Stephanie J. London

List of Publications by Year
in descending order

367 papers	57,102 citations	2671 95 h-index	1187 228 g-index
385 all docs	385 docs citations	385 times ranked	73260 citing authors

#	Article	IF	Citations
1	Maternal haemoglobin levels in pregnancy and child DNA methylation: a study in the pregnancy and childhood epigenetics consortium. Epigenetics, 2022, 17, 19-31.	1.3	3
2	Association of clonal hematopoiesis with chronic obstructive pulmonary disease. Blood, 2022, 139, 357-368.	0.6	106
3	Association between ambient particulate matter exposure and semen quality in fertile men. Environmental Health, 2022, 21, 16.	1.7	23
4	Meta-analysis of epigenome-wide associations between DNA methylation at birth and childhood cognitive skills. Molecular Psychiatry, 2022, 27, 2126-2135.	4.1	13
5	Meta-analysis of epigenome-wide association studies in newborns and children show widespread sex differences in blood DNA methylation. Mutation Research - Reviews in Mutation Research, 2022, 789, 108415.	2.4	24
6	Associations between the Maternal Exposome and Metabolome during Pregnancy. Environmental Health Perspectives, 2022, 130, 37003.	2.8	15
7	Polygenic transcriptome risk scores for COPD and lung function improve cross-ethnic portability of prediction in the NHLBI TOPMed program. American Journal of Human Genetics, 2022, 109, 857-870.	2.6	7
8	Lung function impairment and risk of incident heart failure: the NHLBI Pooled Cohorts Study. European Heart Journal, 2022, 43, 2196-2208.	1.0	12
9	Metabolomic Associations of Asthma in the Hispanic Community Health Study/Study of Latinos. Metabolites, 2022, 12, 359.	1.3	1
10	Pulmonary Function and Blood DNA Methylation: A Multiancestry Epigenome-Wide Association Meta-analysis. American Journal of Respiratory and Critical Care Medicine, 2022, 206, 321-336.	2.5	15
11	Biomass fuel use and birth weight among term births in Nigeria. PLOS Global Public Health, 2022, 2, e0000419.	0.5	0
12	Assessing the contribution of rare genetic variants to phenotypes of chronic obstructive pulmonary disease using whole-genome sequence data. Human Molecular Genetics, 2022, 31, 3873-3885.	1.4	2
13	Longitudinal associations of DNA methylation and sleep in children: a meta-analysis. Clinical Epigenetics, 2022, 14, .	1.8	6
14	House dust microbiota in relation to adult asthma and atopy in a US farming population. Journal of Allergy and Clinical Immunology, 2021, 147, 910-920.	1.5	21
15	Lrp1 Regulation of Pulmonary Function. Follow-Up of Human GWAS in Mice. American Journal of Respiratory Cell and Molecular Biology, 2021, 64, 368-378.	1.4	7
16	Maternal anxiety during pregnancy and newborn epigenome-wide DNA methylation. Molecular Psychiatry, 2021, 26, 1832-1845.	4.1	24
17	Epigenome-wide association study of kidney function identifies trans-ethnic and ethnic-specific loci. Genome Medicine, 2021, 13, 74.	3.6	20
18	Interaction between Genetic Risk Scores for reduced pulmonary function and smoking, asthma and endotoxin. Thorax, 2021, 76, 1219-1226.	2.7	7

