# Jian-Hua Chen

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/7336905/publications.pdf

Version: 2024-02-01

| 83 | 3,542 | 257101 24 | 155451 55 |
|---|---|---|---|
| papers | citations | h-index | g-index |

| 85 | 85 | 85 | 7300 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Patients with mental health disorders in the COVID-19 epidemic. Lancet Psychiatry,the, 2020, 7, e21. | 3.7 | 1,053 |
| 2 | Genome-wide association analysis identifies 30 new susceptibility loci for schizophrenia. Nature Genetics, 2017, 49, 1576-1583. | 9.4 | 395 |
| 3 | Recurrent gain-of-function USP8 mutations in Cushing's disease. Cell Research, 2015, 25, 306-317. | 5.7 | 263 |
| 4 | Rethinking online mental health services in China during the COVID-19 epidemic. Asian Journal of Psychiatry, 2020, 50, 102015. | 0.9 | 151 |
| 5 | The genome-wide mutational landscape of pituitary adenomas. Cell Research, 2016, 26, 1255-1259. | 5.7 | 137 |
| 6 | Identification of recurrent USP48 and BRAF mutations in Cushingâ€™s disease. Nature Communications, 2018, 9, 3171. | 5.8 | 106 |
| 7 | Genome-wide association analysis identifies three new risk loci for gout arthritis in Han Chinese. Nature Communications, 2015, 6, 7041. | 5.8 | 88 |
| 8 | SHEsisPlus, a toolset for genetic studies on polyploid species. Scientific Reports, 2016, 6, 24095. | 1.6 | 77 |
| 9 | DNA origami-based shape IDs for single-molecule nanomechanical genotyping. Nature Communications, 2017, 8, 14738. | 5.8 | 73 |
| 10 | Workplace factors associated with mental health of healthcare workers during the COVID-19 pandemic: an international cross-sectional study. BMC Health Services Research, 2021, 21, 262. | 0.9 | 71 |
| 11 | Germline Mutations in CDH23, Encoding Cadherin-Related 23, Are Associated with Both Familial and Sporadic Pituitary Adenomas. American Journal of Human Genetics, 2017, 100, 817-823. | 2.6 | 57 |
| 12 | Susceptibility loci for metabolic syndrome and metabolic components identified in Han Chinese: a multiâ€stage genomeâ€wide association study. Journal of Cellular and Molecular Medicine, 2017, 21, 1106-1116. | 1.6 | 56 |
| 13 | Genome-wide Analysis of the Role of Copy Number Variation in Schizophrenia Risk in Chinese. Biological Psychiatry, 2016, 80, 331-337. | 0.7 | 55 |
| 14 | Wholeâ€exome sequencing of oral mucosal melanoma reveals mutational profile and therapeutic targets. Journal of Pathology, 2018, 244, 358-366. | 2.1 | 52 |
| 15 | Genome-wide association study identifies two risk loci for tuberculosis in Han Chinese. Nature Communications, 2018, 9, 4072. | 5.8 | 51 |
| 16 | Genome-Wide Association Study of Bladder Cancer in a Chinese Cohort Reveals a New Susceptibility Locus at 5q12.3. Cancer Research, 2016, 76, 3277-3284. | 0.4 | 46 |
| 17 | Common variants at 10p12.31, 10q21.1 and 13q12.13 are associated with sporadic pituitary adenoma. Nature Genetics, 2015, 47, 793-797. | 9.4 | 43 |
| 18 | Body Mass Index and Polycystic Ovary Syndrome: A 2-Sample Bidirectional Mendelian Randomization Study. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 1778-1784. | 1.8 | 39 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Significant association of GRM7 and GRM8 genes with schizophrenia and major depressive disorder in the Han Chinese population. European Neuropsychopharmacology, 2016, 26, 136-146. | 0.3 | 35 |
| 20 | The GSK3B gene confers risk for both major depressive disorder and schizophrenia in the Han Chinese population. Journal of Affective Disorders, 2015, 185, 149-155. | 2.0 | 34 |
| 21 | Glucose and Insulin-Related Traits, Type 2 Diabetes and Risk of Schizophrenia: A Mendelian Randomization Study. EBioMedicine, 2018, 34, 182-188. | 2.7 | 34 |
| 22 | Genome-wide association study of cervical cancer suggests a role forARRDC3gene in human papillomavirus infection. Human Molecular Genetics, 2019, 28, 341-348. | 1.4 | 33 |
| 23 | Acupuncture improves the symptoms, intestinal microbiota, and inflammation of patients with mild to moderate Crohn's disease: A randomized controlled trial. EClinicalMedicine, 2022, 45, 101300. | 3.2 | 30 |
| 24 | Loci with genome-wide associations with schizophrenia in the Han Chinese population. British Journal of Psychiatry, 2015, 207, 490-494. | 1.7 | 29 |
| 25 | Alcohol Consumption in China Before and During COVID-19: Preliminary Results From an Online Retrospective Survey. Frontiers in Psychiatry, 2020, 11, 597826. | 1.3 | 29 |
| 26 | Genetic association between NRG1 and schizophrenia, major depressive disorder, bipolar disorder in Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 468-478. | 1.1 | 26 |
| 27 | Association between SREBF2 gene polymorphisms and metabolic syndrome in clozapine-treated patients with schizophrenia. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2015, 56, 136-141. | 2.5 | 19 |
| 28 | Network pharmacology-based identification for therapeutic mechanism of Ling-Gui-Zhu-Gan decoction in the metabolic syndrome induced by antipsychotic drugs. Computers in Biology and Medicine, 2019, 110, 1-7. | 3.9 | 19 |
| 29 | Enhancement of CTLs induced by DCs loaded with ubiquitinated hepatitis B virus core antigen. World Journal of Gastroenterology, 2012, 18, 1319. | 1.4 | 18 |
| 30 | ITIH family genes confer risk to schizophrenia and major depressive disorder in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2014, 51, 34-38. | 2.5 | 17 |
| 31 | Bioinformatics Analysis of the Effects of Tobacco Smoke on Gene Expression. PLoS ONE, 2015, 10, e0143377. | 1.1 | 17 |
| 32 | The NVL gene confers risk for both major depressive disorder and schizophrenia in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2015, 62, 7-13. | 2.5 | 17 |
| 33 | The comorbidity of mental and physical disorders with self-reported chronic back or neck pain: Results from the China Mental Health Survey. Journal of Affective Disorders, 2020, 260, 334-341. | 2.0 | 17 |
| 34 | Both HLA class I and II regions identified as genome-wide significant susceptibility loci for adult-onset Still's disease in Chinese individuals. Annals of the Rheumatic Diseases, 2020, 79, 161-163. | 0.5 | 17 |
| 35 | Endothelial Nitric Oxide Synthase (eNOS) T-786C, 4a4b, and G894T Polymorphisms and Male Infertility: Study for Idiopathic Asthenozoospermia and Meta-Analysis1. Biology of Reproduction, 2015, 92, 38. | 1.2 | 16 |
| 36 | The protective effects of Mogroside V and its metabolite 11-oxo-mogrol of intestinal microbiota against MK801-induced neuronal damages. Psychopharmacology, 2020, 237, 1011-1026. | 1.5 | 16 |

