## Donglei Hu

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2562631/publications.pdf

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87 papers	5,447 citations	33 h-index	91884 69 g-index
90	90	90	11226
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Meta-analysis of genome-wide association studies of asthma in ethnically diverse North American populations. Nature Genetics, 2011, 43, 887-892.	21.4	736
2	DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium Meta-analysis. American Journal of Human Genetics, 2016, 98, 680-696.	6.2	717
3	Multi-ancestry genome-wide association study of 21,000 cases and 95,000 controls identifies new risk loci for atopic dermatitis. Nature Genetics, 2015, 47, 1449-1456.	21.4	529
4	Genetic Ancestry in Lung-Function Predictions. New England Journal of Medicine, 2010, 363, 321-330.	27.0	230
5	Sparse PCA corrects for cell type heterogeneity in epigenome-wide association studies. Nature Methods, 2016, 13, 443-445.	19.0	205
6	Differential methylation between ethnic sub-groups reflects the effect of genetic ancestry and environmental exposures. ELife, 2017, 6, .	6.0	153
7	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 2062-2074.	2.9	147
8	Breast cancer risk prediction using a clinical risk model and polygenic risk score. Breast Cancer Research and Treatment, 2016, 159, 513-525.	2.5	129
9	Genome-wide association study of breast cancer in Latinas identifies novel protective variants on 6q25. Nature Communications, 2014, 5, 5260.	12.8	123
10	Genome-wide methylation data mirror ancestry information. Epigenetics and Chromatin, 2017, 10, 1.	3.9	120
11	Genetic ancestry influences asthma susceptibility and lung function among Latinos. Journal of Allergy and Clinical Immunology, 2015, 135, 228-235.	2.9	113
12	Genome-wide association study and admixture mapping identify different asthma-associated loci in Latinos: The Genes-environments & Samp; Admixture in Latino Americans study. Journal of Allergy and Clinical Immunology, 2014, 134, 295-305.	2.9	106
13	Whole-Genome Sequencing of Pharmacogenetic Drug Response in Racially Diverse Children with Asthma. American Journal of Respiratory and Critical Care Medicine, 2018, 197, 1552-1564.	5.6	102
14	Germline genetic contribution to the immune landscape of cancer. Immunity, 2021, 54, 367-386.e8.	14.3	95
15	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. Nature Communications, 2018, 9, 2976.	12.8	85
16	The TAM family receptor tyrosine kinase TYRO3 is a negative regulator of type 2 immunity. Science, 2016, 352, 99-103.	12.6	67
17	Serum Insulinâ€Like Growth Factorâ€1 Binding Proteins 1 and 2 and Mortality in Older Adults: The Health, Aging, and Body Composition Study. Journal of the American Geriatrics Society, 2009, 57, 1213-1218.	2.6	63
18	Functional genomics of CDHR3 confirms its role in HRV-C infection and childhood asthma exacerbations. Journal of Allergy and Clinical Immunology, 2019, 144, 962-971.	2.9	63

#	Article	IF	CITATIONS
19	Cis-eQTL-based trans-ethnic meta-analysis reveals novel genes associated with breast cancer risk. PLoS Genetics, 2017, 13, e1006690.	3.5	61
20	Admixture mapping identifies a locus on 6q25 associated with breast cancer risk in US Latinas. Human Molecular Genetics, 2012, 21, 1907-1917.	2.9	60
21	Epigenetic and transcriptional determinants of the human breast. Nature Communications, 2015, 6, 6351.	12.8	56
22	A genome-wide association and admixture mapping study of bronchodilator drug response in African Americans with asthma. Pharmacogenomics Journal, 2019, 19, 249-259.	2.0	54
23	A Polygenic Risk Score for Breast Cancer in US Latinas and Latin American Women. Journal of the National Cancer Institute, 2020, 112, 590-598.	6.3	53
24	Genome-wide association study and admixture mapping reveal new loci associated with total IgE levels in Latinos. Journal of Allergy and Clinical Immunology, 2015, 135, 1502-1510.	2.9	52
25	A meta-analysis of genome-wide association studies of asthma in PuertoÂRicans. European Respiratory Journal, 2017, 49, 1601505.	6.7	51
26	Dysregulated invertebrate tropomyosin–dectin-1 interaction confers susceptibility to allergic diseases. Science Immunology, 2018, 3, .	11.9	51
27	Genome-wide association study of lung function phenotypes in a founder population. Journal of Allergy and Clinical Immunology, 2014, 133, 248-255.e10.	2.9	50
28	Genomeâ€wide association study of inhaled corticosteroid response in admixed children with asthma. Clinical and Experimental Allergy, 2019, 49, 789-798.	2.9	50
29	COMT ValMet polymorphism is associated with post-traumatic stress disorder and functional outcome following mild traumatic brain injury. Journal of Clinical Neuroscience, 2017, 35, 109-116.	1.5	43
30	Pathogenic and likely pathogenic variants in <i>PALB2</i> , <i>CHEK2</i> , and other known breast cancer susceptibility genes among 1054 <i>BRCA</i> â€negative Hispanics with breast cancer. Cancer, 2019, 125, 2829-2836.	4.1	43
31	Whole-Genome Sequencing of Individuals from a Founder Population Identifies Candidate Genes for Asthma. PLoS ONE, 2014, 9, e104396.	2.5	42
32	On the cross-population generalizability of gene expression prediction models. PLoS Genetics, 2020, 16, e1008927.	3.5	41
33	Immunotherapy-Mediated Thyroid Dysfunction: Genetic Risk and Impact on Outcomes with PD-1 Blockade in Non–Small Cell Lung Cancer. Clinical Cancer Research, 2021, 27, 5131-5140.	7.0	40
34	Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients. Nature Communications, 2015, 6, 7539.	12.8	38
35	A genome-wide association study identifies only two ancestry specific variants associated with spontaneous preterm birth. Scientific Reports, 2018, 8, 226.	3.3	37
36	Ancestry-Dependent Enrichment of Deleterious Homozygotes in Runs of Homozygosity. American Journal of Human Genetics, 2019, 105, 747-762.	6.2	36

