# Anthony J Griswold

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/8103101/publications.pdf

Version: 2024-02-01

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|---|---|---|---|
| 59 papers | 2,276 citations | 430442 18 h-index | 276539 41 g-index |
| 67 all docs | 67 docs citations | 67 times ranked | 4876 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Preksha DhyÄna meditation induces alterations at the transcriptome level in novice and healthy college students. Saudi Journal of Biological Sciences, 2022, 29, 2299-2305. | 1.8 | 3 |
| 2 | Ischemic-Trained Monocytes Improve Arteriogenesis in a Mouse Model of Hindlimb Ischemia. Arteriosclerosis, Thrombosis, and Vascular Biology, 2022, 42, 175-188. | 1.1 | 5 |
| 3 | Genetic architecture of RNA editing regulation in Alzheimerâ€™s disease across diverse ancestral populations. Human Molecular Genetics, 2022, 31, 2876-2886. | 1.4 | 2 |
| 4 | Whole-genome sequencing reveals complex genomic features underlying anti-CD19 CAR T-cell treatment failures in lymphoma. Blood, 2022, 140, 491-503. | 0.6 | 32 |
| 5 | A locus at 19q13.31 significantly reduces the ApoE Îµ4 risk for Alzheimerâ€™s Disease in African Ancestry. PLoS Genetics, 2022, 18, e1009977. | 1.5 | 19 |
| 6 | Activity of the growth hormoneâ€releasing hormone antagonist MIA602 and its underlying mechanisms of action in sarcoidosisâ€like granuloma. Clinical and Translational Immunology, 2021, 10, e1310. | 1.7 | 8 |
| 7 | 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. | 0.4 | 33 |
| 8 | Obesity-Dependent Adipokine Chemerin Suppresses Fatty Acid Oxidation to Confer Ferroptosis Resistance. Cancer Discovery, 2021, 11, 2072-2093. | 7.7 | 43 |
| 9 | Extended survival versus accelerated rejection of nonhuman primate islet allografts: Effect of mesenchymal stem cell source and timing. American Journal of Transplantation, 2021, 21, 3524-3537. | 2.6 | 6 |
| 10 | Identification of a genetic variant underlying familial cases of recurrent benign paroxysmal positional vertigo. PLoS ONE, 2021, 16, e0251386. | 1.1 | 2 |
| 11 | Establishment and Characterization of a Novel Human Ocular Adnexal Sebaceous Carcinoma Cell Line. Translational Vision Science and Technology, 2021, 10, 34. | 1.1 | 4 |
| 12 | 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 |
| 13 | Single-Cell RNA-Sequencing Identifies Infrapatellar Fat Pad Macrophage Polarization in Acute Synovitis/Fat Pad Fibrosis and Cell Therapy. Bioengineering, 2021, 8, 166. | 1.6 | 7 |
| 14 | Using Genomic Techniques in Sports and Exercise Science: Current Status and Future Opportunities. Current Sports Medicine Reports, 2021, 20, 617-623. | 0.5 | 2 |
| 15 | Genomic Drivers of Large B-Cell Lymphoma Resistance to CD19 CAR-T Therapy. Blood, 2021, 138, 42-42. | 0.6 | 10 |
| 16 | 675â€…Genomic drivers of large B-cell lymphoma resistance to CD19 CAR-T therapy. , 2021, 9, A703-A703. | | 0 |
| 17 | Assessment of ADâ€related plasma biomarkers in diverse ancestral populations. Alzheimer's and Dementia, 2021, 17, . | 0.4 | 0 |
| 18 | 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. | 0.4 | 0 |

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|---|---|---|---|
| 19 | 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. | 0.4 | 0 |
| 20 | 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 |
| 21 | Infrapatellar Fat Pad/Synovium Complex in Early-Stage Knee Osteoarthritis: Potential New Target and Source of Therapeutic Mesenchymal Stem/Stromal Cells. Frontiers in Bioengineering and Biotechnology, 2020, 8, 860. | 2.0 | 49 |
| 22 | A multiancestry analysis of Alzheimerâ€™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 |
| 23 | 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. | 0.4 | 0 |
| 24 | African and European local ancestry surrounding Apolipoprotein E has a differential biological effect upon acute amyloid beta exposure in iPSCâ€differentiated astrocytes. Alzheimer's and Dementia, 2020, 16, e045424. | 0.4 | 0 |
| 25 | Transcriptomic characterization of a Puerto Rican Alzheimer disease cohort implicates convergent immuneâ€related pathways. Alzheimer's and Dementia, 2020, 16, e045890. | 0.4 | 0 |
| 26 | Development of a massively parallel reporter assay to identify functional regulatory variants in the PICALM Alzheimer disease associated locus. Alzheimer's and Dementia, 2020, 16, e045908. | 0.4 | 0 |
| 27 | Identification of differential regulation of European versus African local ancestry haplotypes surrounding ApoEÎµ4. Alzheimer's and Dementia, 2020, 16, e046016. | 0.4 | 0 |
| 28 | Structural characterization of rare missense variants within known neurodegenerative disease proteins. Alzheimer's and Dementia, 2020, 16, e046405. | 0.4 | 0 |
| 29 | Single-cell resolution analysis of the human pancreatic ductal progenitor cell niche. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 10876-10887. | 3.3 | 109 |
| 30 | Use of local genetic ancestry to assess TOMM40 -523â€² and risk for Alzheimer disease. Neurology: Genetics, 2020, 6, e404. | 0.9 | 12 |
| 31 | 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. | 1.2 | 6 |
| 32 | Three Brothers With Autism Carry a Stopâ€Gain Mutation in the HPAâ€Axis Gene NR3C2. Autism Research, 2020, 13, 523-531. | 2.1 | 7 |
| 33 | Growth Hormone-Releasing Hormone Receptor Antagonist Modulates Lung Inflammation and Fibrosis due to Bleomycin. Lung, 2019, 197, 541-549. | 1.4 | 29 |
| 34 | coMethDMR: accurate identification of co-methylated and differentially methylated regions in epigenome-wide association studies with continuous phenotypes. Nucleic Acids Research, 2019, 47, e98-e98. | 6.5 | 28 |
| 35 | AUTHOR REPLY. Urology, 2019, 129, 67. | 0.5 | 0 |
| 36 | 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 |

