Gary W Beecham

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5775236/publications.pdf

Version: 2024-02-01

114418 186209 9,002 110 28 63 citations h-index g-index papers 131 131 131 13761 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	Meta-analysis of 74,046 individuals identifies 11 new susceptibility loci for Alzheimer's disease. Nature Genetics, 2013, 45, 1452-1458.	9.4	3,741
2	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates Aβ, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	9.4	1,962
3	Exceptionally low likelihood of Alzheimer's dementia in APOE2 homozygotes from a 5,000-person neuropathological study. Nature Communications, 2020, 11, 667.	5 . 8	246
4	Large-scale GWAS reveals insights into the genetic architecture of same-sex sexual behavior. Science, 2019, 365, .	6.0	245
5	Genome-wide Association Study Implicates a Chromosome 12 Risk Locus for Late-Onset Alzheimer Disease. American Journal of Human Genetics, 2009, 84, 35-43.	2.6	242
6	Sex-Specific Association of Apolipoprotein E With Cerebrospinal Fluid Levels of Tau. JAMA Neurology, 2018, 75, 989.	4.5	223
7	Whole exome sequencing study identifies novel rare and common Alzheimer's-Associated variants involved in immune response and transcriptional regulation. Molecular Psychiatry, 2020, 25, 1859-1875.	4.1	191
8	Convergent genetic and expression data implicate immunity in Alzheimer's disease. Alzheimer's and Dementia, 2015, 11, 658-671.	0.4	173
9	Effects of Multiple Genetic Loci on Age at Onset in Late-Onset Alzheimer Disease. JAMA Neurology, 2014, 71, 1394.	4.5	166
10	Gene-Wide Analysis Detects Two New Susceptibility Genes for Alzheimer's Disease. PLoS ONE, 2014, 9, e94661.	1.1	155
11	The Alzheimer's Disease Sequencing Project: Study design and sample selection. Neurology: Genetics, 2017, 3, e194.	0.9	141
12	Ancestral origin of ApoE $\hat{l}\mu$ 4 Alzheimer disease risk in Puerto Rican and African American populations. PLoS Genetics, 2018, 14, e1007791.	1.5	117
13	Early-Onset Alzheimer's Disease: What Is Missing in Research?. Current Neurology and Neuroscience Reports, 2021, 21, 4.	2.0	88
14	Late-onset vs nonmendelian early-onset Alzheimer disease. Neurology: Genetics, 2020, 6, e512.	0.9	82
15	Repeat expansions in the C9ORF72 gene contribute to Alzheimer's disease in Caucasians. Neurobiology of Aging, 2013, 34, 1519.e5-1519.e12.	1.5	74
16	<i>ABCA7</i> frameshift deletion associated with Alzheimer disease in African Americans. Neurology: Genetics, 2016, 2, e79.	0.9	74
17	<i>C9ORF72</i> Intermediate Repeat Copies Are a Significant Risk Factor for Parkinson Disease. Annals of Human Genetics, 2013, 77, 351-363.	0.3	69
18	Sex differences in the genetic predictors of Alzheimer's pathology. Brain, 2019, 142, 2581-2589.	3.7	65