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37	Genetic Investigation Into the Differential Risk of Atrial Fibrillation Among Black and White Individuals. JAMA Cardiology, $2016,1,442.$	6.1	35
38	COMT Val 158 Met polymorphism is associated with nonverbal cognition following mild traumatic brain injury. Neurogenetics, 2016, 17, 31-41.	1.4	33
39	Correcting for cell-type heterogeneity in DNA methylation: a comprehensive evaluation. Nature Methods, 2017, 14, 218-219.	19.0	33
40	An admixture mapping meta-analysis implicates genetic variation at 18q21 with asthma susceptibility in Latinos. Journal of Allergy and Clinical Immunology, 2019, 143, 957-969.	2.9	33
41	Meta-analysis of GWA studies provides new insights on the genetic architecture of skin pigmentation in recently admixed populations. BMC Genetics, 2019, 20, 59.	2.7	32
42	Identification of novel common breast cancer risk variants at the 6q25 locusÂamong Latinas. Breast Cancer Research, 2019, 21, 3.	5.0	32
43	Identification of a novel locus associated with skin colour in African-admixed populations. Scientific Reports, 2017, 7, 44548.	3.3	31
44	The Effects of Migration and Assortative Mating on Admixture Linkage Disequilibrium. Genetics, 2017, 205, 375-383.	2.9	31
45	Secondhand smoke exposure and asthma outcomes among African-American and Latino children with asthma. Thorax, 2018, 73, 1041-1048.	5.6	30
46	Human Epidermal Growth Factor Receptor 2–Positive Breast Cancer Is Associated with Indigenous American Ancestry in Latin American Women. Cancer Research, 2020, 80, 1893-1901.	0.9	29
47	Association of imputed prostate cancer transcriptome with disease risk reveals novel mechanisms. Nature Communications, 2019, 10, 3107.	12.8	28
48	Multiple breast cancer risk variants are associated with differential transcript isoform expression in tumors. Human Molecular Genetics, 2015, 24, 7421-7431.	2.9	24
49	DRD2 C957T polymorphism is associated with improved 6-month verbal learning following traumatic brain injury. Neurogenetics, 2017, 18, 29-38.	1.4	24
50	Ancestry and genetic associations with bronchopulmonary dysplasia in preterm infants. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2018, 315, L858-L869.	2.9	24
51	Ambient air pollution, asthma drug response, and telomere length in African American youth. Journal of Allergy and Clinical Immunology, 2019, 144, 839-845.e10.	2.9	24
52	In utero tobacco smoke exposure, DNA methylation, and asthma in Latino children. Environmental Epidemiology, 2019, 3, e048.	3.0	24
53	Cross-ancestry GWAS meta-analysis identifies six breast cancer loci in African and European ancestry women. Nature Communications, 2021, 12, 4198.	12.8	24
54	A genome-wide association study of asthma hospitalizations in adults. Journal of Allergy and Clinical Immunology, 2021, 147, 933-940.	2.9	23