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|---|---|---|---|
| 37 | Polymorphisms in GCKR, SLC17A1 and SLC22A12 were associated with phenotype gout in Han Chinese males: a caseâ€“control study. BMC Medical Genetics, 2015, 16, 66. | 2.1 | 15 |
| 38 | Replication of Gout/Urate Concentrations GWAS Susceptibility Loci Associated with Gout in a Han Chinese Population. Scientific Reports, 2017, 7, 4094. | 1.6 | 15 |
| 39 | Mitigating mental health consequences during the <scp>COVID</scp>â€19 outbreak: Lessons from China. Psychiatry and Clinical Neurosciences, 2020, 74, 407-408. | 1.0 | 13 |
| 40 | Ubiquitin Conjugation of Hepatitis B Virus Core Antigen DNA Vaccine Leads to Enhanced Cell-Mediated Immune Response in BALB/c Mice. Hepatitis Monthly, 2011, 11, 620-8. | 0.1 | 12 |
| 41 | Association betweenSCAPandSREBF1gene polymorphisms and metabolic syndrome in schizophrenia patients treated with atypical antipsychotics. World Journal of Biological Psychiatry, 2016, 17, 467-474. | 1.3 | 12 |
| 42 | Genetic risk between the CACNA1I gene and schizophrenia in Chinese Uygur population. Hereditas, 2018, 155, 5. | 0.5 | 12 |
| 43 | Identification of a novel susceptibility locus at 16q23.1 associated with childhood acute lymphoblastic leukemia in Han Chinese. Human Molecular Genetics, 2016, 25, ddw112. | 1.4 | 10 |
| 44 | A new risk locus in the ZEB2 gene for schizophrenia in the Han Chinese population. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2016, 66, 97-103. | 2.5 | 10 |
| 45 | Polymorphisms in NRGN are associated with schizophrenia, major depressive disorder and bipolar disorder in the Han Chinese population. Journal of Affective Disorders, 2016, 194, 180-187. | 2.0 | 10 |
| 46 | Genomeâ€wide twoâ€locus interaction analysis identifies multiple epistatic SNP pairs that confer risk of prostate cancer: A crossâ€population study. International Journal of Cancer, 2017, 140, 2075-2084. | 2.3 | 10 |
| 47 | SNX29, a new susceptibility gene shared with major mental disorders in Han Chinese population. World Journal of Biological Psychiatry, 2021, 22, 526-534. | 1.3 | 10 |
| 48 | Common variants in ZMIZ1 and near NGF confer risk for primary dysmenorrhoea. Nature Communications, 2017, 8, 14900. | 5.8 | 9 |
| 49 | Association between the variability of theABCA13gene and the risk of major depressive disorder and schizophrenia in the Han Chinese population. World Journal of Biological Psychiatry, 2017, 18, 550-556. | 1.3 | 9 |
| 50 | PPARG Polymorphisms Are Associated with Unexplained Mild Vision Loss in Patients with Type 2 Diabetes Mellitus. Journal of Ophthalmology, 2019, 2019, 1-7. | 0.6 | 9 |
| 51 | Network pharmacology-based exploration of therapeutic mechanism of Liu-Yu-Tang in atypical antipsychotic drug-induced metabolic syndrome. Computers in Biology and Medicine, 2021, 134, 104452. | 3.9 | 9 |
| 52 | The Effect of Yijinjing on the Cognitive Function of Patients With Chronic Schizophrenia. Frontiers in Psychiatry, 2021, 12, 739364. | 1.3 | 9 |
| 53 | ACTN3 is associated with childrenâ€™s physical fitness in Han Chinese. Molecular Genetics and Genomics, 2019, 294, 47-56. | 1.0 | 8 |
| 54 | The amino acid variants in HLA II molecules explain the major association with adult-onset Still's disease in the Han Chinese population. Journal of Autoimmunity, 2021, 116, 102562. | 3.0 | 8 |