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|---|---|---|---|
| 37 | Transcriptomics of Human Arteriovenous Fistula Failure: Genes Associated With Nonmaturation. American Journal of Kidney Diseases, 2019, 74, 73-81. | 2.1 | 28 |
| 38 | Whole-Exome Sequencing Identifies Novel Heterozygous Mutation in RAF1 in Family With Neonatal Testicular Torsion. Urology, 2019, 129, 60-67. | 0.5 | 4 |
| 39 | Whole Exome Sequencing Identifies a Rare Nonsense Mutation in FAM47C as a Possible Cause of Severe Oligospermia in Brothers With Varicocele. Urology, 2019, 129, 71-73. | 0.5 | 10 |
| 40 | Mycobacterium abscessusâ€"Bronchial Epithelial Cells Cross-Talk Through Type I Interferon Signaling. Frontiers in Immunology, 2019, 10, 2888. | 2.2 | 17 |
| 41 | Identification of rare noncoding sequence variants in gamma-aminobutyric acid A receptor, alpha 4 subunit in autism spectrum disorder. Neurogenetics, 2018, 19, 17-26. | 0.7 | 5 |
| 42 | Transcriptomic analysis of synovial extracellular RNA following knee trauma: A pilot study. Journal of Orthopaedic Research, 2018, 36, 1659-1665. | 1.2 | 11 |
| 43 | P2â€106: AFRICAN AMERICAN WHOLE EXOME SEQUENCING SUGGESTS RISK CODING VARIANTS IN IDH1 GENE. Alzheimer's and Dementia, 2018, 14, P709. | 0.4 | 0 |
| 44 | 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 |
| 45 | Whole Exome Sequencing of a Consanguineous Turkish Family Identifies a Mutation in GTF2H3 in Brothers With Spermatogenic Failure. Urology, 2018, 120, 86-89. | 0.5 | 9 |
| 46 | Global Gene Expression Change Induced by Major Thoracoabdominal Surgery. Annals of Surgery, 2017, 266, 981-987. | 2.1 | 8 |
| 47 | Osteoarthritic Extracellular RNA Biomarkers in Synovial Fluid. Medicine and Science in Sports and Exercise, 2017, 49, 87-88. | 0.2 | 0 |
| 48 | Targeted massively parallel sequencing of autism spectrum disorder-associated genes in a case control cohort reveals rare loss-of-function risk variants. Molecular Autism, 2015, 6, 43. | 2.6 | 57 |
| 49 | Two knockdown models of the autism genes SYNGAP1 and SHANK3 in zebrafish produce similar behavioral phenotypes associated with embryonic disruptions of brain morphogenesis. Human Molecular Genetics, 2015, 24, 4006-4023. | 1.4 | 67 |
| 50 | Global Gene Expression Change Induced by Major Thoracoabdominal Surgery. Journal of the American College of Surgeons, 2015, 221, S70. | 0.2 | 0 |
| 51 | Detection of canonical A-to-G editing events at 3â€² UTRs and microRNA target sites in human lungs using next-generation sequencing. Oncotarget, 2015, 6, 35726-35736. | 0.8 | 15 |
| 52 | Convergence of Genes and Cellular Pathways Dysregulated in Autism Spectrum Disorders. American Journal of Human Genetics, 2014, 94, 677-694. | 2.6 | 819 |
| 53 | Evaluation of copy number variations reveals novel candidate genes in autism spectrum disorder-associated pathways. Human Molecular Genetics, 2012, 21, 3513-3523. | 1.4 | 158 |
| 54 | A de novo 1.5â€‰Mb microdeletion on chromosome 14q23.2â€23.3 in a patient with autism and spherocytosis. Autism Research, 2011, 4, 221-227. | 2.1 | 31 |

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|---|---|---|---|
| 55 | A noise-reduction GWAS analysis implicates altered regulation of neurite outgrowth and guidance in autism. Molecular Autism, 2011, 2, 1. | 2.6 | 191 |
| 56 | A Genomeâ€wide Association Study of Autism Reveals a Common Novel Risk Locus at 5p14.1. Annals of Human Genetics, 2009, 73, 263-273. | 0.3 | 207 |
| 57 | Overexpression of frataxin in the mitochondria increases resistance to oxidative stress and extends lifespan in Drosophila. FEBS Letters, 2008, 582, 715-719. | 1.3 | 66 |
| 58 | Sir2 mediates apoptosis through JNK-dependent pathways in Drosophila. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 8673-8678. | 3.3 | 65 |
| 59 | Individual histone deacetylases in Drosophila modulate transcription of distinct genes. Genomics, 2005, 86, 606-617. | 1.3 | 67 |