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List of Publications by Year in descending order

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Version: 2024-02-01

24
papers

9,265
citations

471509

17
h-index

642732

23
g-index

27
all docs

27
docs citations

27
times ranked

18030
citing authors

#	ARTICLE	IF	CITATIONS
1	Early Changes in Immune Cell Count, Metabolism, and Function Following Sleeve Gastrectomy: A Prospective Human Study. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2022, 107, e619-e630.	3.6	11
2	Asthma Susceptibility Gene <i>ORMDL3</i> Promotes Autophagy in Human Bronchial Epithelium. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2022, 66, 661-670.	2.9	6
3	Expression of SMARCD1 interacts with age in association with asthma control on inhaled corticosteroid therapy. <i>Respiratory Research</i> , 2020, 21, 31.	3.6	6
4	Pathway analysis of a genome-wide gene by air pollution interaction study in asthmatic children. <i>Journal of Exposure Science and Environmental Epidemiology</i> , 2019, 29, 539-547.	3.9	13
5	Meta-analysis of genome-wide association studies for body fat distribution in 694,649 individuals of European ancestry. <i>Human Molecular Genetics</i> , 2019, 28, 166-174.	2.9	752
6	The eMERGE genotype set of 83,717 subjects imputed to ~40 million variants genome wide and association with the herpes zoster medical record phenotype. <i>Genetic Epidemiology</i> , 2019, 43, 63-81.	1.3	63
7	Role of local CpG DNA methylation in mediating the 17q21 asthma susceptibility gasdermin B (<i>GSDMB</i>)/ <i>ORMDL</i> sphingolipid biosynthesis regulator 3 (<i>ORMDL3</i>) expression quantitative trait locus. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 2282-2286.e6.	2.9	20
8	TREM-1 Response Signatures Common to Expression Profiles of Both Asthma Affection and Asthma Control. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2018, 198, 401-404.	5.6	0
9	Gene Coexpression Networks in Whole Blood Implicate Multiple Interrelated Molecular Pathways in Obesity in People with Asthma. <i>Obesity</i> , 2018, 26, 1938-1948.	3.0	11
10	An Integrative Transcriptomic and Metabolomic Study of Lung Function in Children With Asthma. <i>Chest</i> , 2018, 154, 335-348.	0.8	52
11	Novel eosinophilic gene expression networks associated with IgE in two distinct asthma populations. <i>Clinical and Experimental Allergy</i> , 2018, 48, 1654-1664.	2.9	22
12	Limited statistical evidence for shared genetic effects of eQTLs and autoimmune-disease-associated loci in three major immune-cell types. <i>Nature Genetics</i> , 2017, 49, 600-605.	21.4	205
13	Across-cohort QC analyses of GWAS summary statistics from complex traits. <i>European Journal of Human Genetics</i> , 2017, 25, 137-146.	2.8	18
14	Gene Expression Profiling in Blood Provides Reproducible Molecular Insights into Asthma Control. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2017, 195, 179-188.	5.6	49
15	Genetics and Genomics of Longitudinal Lung Function Patterns in Individuals with Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2016, 194, 1465-1474.	5.6	20
16	Stress and Bronchodilator Response in Children with Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2015, 192, 47-56.	5.6	99
17	New genetic loci link adipose and insulin biology to body fat distribution. <i>Nature</i> , 2015, 518, 187-196.	27.8	1,328
18	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015, 518, 197-206.	27.8	3,823

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19	Genome-wide interaction studies reveal sex-specific asthma risk alleles. <i>Human Molecular Genetics</i> , 2014, 23, 5251-5259.	2.9	70
20	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014, 46, 1173-1186.	21.4	1,818
21	Quality control and conduct of genome-wide association meta-analyses. <i>Nature Protocols</i> , 2014, 9, 1192-1212.	12.0	398
22	Sex-stratified Genome-wide Association Studies Including 270,000 Individuals Show Sexual Dimorphism in Genetic Loci for Anthropometric Traits. <i>PLoS Genetics</i> , 2013, 9, e1003500.	3.5	371
23	Population-specific coding variant underlies genome-wide association with adiponectin level. <i>Human Molecular Genetics</i> , 2012, 21, 463-471.	2.9	37
24	Genome-wide Association Study of Anthropometric Traits and Evidence of Interactions With Age and Study Year in Filipino Women. <i>Obesity</i> , 2011, 19, 1019-1027.	3.0	72