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19	Association of medically assisted reproduction with offspring cord blood DNA methylation across cohorts. Human Reproduction, 2021, 36, 2403-2413.	0.4	17
20	A systematic analysis of protein-altering exonic variants in chronic obstructive pulmonary disease. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2021, 321, L130-L143.	1.3	11
21	Placental DNA methylation signatures of maternal smoking during pregnancy and potential impacts on fetal growth. Nature Communications, 2021, 12, 5095.	5.8	41
22	Newborn DNA Methylation Signatures Related to Prenatal Smoking Exposures in the PACE Consortium. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
23	Rare and low-frequency exonic variants and gene-by-smoking interactions in pulmonary function. Scientific Reports, 2021, 11, 19365.	1.6	2
24	Epigenome-Wide DNA Methylation and Pesticide Use in the Agricultural Lung Health Study. Environmental Health Perspectives, 2021, 129, 97008.	2.8	20
25	Increasing the Impact of Environmental Epidemiology in the Global Burden of Disease Project. Epidemiology, 2021, 32, 1-5.	1.2	3
26	Association Between Preserved Ratio Impaired Spirometry and Clinical Outcomes in US Adults. JAMA - Journal of the American Medical Association, 2021, 326, 2287.	3.8	74
27	Lung Function and Dementia Risk in the Atherosclerosis Risk in Communities (ARIC) Study. Innovation in Aging, 2021, 5, 651-651.	0.0	0
28	Lung function decline in former smokers and low-intensity current smokers: a secondary data analysis of the NHLBI Pooled Cohorts Study. Lancet Respiratory Medicine,the, 2020, 8, 34-44.	5.2	96
29	Whole genome sequence analysis of pulmonary function and COPD in 19,996 multi-ethnic participants. Nature Communications, 2020, 11, 5182.	5.8	32
30	DNA methylation and body mass index from birth to adolescence: meta-analyses of epigenome-wide association studies. Genome Medicine, 2020, 12, 105.	3.6	41
31	Long-term exposure to PM10 and NO2 in relation to lung function and imaging phenotypes in a COPD cohort. Respiratory Research, 2020, 21, 247.	1.4	20
32	Cord blood DNA methylation reflects cord blood C-reactive protein levels but not maternal levels: a longitudinal study and meta-analysis. Clinical Epigenetics, 2020, 12, 60.	1.8	9
33	Lung Development Genes and Adult Lung Function. American Journal of Respiratory and Critical Care Medicine, 2020, 202, 853-865.	2.5	23
34	Epigenome-wide association study of DNA methylation and adult asthma in the Agricultural Lung Health Study. European Respiratory Journal, 2020, 56, 2000217.	3.1	40
35	Epigenetic biomarkers and preterm birth. Environmental Epigenetics, 2020, 6, dvaa005.	0.9	19
36	Identifying potential causal effects of age at menarche: a Mendelian randomization phenome-wide association study. BMC Medicine, 2020, 18, 71.	2.3	27

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37	Epigenome-wide meta-analysis of blood DNA methylation in newborns and children identifies numerous loci related to gestational age. Genome Medicine, 2020, 12, 25.	3.6	81
38	Child Electronic Growth Monitoring System: An innovative and sustainable approach for establishing the Kaduna Infant Development (KID) Study in Nigeria. Paediatric and Perinatal Epidemiology, 2020, 34, 532-543.	0.8	6
39	Association of Nonobstructive Chronic Bronchitis With Respiratory Health Outcomes in Adults. JAMA Internal Medicine, 2020, 180, 676.	2.6	33
40	Outdoor Air Pollution and New-Onset Airway Disease. An Official American Thoracic Society Workshop Report. Annals of the American Thoracic Society, 2020, 17, 387-398.	1.5	120
41	Association of intimate partner violence during pregnancy and birth weight among term births: a cross-sectional study in Kaduna, Northwestern Nigeria. BMJ Open, 2020, 10, e036320.	0.8	6
42	Methylation, smoking, and reduced lung function. European Respiratory Journal, 2019, 54, 1900920.	3.1	8
43	Improving and Expanding Estimates of the Global Burden of Disease Due to Environmental Health Risk Factors. Environmental Health Perspectives, 2019, 127, 105001.	2.8	73
44	Associations of autozygosity with a broad range of human phenotypes. Nature Communications, 2019, 10, 4957.	5.8	84
45	Comparison of smoking-related DNA methylation between newborns from prenatal exposure and adults from personal smoking. Epigenomics, 2019, 11, 1487-1500.	1.0	64
46	Epigenome-wide meta-analysis of DNA methylation and childhood asthma. Journal of Allergy and Clinical Immunology, 2019, 143, 2062-2074.	1.5	147
47	Maternal levels of perfluoroalkyl substances (PFASs) during pregnancy and childhood allergy and asthma related outcomes and infections in the Norwegian Mother and Child (MoBa) cohort. Environment International, 2019, 124, 462-472.	4.8	64
48	Hypertensive Disorders of Pregnancy and DNA Methylation in Newborns. Hypertension, 2019, 74, 375-383.	1.3	73
49	Prenatal Particulate Air Pollution and DNA Methylation in Newborns: An Epigenome-Wide Meta-Analysis. Environmental Health Perspectives, 2019, 127, 57012.	2.8	111
50	Integrated analysis of environmental and genetic influences on cord blood DNA methylation in new-borns. Nature Communications, 2019, 10, 2548.	5.8	94
51	Genomic interactions with exposure to inhaled pollutants. Journal of Allergy and Clinical Immunology, 2019, 143, 2011-2013.e1.	1.5	6
52	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	5.8	140
53	Animal production, insecticide use and self-reported symptoms and diagnoses of COPD, including chronic bronchitis, in the Agricultural Health Study. Environment International, 2019, 127, 764-772.	4.8	17
54	DNA methylation signature of smoking in lung cancer is enriched for exposure signatures in newborn and adult blood. Scientific Reports, 2019, 9, 4576.	1.6	32

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55	Genome-wide DNA methylation and long-term ambient air pollution exposure in Korean adults. Clinical Epigenetics, 2019, 11, 37.	1.8	76
56	Metabolomics Identifies Novel Blood Biomarkers of Pulmonary Function and COPD in the General Population. Metabolites, 2019, 9, 61.	1.3	30
57	Genetic landscape of chronic obstructive pulmonary disease identifies heterogeneous cell-type and phenotype associations. Nature Genetics, 2019, 51, 494-505.	9.4	257
58	Semen quality and cigarette smoking in a cohort of healthy fertile men. Environmental Epidemiology, 2019, 3, e055.	1.4	20
59	Omega-3 Fatty Acids and Genome-Wide Interaction Analyses Reveal DPP10â€“Pulmonary Function Association. American Journal of Respiratory and Critical Care Medicine, 2019, 199, 631-642.	2.5	14
60	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.	1.5	33
61	Maternal history of miscarriages and measures of fertility in relation to childhood asthma. Thorax, 2019, 74, 106-113.	2.7	13
62	Diet Pattern and Respiratory Morbidity in the Atherosclerosis Risk in Communities Study. Annals of the American Thoracic Society, 2018, 15, 675-682.	1.5	40
63	Role of local CpG DNA methylation in mediating the 17q21 asthma susceptibility gasdermin B (GSDMB)/ORMDL sphingolipid biosynthesis regulator 3 (ORMDL3) expression quantitative trait locus. Journal of Allergy and Clinical Immunology, 2018, 141, 2282-2286.e6.	1.5	20
64	A Genome-Wide Association Study in Hispanics/Latinos Identifies Novel Signals for Lung Function. The Hispanic Community Health Study/Study of Latinos. American Journal of Respiratory and Critical Care Medicine, 2018, 198, 208-219.	2.5	37
65	Association of Maternal Psychosocial Stress With Increased Risk of Asthma Development in Offspring. American Journal of Epidemiology, 2018, 187, 1199-1209.	1.6	30
66	Multiancestry association study identifies new asthma risk loci that colocalize with immune-cell enhancer marks. Nature Genetics, 2018, 50, 42-53.	9.4	426
67	Preeclampsia and Hypertension During Pregnancy in Areas with Relatively Low Levels of Traffic Air Pollution. Maternal and Child Health Journal, 2018, 22, 512-519.	0.7	19
68	Vitamin A and D intake in pregnancy, infant supplementation, and asthma development: the Norwegian Mother and Child Cohort. American Journal of Clinical Nutrition, 2018, 107, 789-798.	2.2	32
69	A DNA methylation biomarker of alcohol consumption. Molecular Psychiatry, 2018, 23, 422-433.	4.1	280
70	Raw milk consumption and other early-life farm exposures and adult pulmonary function in the Agricultural Lung Health Study. Thorax, 2018, 73, 279-282.	2.7	19
71	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	0.9	105
72	Sleep apnea and pesticide exposure in a study of US farmers. Sleep Health, 2018, 4, 20-26.	1.3	21

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73	Maternal alcohol consumption and offspring DNA methylation: findings from six general population-based birth cohorts. Epigenomics, 2018, 10, 27-42.	1.0	58
74	Vitamin D and Risk of Pregnancy-Related Hypertensive Disorders: Mendelian Randomization Study. Obstetrical and Gynecological Survey, 2018, 73, 617-619.	0.2	0
75	Genetic-Epigenetic Interactions in Asthma Revealed by a Genome-Wide Gene-Centric Search. Human Heredity, 2018, 83, 130-152.	0.4	18
76	Gene Coexpression Networks in Whole Blood Implicate Multiple Interrelated Molecular Pathways in Obesity in People with Asthma. Obesity, 2018, 26, 1938-1948.	1.5	11
77	Ambient Air Pollution and Chronic Bronchitis in a Cohort of U.S. Women. Environmental Health Perspectives, 2018, 126, 027005.	2.8	55
78	Exposures Related to House Dust Microbiota in a U.S. Farming Population. Environmental Health Perspectives, 2018, 126, 067001.	2.8	23
79	Meta-analysis across Cohorts for Heart and Aging Research in Genomic Epidemiology (CHARGE) consortium provides evidence for an association of serum vitamin D with pulmonary function. British Journal of Nutrition, 2018, 120, 1159-1170.	1.2	9
80	Harmonization of Respiratory Data From 9 US Population-Based Cohorts. American Journal of Epidemiology, 2018, 187, 2265-2278.	1.6	46
81	Declining Lung Function and Cardiovascular Risk. Journal of the American College of Cardiology, 2018, 72, 1109-1122.	1.2	74
82	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. Nature Communications, 2018, 9, 2976.	5.8	85
83	Prenatal iron exposure and childhood type 1 diabetes. Scientific Reports, 2018, 8, 9067.	1.6	25
84	Vitamin D and risk of pregnancy related hypertensive disorders: mendelian randomisation study. BMJ: British Medical Journal, 2018, 361, k2167.	2.4	31
85	Meta-analysis of exome array data identifies six novel genetic loci for lung function. Wellcome Open Research, 2018, 3, 4.	0.9	19
86	Evidence for large-scale gene-by-smoking interaction effects on pulmonary function. International Journal of Epidemiology, 2017, 46, dyw318.	0.9	36
87	Misclassified exposure in epigenetic mediation analyses. Does DNA methylation mediate effects of smoking on birthweight?. Epigenomics, 2017, 9, 253-265.	1.0	42
88	Genetic loci associated with chronic obstructive pulmonary disease overlap with loci for lung function and pulmonary fibrosis. Nature Genetics, 2017, 49, 426-432.	9.4	306
89	House Dust Endotoxin Levels Are Associated with Adult Asthma in a U.S. Farming Population. Annals of the American Thoracic Society, 2017, 14, 324-331.	1.5	47
90	Classifying oxidative stress by F2-isoprostane levels across human diseases: A meta-analysis. Redox Biology, 2017, 12, 582-599.	3.9	134

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91	Epigenome-wide association study of chronic obstructive pulmonary disease and lung function in Koreans. Epigenomics, 2017, 9, 971-984.	1.0	39
92	Adverse Experiences in Childhood and Sexually Transmitted Infection Risk From Adolescence Into Adulthood. Sexually Transmitted Diseases, 2017, 44, 524-532.	0.8	42
93	Vagal innervation is required for pulmonary function phenotype in Htr4^{â^'/â^'} mice. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2017, 312, L520-L530.	1.3	2
94	Maternal BMI at the start of pregnancy and offspring epigenome-wide DNA methylation: findings from the pregnancy and childhood epigenetics (PACE) consortium. Human Molecular Genetics, 2017, 26, 4067-4085.	1.4	211
95	Global, regional, and national incidence, prevalence, and years lived with disability for 328 diseases and injuries for 195 countries, 1990â€“2016: a systematic analysis for the Global Burden of Disease Study 2016. Lancet, The, 2017, 390, 1211-1259.	6.3	5,578
96	Early-life farm exposures and adult asthma and atopy in the Agricultural Lung Health Study. Journal of Allergy and Clinical Immunology, 2017, 140, 249-256.e14.	1.5	61
97	Maternal Folate Intake during Pregnancy and Childhood Asthma in a Population-based Cohort. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 221-228.	2.5	44
98	Gene Expression Profiling in Blood Provides Reproducible Molecular Insights into Asthma Control. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 179-188.	2.5	49
99	Genome-Wide Association Analysis of the Sense of Smell in U.S. Older Adults: Identification of Novel Risk Loci in African-Americans and European-Americans. Molecular Neurobiology, 2017, 54, 8021-8032.	1.9	17
100	Correcting Oxidative Stress Measurements using the 8-iso-PGF 2Î± /PGF 2Î± