

Xiaodong Zheng

List of Publications by Year in descending order

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Version: 2024-02-01

57
papers

1,625
citations

516710

16
h-index

315739

38
g-index

59
all docs

59
docs citations

59
times ranked

3439
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Case Report: Challenges in the Diagnosis of a Case of Mal de Meleda and a Therapeutic Attempt of Ixekizumab and Adalimumab. <i>Frontiers in Medicine</i> , 2022, 9, 821301. | 2.6 | 2 |
| 2 | Gene interaction analysis of psoriasis in Chinese Han population. <i>Molecular Genetics & Genomic Medicine</i> , 2022, , e1858. | 1.2 | 1 |
| 3 | Genome-wide association study of 7661 Chinese Han individuals and fine-mapping major histocompatibility complex identifies HLA-DRB1 as associated with IgA vasculitis. <i>Journal of Clinical Laboratory Analysis</i> , 2022, 36, e24457. | 2.1 | 5 |
| 4 | Bach2 overexpression represses Th9 cell differentiation by suppressing IRF4 expression in systemic lupus erythematosus. <i>FEBS Open Bio</i> , 2021, 11, 395-403. | 2.3 | 12 |
| 5 | Meta-analysis of 208370 East Asians identifies 113 susceptibility loci for systemic lupus erythematosus. <i>Annals of the Rheumatic Diseases</i> , 2021, 80, 632-640. | 0.9 | 103 |
| 6 | <i>CYP2S1</i> might regulate proliferation and immune response of keratinocyte in psoriasis. <i>Epigenetics</i> , 2021, 16, 618-628. | 2.7 | 7 |
| 7 | A high stem to leaf ratio reduced rainfall use efficiency under altered rainfall patterns in a semi-arid grassland in northeast China. <i>Plant Biology</i> , 2021, 23, 760-769. | 3.8 | 1 |
| 8 | Assay for Transposase-Accessible Chromatin Using Sequencing Analysis Reveals a Widespread Increase in Chromatin Accessibility in Psoriasis. <i>Journal of Investigative Dermatology</i> , 2021, 141, 1745-1753. | 0.7 | 8 |
| 9 | An in-depth analysis reveals two new genetic variants on 22q11.2 associated with vitiligo in the Chinese Han population. <i>Molecular Biology Reports</i> , 2021, 48, 5955-5964. | 2.3 | 1 |
| 10 | A Genome-wide association study identified <i>HLA-C</i> associated with the effectiveness of methotrexate for psoriasis treatment. <i>Journal of the European Academy of Dermatology and Venereology</i> , 2021, 35, e898-e900. | 2.4 | 5 |
| 11 | AURKA facilitates the psoriasis-related inflammation by impeding autophagy-mediated AIM2 inflammasome suppression. <i>Immunology Letters</i> , 2021, 240, 98-105. | 2.5 | 8 |
| 12 | Exome-Wide Rare Loss-of-Function Variant Enrichment Study of 21,347 Han Chinese Individuals Identifies Four Susceptibility Genes for Psoriasis. <i>Journal of Investigative Dermatology</i> , 2020, 140, 799-805.e1. | 0.7 | 6 |
| 13 | Identification of a Novel Mutation in SASH1 Gene in a Chinese Family With Dyschromatosis Universalis Hereditaria and Genotype-Phenotype Correlation Analysis. <i>Frontiers in Genetics</i> , 2020, 11, 841. | 2.3 | 9 |
| 14 | Loss-of-function variants in FSIP1 identified by targeted sequencing are associated with one particular subtype of mucosal melanoma. <i>Gene</i> , 2020, 759, 144964. | 2.2 | 3 |
| 15 | Changes in the hepatitis B surface antibody in childhood acute lymphocytic leukaemia survivors after treatment with the CCLG-ALL 2008 protocol. <i>Clinical and Experimental Immunology</i> , 2020, 203, 80-86. | 2.6 | 2 |
| 16 | Effects of laying breeder hens dietary β -carotene, curcumin, allicin, and sodium butyrate supplementation on the jejunal microbiota and immune response of their offspring chicks. <i>Poultry Science</i> , 2020, 99, 3807-3816. | 3.4 | 13 |
| 17 | Quantitative proteomics analysis of young and elderly skin with DIA mass spectrometry reveals new skin aging-related proteins. <i>Aging</i> , 2020, 12, 13529-13554. | 3.1 | 21 |
| 18 | Association Study Reveals One Susceptibility Locus with Vitiligo in the Chinese Han Population. <i>Genetic Testing and Molecular Biomarkers</i> , 2019, 23, 791-796. | 0.7 | 11 |

