

Otto Valladares

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

8,510
citations

430874

18
h-index

501196

28
g-index

44
all docs

44
docs citations

44
times ranked

12596
citing authors

#	ARTICLE	IF	CITATIONS
1	Alzheimer's Disease Variant Portal: A Catalog of Genetic Findings for Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2022, 86, 461-477.	2.6	4
2	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	21.4	700
3	Copy Number Variation Identification on 3,800 Alzheimer's Disease Whole Genome Sequencing Data from the Alzheimer's Disease Sequencing Project. <i>Frontiers in Genetics</i> , 2021, 12, 752390.	2.3	4
4	Genetically regulated expression in late-onset Alzheimer's disease implicates risk genes within known and novel loci. <i>Translational Psychiatry</i> , 2021, 11, 618.	4.8	17
5	NIA genetics of Alzheimer's disease data storage site (NIAGADS): 2021 update.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e052258.	0.8	0
6	Copy number variation (CNV) identification and association study on 3,928 Alzheimer's disease whole genome sequencing data from the Alzheimer's Disease Sequencing Project (ADSP).. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e052721.	0.8	0
7	Introducing the NIAGADS Alzheimer's GenomicsDB API: A toolkit for remote exploration of Alzheimer's disease genetics.. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e053963.	0.8	0
8	Characterization of regulatory roles of genetic signals curated from more than 200 GWA studies in the Alzheimer's Disease Variant Portal (ADVP).. <i>Alzheimer's and Dementia</i> , 2021, 17 Suppl 3, e054255.	0.8	0
9	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.	7.9	191
10	SparkINFERNO: a scalable high-throughput pipeline for inferring molecular mechanisms of non-coding genetic variants. <i>Bioinformatics</i> , 2020, 36, 3879-3881.	4.1	7
11	Alzheimer's disease variant portal (ADVP): Harmonized genetics data and evidence collection for Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2020, 16, e044090.	0.8	0
12	NIA genetics of Alzheimer's disease data storage site (NIAGADS): Update 2020. <i>Alzheimer's and Dementia</i> , 2020, 16, e044284.	0.8	1
13	HIPPIE2: a method for fine-scale identification of physically interacting chromatin regions. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa022.	3.2	2
14	DASHR 2.0: integrated database of human small non-coding RNA genes and mature products. <i>Bioinformatics</i> , 2019, 35, 1033-1039.	4.1	46
15	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates A β , tau, immunity and lipid processing. <i>Nature Genetics</i> , 2019, 51, 414-430.	21.4	1,962
16	VCPA: genomic variant calling pipeline and data management tool for Alzheimer's Disease Sequencing Project. <i>Bioinformatics</i> , 2019, 35, 1768-1770.	4.1	23
17	P4044: THE GCAD CLOUD-BASED WORKFLOW FOR PROCESSING WHOLE EXOME AND WHOLE GENOME DATA FROM THE ALZHEIMER'S DISEASE SEQUENCING PROJECT. <i>Alzheimer's and Dementia</i> , 2018, 14, P1450.	0.8	0
18	P1149: THE ALZHEIMER'S DISEASE SEQUENCING PROJECT (ADSP) DATA UPDATE 2018. <i>Alzheimer's and Dementia</i> , 2018, 14, P333.	0.8	0

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19	P3-130: NIA GENETICS OF ALZHEIMER'S DISEASE DATA STORAGE SITE (NIAGADS): ALZHEIMER'S GENOMICS DATABASE. <i>Alzheimer's and Dementia</i> , 2018, 14, P1117.	0.8	0
20	P1-157: NIA GENETICS OF ALZHEIMER'S DISEASE DATA STORAGE SITE (NIAGADS): UPDATE 2018. <i>Alzheimer's and Dementia</i> , 2018, 14, P337.	0.8	0
21	INFERNO: inferring the molecular mechanisms of noncoding genetic variants. <i>Nucleic Acids Research</i> , 2018, 46, 8740-8753.	14.5	46
22	Rare coding variants in <i>PLCG2</i> , <i>ABI3</i> , and <i>TREM2</i> implicate microglial-mediated innate immunity in Alzheimer's disease. <i>Nature Genetics</i> , 2017, 49, 1373-1384.	21.4	783
23	[O1-03-01]: GENOME-WIDE RARE VARIANT IMPUTATION AND TISSUE-SPECIFIC TRANSCRIPTOMIC ANALYSIS IDENTIFY NOVEL RARE VARIANT CANDIDATE LOCI IN LATE-ONSET ALZHEIMER'S DISEASE: THE ALZHEIMER'S DISEASE GENETICS CONSORTIUM. <i>Alzheimer's and Dementia</i> , 2017, 13, P189.	0.8	4
24	[P3-097]: NIA GENETICS OF ALZHEIMER'S DISEASE DATA STORAGE SITE (NIAGADS): 2017. <i>Alzheimer's and Dementia</i> , 2017, 13, P971.	0.8	0
25	[P3-090]: THE ALZHEIMER'S DISEASE SEQUENCING PROJECT (ADSP) DATA UPDATE 2017. <i>Alzheimer's and Dementia</i> , 2017, 13, P968.	0.8	0
26	P2-097: The Alzheimer's Disease Sequencing Project (ADSP): Data Production, Management, and Availability. <i>Alzheimer's and Dementia</i> , 2016, 12, P648.	0.8	0
27	P3-093: NIA Genetics of Alzheimer's Disease Data Storage Site (NIAGADS): 2016 Update. , 2016, 12, P855-P856.		0
28	O1-03-05: High-Resolution Imputation in Genome-Wide Association Studies of Late-Onset Alzheimer's Disease Identifies Novel Rare Variant Associations. , 2016, 12, P178-P179.		0
29	NIAGADS: The NIA Genetics of Alzheimer's Disease Data Storage Site. <i>Alzheimer's and Dementia</i> , 2016, 12, 1200-1203.	0.8	24
30	DASHR: database of small human noncoding RNAs. <i>Nucleic Acids Research</i> , 2016, 44, D216-D222.	14.5	74
31	Global and local ancestry in African-Americans: Implications for Alzheimer's disease risk. <i>Alzheimer's and Dementia</i> , 2016, 12, 233-243.	0.8	42
32	Rarity of the Alzheimer Disease-Protective <i>APP</i> A673T Variant in the United States. <i>JAMA Neurology</i> , 2015, 72, 209.	9.0	41
33	The role of <i>TREM2</i> R47H as a risk factor for Alzheimer's disease, frontotemporal lobar degeneration, amyotrophic lateral sclerosis, and Parkinson's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 1407-1416.	0.8	152
34	Association of Long Runs of Homozygosity With Alzheimer Disease Among African American Individuals. <i>JAMA Neurology</i> , 2015, 72, 1313.	9.0	39
35	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 658-671.	0.8	173
36	HIPPIE: a high-throughput identification pipeline for promoter interacting enhancer elements. <i>Bioinformatics</i> , 2015, 31, 1290-1292.	4.1	52

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37	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. JAMA Neurology, 2014, 71, 1394.	9.0	166
38	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661.	2.5	155
39	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. Nature Genetics, 2013, 45, 1452-1458.	21.4	3,741