

Otto Valladares

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1463637/publications.pdf>

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	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. <i>Nature Genetics</i> , 2013, 45, 1452-1458.	21.4	3,741
2	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
3	Rare coding variants in <i>PLCC2</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
4	New insights into the genetic etiology of Alzheimer's disease and related dementias. <i>Nature Genetics</i> , 2022, 54, 412-436.	21.4	700
5	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
6	Convergent genetic and expression data implicate immunity in Alzheimer's disease. <i>Alzheimer's and Dementia</i> , 2015, 11, 658-671.	0.8	173
7	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. <i>JAMA Neurology</i> , 2014, 71, 1394.	9.0	166
8	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. <i>PLoS ONE</i> , 2014, 9, e94661.	2.5	155
9	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
10	DASHR: database of small human noncoding RNAs. <i>Nucleic Acids Research</i> , 2016, 44, D216-D222.	14.5	74
11	HIPPIE: a high-throughput identification pipeline for promoter interacting enhancer elements. <i>Bioinformatics</i> , 2015, 31, 1290-1292.	4.1	52
12	INFERNO: inferring the molecular mechanisms of noncoding genetic variants. <i>Nucleic Acids Research</i> , 2018, 46, 8740-8753.	14.5	46
13	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
14	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
15	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
16	Association of Long Runs of Homozygosity With Alzheimer Disease Among African American Individuals. <i>JAMA Neurology</i> , 2015, 72, 1313.	9.0	39
17	NIAGADS: The NIA Genetics of Alzheimer's Disease Data Storage Site. <i>Alzheimer's and Dementia</i> , 2016, 12, 1200-1203.	0.8	24
18	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

#	ARTICLE	IF	CITATIONS
19	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
20	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
21	[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
22	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
23	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
24	HIPPIE2: a method for fine-scale identification of physically interacting chromatin regions. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa022.	3.2	2
25	NIA genetics of Alzheimer's disease data storage site (NIAGADS): Update 2020. <i>Alzheimer's and Dementia</i> , 2020, 16, e044284.	0.8	1
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	[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
30	[P3-090]: THE ALZHEIMER'S DISEASE SEQUENCING PROJECT (ADSP) DATA UPDATE 2017. <i>Alzheimer's and Dementia</i> , 2017, 13, P968.	0.8	0
31	P4-044: 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
32	P1-149: THE ALZHEIMER'S DISEASE SEQUENCING PROJECT (ADSP) DATA UPDATE 2018. <i>Alzheimer's and Dementia</i> , 2018, 14, P333.	0.8	0
33	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
34	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
35	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
36	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

#	ARTICLE	IF	CITATIONS
37	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).. Alzheimer's and Dementia, 2021, 17 Suppl 3, e052721.	0.8	0
38	Introducing the NIAGADS Alzheimer's GenomicsDB API: A toolkit for remote exploration of Alzheimer's disease genetics.. Alzheimer's and Dementia, 2021, 17 Suppl 3, e053963.	0.8	0
39	Characterization of regulatory roles of genetic signals curated from more than 200 CWA studies in the Alzheimer's Disease Variant Portal (ADVP).. Alzheimer's and Dementia, 2021, 17 Suppl 3, e054255.	0.8	0