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|----|---|------|-----------|
| 19 | DNA methylation profile of psoriatic skins from different body locations. <i>Epigenomics</i> , 2019, 11, 1613-1625. | 2.1 | 3 |
| 20 | Physical origin of the expansion of polymer coils in a binary solvent in the vicinity of its demixing critical point. <i>Molecular Physics</i> , 2019, 117, 3806-3811. | 1.7 | 3 |
| 21 | HLA-C*01:02 and HLA-A*02:07 Confer Risk Specific for Psoriatic Patients in Southern China. <i>Journal of Investigative Dermatology</i> , 2019, 139, 2045-2048.e4. | 0.7 | 8 |
| 22 | A KRT16 mutation in the first Chinese pedigree with Pachyonychia congenita and review of the literatures. <i>Journal of Cosmetic Dermatology</i> , 2019, 18, 1930-1934. | 1.6 | 4 |
| 23 | Association analysis of the major histocompatibility complex region in psoriasis vulgaris. <i>British Journal of Dermatology</i> , 2019, 180, 1553-1554. | 1.5 | 0 |
| 24 | Fine mapping and subphenotyping implicates <i>ADRA1B</i> gene variants in psoriasis susceptibility in a Chinese population. <i>Epigenomics</i> , 2019, 11, 455-467. | 2.1 | 10 |
| 25 | <sc>HLA</sc>â€<sc>DQ</sc>Î²1 amino acid position 87 and <sc>DQB</sc>1*0301 are associated with Chinese Han <sc>SLE</sc>. <i>Molecular Genetics & Genomic Medicine</i> , 2018, 6, 541-546. | 1.2 | 5 |
| 26 | DNA methylation-based subclassification of psoriasis in the Chinese Han population. <i>Frontiers of Medicine</i> , 2018, 12, 717-725. | 3.4 | 8 |
| 27 | Association of the novel susceptible locus rs9266150 with clinical features of psoriasis vulgaris in the Chinese Han population. <i>Experimental Dermatology</i> , 2018, 27, 748-753. | 2.9 | 1 |
| 28 | Exome-wide association study identifies four novel loci for systemic lupus erythematosus in Han Chinese population. <i>Annals of the Rheumatic Diseases</i> , 2018, 77, 417-417. | 0.9 | 50 |
| 29 | A catalog of potential putative functional variants in psoriasis genome-wide association regions. <i>PLoS ONE</i> , 2018, 13, e0196635. | 2.5 | 5 |
| 30 | A novel insertion mutation of <i>CDSN</i> responsible for hypotrichosis simplex of scalp in a Chinese family. <i>Clinical and Experimental Dermatology</i> , 2018, 43, 722-723. | 1.3 | 3 |
| 31 | Genome-wide analyses of non-syndromic cleft lip with palate identify 14 novel loci and genetic heterogeneity. <i>Nature Communications</i> , 2017, 8, 14364. | 12.8 | 207 |
| 32 | A genome-wide association study identifies six novel risk loci for primary biliary cholangitis. <i>Nature Communications</i> , 2017, 8, 14828. | 12.8 | 102 |
| 33 | Correlation analysis of the <i>HLAâ€PB1*05:01</i> and <i>BTNL2</i> genes within the histocompatibility complex region with a clinical phenotype of psoriasis vulgaris in the Chinese Han population. <i>Journal of Gene Medicine</i> , 2017, 19, e2961. | 2.8 | 4 |
| 34 | Integrative analyses reveal biological pathways and key genes in psoriasis. <i>British Journal of Dermatology</i> , 2017, 177, 1349-1357. | 1.5 | 30 |
| 35 | Integration of expression quantitative trait loci and pleiotropy identifies a novel psoriasis susceptibility gene, <i>PTPN1</i> . <i>Journal of Gene Medicine</i> , 2017, 19, e2939. | 2.8 | 5 |
| 36 | Genotype combination contributes to psoriasis: An exhaustive algorithm perspective. <i>PLoS ONE</i> , 2017, 12, e0186067. | 2.5 | 5 |

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|----|--|------|-----------|
| 37 | Common susceptibility variants are shared between schizophrenia and psoriasis in the Han Chinese population. <i>Journal of Psychiatry and Neuroscience</i> , 2016, 41, 413-421. | 2.4 | 19 |
| 38 | Genetic Susceptibility to Vitiligo: GWAS Approaches for Identifying Vitiligo Susceptibility Genes and Loci. <i>Frontiers in Genetics</i> , 2016, 7, 3. | 2.3 | 69 |
| 39 | Epigenome-wide association data implicates DNA methylation-mediated genetic risk in psoriasis. <i>Clinical Epigenetics</i> , 2016, 8, 131. | 4.1 | 31 |
| 40 | Deep sequencing of the MHC region in the Chinese population contributes to studies of complex disease. <i>Nature Genetics</i> , 2016, 48, 740-746. | 21.4 | 188 |
| 41 | Epigenome-Wide Association Analysis Identified Nine Skin DNA Methylation Loci for Psoriasis. <i>Journal of Investigative Dermatology</i> , 2016, 136, 779-787. | 0.7 | 75 |
| 42 | A Genetic Variant rs1020760 at <i>NFKB1</i> is Associated with Clinical Features of Psoriasis Vulgaris in a Han Chinese Population. <i>Annals of Human Genetics</i> , 2016, 80, 197-202. | 0.8 | 5 |
| 43 | Downregulated expression of <i>LBH</i> mRNA in peripheral blood mononuclear cells from patients with systemic lupus erythematosus. <i>Journal of Dermatology</i> , 2016, 43, 99-102. | 1.2 | 14 |
| 44 | Identification of cell types, tissues and pathways affected by risk loci in psoriasis. <i>Molecular Genetics and Genomics</i> , 2016, 291, 1005-1012. | 2.1 | 3 |
| 45 | Appropriate Osmotic Balance Duration for Different Volumes of Ovarian Tissue in Vitrification Solution: a Study of Ovary Tissue Vitrification and Transplantation in Sheep. <i>Cryo-Letters</i> , 2016, 37, 365-378. | 0.3 | 0 |
| 46 | Discovery of a novel genetic susceptibility locus on X chromosome for systemic lupus erythematosus. <i>Arthritis Research and Therapy</i> , 2015, 17, 349. | 3.5 | 26 |
| 47 | Rs4948496 within <i>ARID5B</i> gene is associated with clinical features of systemic lupus erythematosus in the Chinese Han population. <i>Journal of Dermatology</i> , 2015, 42, 608-612. | 1.2 | 2 |
| 48 | A genetic coding variant rs72474224 in <i>GJB2</i> is associated with clinical features of psoriasis vulgaris in a Chinese Han population. <i>Tissue Antigens</i> , 2015, 86, 134-138. | 1.0 | 8 |
| 49 | Whole-exome SNP array identifies 15 new susceptibility loci for psoriasis. <i>Nature Communications</i> , 2015, 6, 6793. | 12.8 | 118 |
| 50 | Association analysis of allergic sensitization susceptibility loci with atopic dermatitis in Chinese population. <i>Journal of Dermatological Science</i> , 2015, 80, 217-220. | 1.9 | 1 |
| 51 | Rapid diagnosis of Fusarium root rot in soybean caused by <i>Fusarium equiseti</i> or <i>Fusarium graminearum</i> using loop-mediated isothermal amplification (LAMP) assays. <i>Australasian Plant Pathology</i> , 2015, 44, 437-443. | 1.0 | 23 |
| 52 | A new magnetorheological elastomer isolator in shear-compression mixed mode. <i>Journal of Intelligent Material Systems and Structures</i> , 2015, 26, 1290-1300. | 2.5 | 79 |
| 53 | A large-scale screen for coding variants predisposing to psoriasis. <i>Nature Genetics</i> , 2014, 46, 45-50. | 21.4 | 183 |
| 54 | Sequencing-based approach identified three new susceptibility loci for psoriasis. <i>Nature Communications</i> , 2014, 5, 4331. | 12.8 | 67 |

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|----|--|-----|-----------|
| 55 | Exome sequencing identifies SLC17A9 pathogenic gene in two Chinese pedigrees with disseminated superficial actinic porokeratosis. <i>Journal of Medical Genetics</i> , 2014, 51, 699-704. | 3.2 | 32 |
| 56 | Molecular Characterization and Tissue-specific Expression of a Novel FKBP38 Gene in the Cashmere Goat (<i>Capra hircus</i>). <i>Asian-Australasian Journal of Animal Sciences</i> , 2012, 25, 758-763. | 2.4 | 2 |
| 57 | First Report of Elm Yellow's Phytoplasma Infecting Clover in China. <i>Plant Disease</i> , 2009, 93, 321-321. | 1.4 | 9 |