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55	Joint relative risks for estrogen receptor-positive breast cancer from a clinical model, polygenic risk score, and sex hormones. Breast Cancer Research and Treatment, 2017, 166, 603-612.	2.5	22
56	Telomere Length and the Risk of Atrial Fibrillation. Circulation: Arrhythmia and Electrophysiology, 2014, 7, 1026-1032.	4.8	21
57	Genetic Determinants of Telomere Length in African American Youth. Scientific Reports, 2018, 8, 13265.	3.3	20
58	European genetic ancestry associated with risk of childhood ependymoma. Neuro-Oncology, 2020, 22, 1637-1646.	1.2	16
59	Racial/ethnic differences in eligibility for asthma biologics among pediatric populations. Journal of Allergy and Clinical Immunology, 2021, 148, 1324-1331.e12.	2.9	16
60	Asthma and its relationship to mitochondrial copy number: Results from the Asthma Translational Genomics Collaborative (ATGC) of the Trans-Omics for Precision Medicine (TOPMed) program. PLoS ONE, 2020, 15, e0242364.	2.5	16
61	Intrahepatic Cholestasis of Pregnancy (ICP) in U.S. Latinas and Chileans: Clinical features, Ancestry Analysis, and Admixture Mapping. PLoS ONE, 2015, 10, e0131211.	2.5	15
62	A genome-wide association study of severe asthma exacerbations in Latino children and adolescents. European Respiratory Journal, 2021, 57, 2002693.	6.7	15
63	Confounding in Genetic Association Studies and Its Solutions. Methods in Molecular Biology, 2008, 448, 31-39.	0.9	13
64	Racial Differences in Left Atrial Size: Results from the Coronary Artery Risk Development in Young Adults (CARDIA) Study. PLoS ONE, 2016, 11, e0151559.	2.5	13
65	Adapt-Mix: learning local genetic correlation structure improves summary statistics-based analyses. Bioinformatics, 2015, 31, i181-i189.	4.1	12
66	Epigenome-wide association study of lung function in Latino children and youth with asthma. Clinical Epigenetics, 2022, 14, 9.	4.1	12
67	Lung Function in African American Children with Asthma Is Associated with Novel Regulatory Variants of the KIT Ligand <i>KITLG/SCF</i> and Gene-By-Air-Pollution Interaction. Genetics, 2020, 215, 869-886.	2.9	11
68	Whole-Genome Sequencing Identifies Novel Functional Loci Associated with Lung Function in Puerto Rican Youth. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 962-972.	5.6	11
69	NLRP1 variant M1184V decreases inflammasome activation in the context of DPP9 inhibition and asthma severity. Journal of Allergy and Clinical Immunology, 2021, 147, 2134-2145.e20.	2.9	11
70	A genome-wide study of DNA methylation in white blood cells and asthma in Latino children and youth. Epigenetics, 2021, 16, 577-585.	2.7	10
71	Lymph node–resident dendritic cells drive T <sub>H</sub> 2 cell development involving MARCH1. Science Immunology, 2021, 6, eabh0707.	11.9	10
72	Pharmacogenetic studies of long-acting beta agonist and inhaled corticosteroid responsiveness in randomised controlled trials of individuals of African descent with asthma. The Lancet Child and Adolescent Health, 2021, 5, 862-872.	5.6	10

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73	Breastfeeding associated with higher lung function in African American youths with asthma. Journal of Asthma, 2017, 54, 856-865.	1.7	7
74	Role of Sex on the Genetic Susceptibility to Childhood Asthma in Latinos and African Americans. Journal of Personalized Medicine, 2021, 11, 1140.	2.5	7
75	Whole Genome Sequencing Identifies CRISPLD2 as a Lung Function Gene in Children With Asthma. Chest, 2019, 156, 1068-1079.	0.8	5
76	Association of a PAI-1 Gene Polymorphism and Early Life Infections with Asthma Risk, Exacerbations, and Reduced Lung Function. PLoS ONE, 2016, 11, e0157848.	2.5	5
77	Integrative genomic analysis in African American children with asthma finds three novel loci associated with lung function. Genetic Epidemiology, 2021, 45, 190-208.	1.3	4
78	Genetic loci determining total immunoglobulin E levels from birth through adulthood. Allergy: European Journal of Allergy and Clinical Immunology, 2019, 74, 621-625.	5.7	2
79	Native American Ancestry and Air Pollution Interact to Impact Bronchodilator Response in Puerto Rican Children with Asthma. Ethnicity and Disease, 2021, 31, 77-88.	2.3	2
80	Association between Ancestry-Specific 6q25 Variants and Breast Cancer Subtypes in Peruvian Women. Cancer Epidemiology Biomarkers and Prevention, 0, , OF1-OF8.	2.5	2
81	Disentangling the impact of alcohol use and hepatitis C on insulin action in Latino individuals. Alcoholism: Clinical and Experimental Research, 2022, 46, 87-99.	2.4	0
82	On the cross-population generalizability of gene expression prediction models., 2020, 16, e1008927.		0
83	On the cross-population generalizability of gene expression prediction models., 2020, 16, e1008927.		O
84	On the cross-population generalizability of gene expression prediction models., 2020, 16, e1008927.		0
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86	On the cross-population generalizability of gene expression prediction models., 2020, 16, e1008927.		0
87	On the cross-population generalizability of gene expression prediction models., 2020, 16, e1008927.		0