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|---|---|---|---|
| 55 | Diacylglycerol kinase Îº (<scp>DGKK</scp>) variants and hypospadias in <scp>H</scp>an <scp>C</scp>hinese: association and metaâ€analysis. BJU International, 2015, 116, 634-640. | 1.3 | 7 |
| 56 | SLC39A8 is a risk factor for schizophrenia in Uygur Chinese: a case-control study. BMC Psychiatry, 2019, 19, 293. | 1.1 | 7 |
| 57 | Fineâ€mapping of ZDHHC2 identifies risk variants for schizophrenia in the Han Chinese population. Molecular Genetics & Genomic Medicine, 2020, 8, e1190. | 0.6 | 7 |
| 58 | Association study of NDST3 gene for schizophrenia, bipolar disorder, major depressive disorder in the Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2018, 177, 3-9. | 1.1 | 6 |
| 59 | Identification of rare and common variants in BNIP3L: a schizophrenia susceptibility gene. Human Genomics, 2020, 14, 16. | 1.4 | 6 |
| 60 | Genome-wide analysis of DNA methylation identifies S100A13 as an epigenetic biomarker in individuals with chronic (â‰¥ 30â€‰years) type 2 diabetes without diabetic retinopathy. Clinical Epigenetics, 2020, 12, 77. | 1.8 | 6 |
| 61 | Analysis of association between common variants of uncoupling proteins genes and diabetic retinopathy in a Chinese population. BMC Medical Genetics, 2020, 21, 25. | 2.1 | 6 |
| 62 | Cigarette smoking and schizophrenia: Mendelian randomisation study. British Journal of Psychiatry, 2021, 218, 98-103. | 1.7 | 6 |
| 63 | Identification of SHANK2 Pathogenic Variants in a Chinese Uygur Population with Schizophrenia. Journal of Molecular Neuroscience, 2021, 71, 1-8. | 1.1 | 6 |
| 64 | Common variants in FAN1, located in 15q13.3, confer risk for schizophrenia and bipolar disorder in Han Chinese. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2020, 103, 109973. | 2.5 | 5 |
| 65 | Genetic risk of clozapine-induced leukopenia and neutropenia: a genome-wide association study. Translational Psychiatry, 2021, 11, 343. | 2.4 | 5 |
| 66 | Long-term effect of moxibustion on irritable bowel syndrome with diarrhea: a randomized clinical trial. Therapeutic Advances in Gastroenterology, 2022, 15, 175628482210751. | 1.4 | 5 |
| 67 | The Relationship between Alcohol Consumption and Gout: A Mendelian Randomization Study. Genes, 2022, 13, 557. | 1.0 | 5 |
| 68 | Analysis of association between common variants in theSLCO6A1gene with schizophrenia, bipolar disorder and major depressive disorder in the Han Chinese population. World Journal of Biological Psychiatry, 2016, 17, 140-146. | 1.3 | 4 |
| 69 | Association of fat mass and obesity-associated and retinitis pigmentosa guanosine triphosphatase (GTPase) regulator-interacting protein-1 like polymorphisms with body mass index in Chinese women. Endocrine Journal, 2018, 65, 783-791. | 0.7 | 4 |
| 70 | Rare and common variants analysis of the EMB gene in patients with schizophrenia. BMC Psychiatry, 2020, 20, 135. | 1.1 | 4 |
| 71 | Patient health questionnaire-15 (PHQ-15) to distinguish bipolar II disorder from major depressive disorder. Psychiatry Research, 2020, 290, 113026. | 1.7 | 4 |
| 72 | Common variants in QPCT gene confer risk of schizophrenia in the Han Chinese population. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2016, 171, 237-242. | 1.1 | 3 |