#	Article	IF	CITATIONS
19	Integrated Whole Transcriptome and DNA Methylation Analysis Identifies Gene Networks Specific to Late-Onset Alzheimer's Disease. Journal of Alzheimer's Disease, 2015, 44, 977-987.	1.2	62
20	Genome-wide brain DNA methylation analysis suggests epigenetic reprogramming in Parkinson disease. Neurology: Genetics, 2019, 5, e342.	0.9	50
21	<i>PARK10</i> is a major locus for sporadic neuropathologically confirmed Parkinson disease. Neurology, 2015, 84, 972-980.	1.5	48
22	Genome-Wide Association Study of Male Sexual Orientation. Scientific Reports, 2017, 7, 16950.	1.6	44
23	Global and local ancestry in Africanâ€Americans: Implications for Alzheimer's disease risk. Alzheimer's and Dementia, 2016, 12, 233-243.	0.4	42
24	Segregation of a rare <i>TTC3</i> variant in an extended family with late-onset Alzheimer disease. Neurology: Genetics, 2016, 2, e41.	0.9	41
25	Early-Onset Alzheimer Disease and Candidate Risk Genes Involved in Endolysosomal Transport. JAMA Neurology, 2017, 74, 1113.	4.5	41
26	A rare missense variant of <i>CASP7</i> is associated with familial lateâ€onset Alzheimer's disease. Alzheimer's and Dementia, 2019, 15, 441-452.	0.4	39
27	Properties of global―and localâ€ancestry adjustments in genetic association tests in admixed populations. Genetic Epidemiology, 2018, 42, 214-229.	0.6	37
28	Variation in <i>SIPA1L2</i> is correlated with phenotype modification in Charcot– Marie– Tooth disease type 1A. Annals of Neurology, 2019, 85, 316-330.	2.8	33
29	Increased <i>APOE</i> \hat{l} µ4 expression is associated with the difference in Alzheimer's disease risk from diverse ancestral backgrounds. Alzheimer's and Dementia, 2021, 17, 1179-1188.	0.4	33
30	Overlap between Parkinson disease and Alzheimer disease in <i>ABCA7</i> functional variants. Neurology: Genetics, 2016, 2, e44.	0.9	31
31	Genomewide Association Studies of <scp><i>LRRK2</i></scp> Modifiers of Parkinson's Disease. Annals of Neurology, 2021, 90, 76-88.	2.8	30
32	Rare genetic variation implicated in non-Hispanic white families with Alzheimer disease. Neurology: Genetics, 2018, 4, e286.	0.9	27
33	Genome-wide pleiotropy analysis of neuropathological traits related to Alzheimer's disease. Alzheimer's Research and Therapy, 2018, 10, 22.	3.0	27
34	Genomic evidence consistent with antagonistic pleiotropy may help explain the evolutionary maintenance of same-sex sexual behaviour in humans. Nature Human Behaviour, 2021, 5, 1251-1258.	6.2	27
35	Genomeâ€wide linkage analyses of nonâ€Hispanic white families identify novel loci for familial lateâ€onset Alzheimer's disease. Alzheimer's and Dementia, 2016, 12, 2-10.	0.4	24
36	Haplotype-specific modulation of a SOX10/CREB response element at the Charcot–Marie–Tooth disease type 4C locus SH3TC2. Human Molecular Genetics, 2014, 23, 5171-5187.	1.4	21

#	Article	IF	CITATIONS
37	PCDH11X variation is not associated with late-onset Alzheimer disease susceptibility. Psychiatric Genetics, 2010, 20, 321-324.	0.6	19
38	RNA editing alterations in a multi-ethnic Alzheimer disease cohort converge on immune and endocytic molecular pathways. Human Molecular Genetics, 2019, 28, 3053-3061.	1.4	19
39	Modifier Gene Candidates in Charcot-Marie-Tooth Disease Type 1A: A Case-Only Genome-Wide Association Study. Journal of Neuromuscular Diseases, 2019, 6, 201-211.	1.1	19
40	The Utility of the National Alzheimer's Coordinating Center's Database for the Rapid Assessment of Evolving Neuropathologic Conditions. Alzheimer Disease and Associated Disorders, 2020, 34, 105-111.	0.6	19
41	A locus at 19q13.31 significantly reduces the ApoE Îμ4 risk for Alzheimer's Disease in African Ancestry. PLoS Genetics, 2022, 18, e1009977.	1.5	19
42	<i>APOE</i> is not Associated with Alzheimer Disease: a Cautionary tale of Genotype Imputation. Annals of Human Genetics, 2010, 74, 189-194.	0.3	13
43	Genetic Characterization and Influence on Inflammatory Bowel Disease Expression in a Diverse Hispanic South Florida Cohort. Clinical and Translational Gastroenterology, 2017, 8, e87.	1.3	13
44	Use of local genetic ancestry to assess <i>TOMM40</i> -523′ and risk for Alzheimer disease. Neurology: Genetics, 2020, 6, e404.	0.9	12
45	DNA variants in <i>CACNA1C</i> modify Parkinson disease risk only when vitamin D level is deficient. Neurology: Genetics, 2016, 2, e72.	0.9	11
46	The executive prominent/memory prominent spectrum in Alzheimer's disease is highly heritable. Neurobiology of Aging, 2016, 41, 115-121.	1.5	11
47	Dissecting the role of Amerindian genetic ancestry and the ApoE $\hat{l}\mu4$ allele on Alzheimer disease in an admixed Peruvian population. Neurobiology of Aging, 2021, 101, 298.e11-298.e15.	1.5	11
48	The Puerto Rico Alzheimer Disease Initiative (PRADI): A Multisource Ascertainment Approach. Frontiers in Genetics, 2019, 10, 538.	1.1	10
49	Linkage analysis of multiplex Caribbean Hispanic families loaded for unexplained earlyâ€onset cases identifies novel Alzheimer's disease loci. Alzheimer's and Dementia: Diagnosis, Assessment and Disease Monitoring, 2018, 10, 554-562.	1.2	8
50	Response to Comment on "Large-scale GWAS reveals insights into the genetic architecture of same-sex sexual behavior― Science, 2021, 371, .	6.0	5
51	Genome studies must account for history—Response. Science, 2019, 366, 1461-1462.	6.0	4
52	Linkage of Alzheimer disease families with Puerto Rican ancestry identifies a chromosome 9 locus. Neurobiology of Aging, 2021, 104, 115.e1-115.e7.	1.5	4
53	Neuropathological lesions and their contribution to dementia and cognitive impairment in a heterogeneous clinical population. Alzheimer's and Dementia, 2022, 18, 2403-2412.	0.4	4
54	Pedigree Selection and Information Content. Current Protocols in Human Genetics, 2018, 97, e56.	3.5	3

