017, 13, P295.	0.4	0
36	O1-09-01: Genomewide Linkage Analysis Identifies Novel Candidate Genes for Alzheimer's Disease. , 2016, 12, P196-P196.		0

#	ARTICLE	IF	CITATIONS
37	01â€09â€04: Identification of Whole Exome Sequencing Variants Associated with Lateâ€Onset Alzheimer's Disease in the Cohorts for Heart and Aging Research in Genomic Epidemiology (Charge) Consortium. Alzheimer's and Dementia, 2016, 12, P197.	0.4	0