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|---|---|---|---|
| 73 | Role played by the SP4 gene in schizophrenia and major depressive disorder in the Han Chinese population. British Journal of Psychiatry, 2016, 208, 441-445. | 1.7 | 3 |
| 74 | Survival prediction of anxious emotion in advanced cancer patients receiving palliative care. Psycho-Oncology, 2017, 26, 1463-1469. | 1.0 | 3 |
| 75 | Polymorphism of the PPARD Gene and Dynamic Balance Performance in Han Chinese Children. Hereditas, 2019, 156, 15. | 0.5 | 3 |
| 76 | VariFAST: a variant filter by automated scoring based on tagged-signatures. BMC Bioinformatics, 2019, 20, 713. | 1.2 | 3 |
| 77 | Common variants in SATB2 are associated with schizophrenia in Uygur Chinese population. Psychiatric Genetics, 2019, 29, 120-126. | 0.6 | 2 |
| 78 | Acute psychological impact of coronavirus disease 2019 outbreak among psychiatric professionals in China: a multicentre, cross-sectional, web-based study. BMJ Open, 2021, 11, e047828. | 0.8 | 2 |
| 79 | Brief mindfulness-based intervention of 'STOP (Stop, Take a Breath, Observe, Proceed) touching your face': a study protocol of a randomised controlled trial. BMJ Open, 2020, 10, e041364. | 0.8 | 2 |
| 80 | RNA-seq co-expression network analysis reveals anxiolytic behavior of mice with Efnb2 knockout in parvalbumin+â€‰neurons. Molecular Brain, 2021, 14, 118. | 1.3 | 1 |
| 81 | Editorial: Neurobiological Biomarkers for Developing Novel Treatments of Substance and Non-substance Addiction. Frontiers in Psychiatry, 2021, 12, 811032. | 1.3 | 1 |
| 82 | Association analysis of potentially functional variants within 8p12 with schizophrenia in the Han Chinese population. World Journal of Biological Psychiatry, 2021, 22, 27-33. | 1.3 | 0 |
| 83 | Rare variations in the SHANK3 gene confers susceptibility to schizophrenia in Uygur Chinese population. Schizophrenia Research, 2021, 228, 597-599. | 1.1 | 0 |