#	Article	IF	CITATIONS
55	Genome-Wide Linkage Study Meta-Analysis of Male Sexual Orientation. Archives of Sexual Behavior, 2021, 50, 3371-3375.	1.2	3
56	Genome-Wide Linkage and Association Study of Childhood Gender Nonconformity in Males. Archives of Sexual Behavior, 2021, 50, 3377-3383.	1.2	3
57	O1-03-03: Identification of Novel Candidate Genes for Early-Onset Alzheimer's Disease Through Integrated Whole-Exome Sequencing and Exome Chip Array Association Analysis., 2016, 12, P177-P178.		2
58	Genetic architecture of RNA editing regulation in Alzheimer's disease across diverse ancestral populations. Human Molecular Genetics, 2022, 31, 2876-2886.	1.4	2
59	APOEâ€stratified genomeâ€wide association analysis identifies novel Alzheimer disease candidate risk loci for African Americans. Alzheimer's and Dementia, 2021, 17, e056383.	0.4	2
60	P1â€018: Rare Deleterious And Lossâ€ofâ€Function Variants in <i>OPRL1</i> and <i>GAS2L2</i> Contribute to the Risk of Lateâ€Onset Alzheimer's Disease: Alzheimer's Disease Sequencing Project Caseâ€Control Stud Alzheimer's and Dementia, 2016, 12, P406.	ly0.4	1
61	[O2–08–03]: WHOLEâ€GENOME SEQUENCING IN FAMILIAL LATEâ€ONSET ALZHEIMER's DISEASE IDENTIFIES VARIATION IN AD CANDIDATE GENES. Alzheimer's and Dementia, 2017, 13, P571.	RARE 0.4	1
62	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.	0.4	1
63	F1â€01â€02: Alzheimer's Disease Sequencing Project: Search for Alzheimer's Disease Resilience Genes That May Modify Disease Susceptibility in Specific Apoe Genotype Backgrounds. Alzheimer's and Dementia, 2016, 12, P162.	0.4	О
64	P2â€077: Alzheimer's Disease Sequencing Project: Search for Alzheimer's Disease Resilience Genes That May Modify Disease Susceptibility in Specific ⟨i⟩Apoe⟨/i⟩ Genotype Backgrounds. Alzheimer's and Dementia, 2016, 12, P638.	0.4	0
65	F1-01-03: Rare Deleterious and Loss-of-Function Variants in OPRL1 and GAS2L2 Contribute to the Risk of Late-Onset Alzheimer's Disease: Alzheimer's Disease Sequencing Project Case-Control Study. , 2016, 12, P163-P163.		О
66	O1â€03â€02: <i>ABCA7</i> Frameshift Deletion Associated with Alzheimer's Disease in African Americans. Alzheimer's and Dementia, 2016, 12, P177.	0.4	0
67	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
68	O1-09-02: Whole Exome Sequencing of Late Onset Multiplex Families Identifies Rare Coding Variants in Known and Novel Alzheimer's Disease Genes. , 2016, 12, P196-P197.		0
69	O1â€09â€03: Whole Genome Sequencing in Familial Lateâ€Onset Alzheimer's Disease Identifies Variations in TTC3 and FSIP2. Alzheimer's and Dementia, 2016, 12, P197.	0.4	О
70	P1â€122: Multivariate Phenotypes Association Study of Neuropathological Features of Alzheimer's Disease and Related Dementias. Alzheimer's and Dementia, 2016, 12, P450.	0.4	0
71	[P3–094]: RESOURCE OF MULTIPLEX AFRICAN AMERICAN FAMILIES FOR WHOLEâ€GENOME SEQUENCING. Alzheimer's and Dementia, 2017, 13, P970.	0.4	О
72	[P2–075]: INFLUENCE OF COMMUNITY ENGAGED FAMILY CONNECTOR IN RECRUITING AND ASCERTAINING AFRICAN AMERICANS' FAMILY MEMBERS FOR GENOMIC RESEARCH. Alzheimer's and Dementia, 2017, 13, P63	3 4 :4	0

#	Article	IF	CITATIONS
73	[P2–102]: THE PUERTO RICO ALZHEIMER DISEASE INITIATIVE (PRADI): A MULTISOURCE ASCERTAINMENT APPROACH. Alzheimer's and Dementia, 2017, 13, P646.	0.4	0
74	[P2–105]: COLLECTION OF MULTIPLEX FAMILIES WITH UNEXPLAINED EARLYâ€ONSET ALZHEIMER's DISEASE FOR GENOMIC RESEARCH. Alzheimer's and Dementia, 2017, 13, P647.	OR 0.4	0
75	[P2–124]: THE PUERTO RICAN ALZHEIMER DISEASE INITIATIVE (PRADI): INITIAL CLINICAL FINDINGS. Alzheimer's and Dementia, 2017, 13, P654.	0.4	0
76	[O2–08–02]: SEX‧PECIFIC ANALYSIS OF THE ADSP CASE ONTROL WHOLEâ€EXOME SEQUENCING DATAlzheimer's and Dementia, 2017, 13, P571.	TASET. 0.4	0
77	P3â€034: CONTINUOUS COMMUNITY ENGAGEMENT IMPROVES RECRUITMENT OF OLDER AFRICAN AMERICANS FOR GENETIC STUDIES IN ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2018, 14, P1077.	0.4	O
78	P1â€156: GENEâ€BASED ANALYSES IN WHOLE GENOME SEQUENCING OF FAMILIAL LATEâ€ONSET ALZHEIMER'S DISEASE. Alzheimer's and Dementia, 2018, 14, P336.	0.4	0
79	P2â€106: AFRICAN AMERICAN WHOLE EXOME SEQUENCING SUGGESTS RISK CODING VARIANTS IN IDH1 GENE. Alzheimer's and Dementia, 2018, 14, P709.	0.4	O
80	P1â€139: THE CONTRIBUTION OF SEXâ€SPECIFIC ASSOCIATIONS IN GENETIC STUDIES OF ALZHEIMER'S DISEASE PATHOLOGY. Alzheimer's and Dementia, 2018, 14, P327.	0.4	0
81	P1â€154: GENOMEâ€WIDE LINKAGE ANALYSES OF AFRICAN AMERICAN FAMILIES SUPPORTS EVIDENCE OF LINKA TO CHROMOSOME 12. Alzheimer's and Dementia, 2018, 14, P336.	GE 0.4	O
82	P2â€121: APOLIPOPROTEIN E AND PHENOTYPIC FEATURES IN HISPANICS. Alzheimer's and Dementia, 2018, 14, P715.	0.4	0
83	P2â€108: WHOLEâ€GENOME SEQUENCING IN NONâ€HISPANIC WHITE FAMILIES IMPLICATES RARE VARIATION IN LATEâ€ONSET ALZHEIMER'S DISEASE RISK. Alzheimer's and Dementia, 2018, 14, P710.	0.4	O
84	O2â€01â€05: MULTIâ€ETHNIC ALZHEIMER'S DISEASE RELATED CHANGES OF RNA EDITING AFFECT IMMUNE REGULATION, ENDOCYTOSIS, AND AMYLOID PRECURSOR PROTEIN CATABOLISM. Alzheimer's and Dementia, 2018, 14, P609.	0.4	0
85	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.	0.4	0
86	Recruitment strategies for the genetics of Alzheimer disease in the Puerto Rican population. Alzheimer's and Dementia, 2020, 16, e043468.	0.4	0
87	Exploring the role of Amerindian genetic ancestry and ApoEε4 gene on Alzheimer disease in the Peruvian population. Alzheimer's and Dementia, 2020, 16, e045012.	0.4	0
88	A multiancestry analysis of Alzheimer's disease coexpressed gene networks identifies a common immune signaling pathway regulated by granulocyteâ€colony stimulating factor (Gâ€CSF). Alzheimer's and Dementia, 2020, 16, e045361.	0.4	0
89	Increased <i>APOEâ€e4</i> 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.	0.4	O
90	Assessing whole genome sequencing variation for Alzheimer's disease in 4707 individuals from the Alzheimer's Disease Sequencing Project (ADSP). Alzheimer's and Dementia, 2020, 16, e045548.	0.4	0

#	Article	IF	CITATIONS
91	Transcriptomic characterization of a Puerto Rican Alzheimer disease cohort implicates convergent immuneâ€related pathways. Alzheimer's and Dementia, 2020, 16, e045890.	0.4	O
92	Southern European genetic ancestry shows reduced APOE E4 risk for Alzheimer disease in Caribbean Hispanic population. Alzheimer's and Dementia, 2020, 16, e045951.	0.4	0
93	Multimodal genomeâ€wide metaâ€analysis of brain amyloidosis reveals heterogeneity across CSF, PET, and pathological amyloid measures. Alzheimer's and Dementia, 2020, 16, e046009.	0.4	O
94	The effect of global ancestry and diabetes on the 3MS score in older Puerto Ricans. Alzheimer's and Dementia, 2020, 16, e046051.	0.4	0
95	Mapping Alzheimer disease–associated regions in the African American population. Alzheimer's and Dementia, 2020, 16, e046072.	0.4	0
96	Education and its effect on risk and age at onset in Alzheimer disease (AD) in African Americans. Alzheimer's and Dementia, 2020, 16, e046078.	0.4	0
97	Recruiting African American males in Alzheimer's disease education and genetics research. Alzheimer's and Dementia, 2020, 16, e046178.	0.4	0
98	[P2â€"113]: THE RELEVANCE OF APOE4 TO ALZHEIMER's DISEASE IN THE PRESENCE OF LOCAL ANCESTRY DIFFERENCES. Alzheimer's and Dementia, 2017, 13, P650.	0.4	0
99	Assessment of ADâ \in related plasma biomarkers in diverse ancestral populations. Alzheimer's and Dementia, 2021, 17, .	0.4	0
100	Does higher educational attainment influence functional capabilities among African Americans with Alzheimer $\hat{a} \in \mathbb{R}^{M}$ s disease?. Alzheimer's and Dementia, 2021, 17, .	0.4	0
101	Transgenic <i>APOEÎμ4/4</i> overexpression induces reactivity in astrocytes with a European <i>APOEÎμ4/4</i> local ancestry, but not in astrocytes with an African <i>APOEÎμ4/4</i> local ancestry. Alzheimer's and Dementia, 2021, 17, e056397.	0.4	0
102	Outreach and recruitment of African Americans for Alzheimer's disease studies during the COVIDâ€19 pandemic. Alzheimer's and Dementia, 2021, 17, .	0.4	0
103	Neuropathologic lesions and comorbidity in Alzheimer disease and related dementias in a heterogeneous clinical population. Alzheimer's and Dementia, 2021, 17, e056249.	0.4	0
104	Heritability analyses show partial genetic overlap between (nonâ€Mendelian) early and late onset Alzheimer disease due to an intriguing APOE effect. Alzheimer's and Dementia, 2021, 17, e056143.	0.4	0
105	Clinical profile of an AlzheimerÂ's disease cohort in the Peruvian population. Alzheimer's and Dementia, 2021, 17, .	0.4	0
106	African locus reduces the effect of ApoE $\hat{l}\mu4$ allele in Alzheimer's disease Alzheimer's and Dementia, 2021, 17 Suppl 3, e056210.	0.4	0
107	Expression quantitative trait loci (eQTL) analysis in a diverse Alzheimer disease cohort reveals ancestry-specific regulatory architectures Alzheimer's and Dementia, 2021, 17 Suppl 3, e056211.	0.4	0
108	Linkage analysis identifies novel loci in early-onset Alzheimer disease in non-Hispanic white families Alzheimer's and Dementia, 2021, 17 Suppl 3, e056427.	0.4	0

#	Article	IF	CITATIONS
109	Admixture mapping identifies novel regions influencing Alzheimer disease in African Americans Alzheimer's and Dementia, 2021, 17 Suppl 3, e056443.	0.4	0
110	A large-scale, whole genome sequencing study of unexplained early-onset Alzheimer disease Alzheimer's and Dementia, 2021, 17 Suppl 3, e056664.	0.4	0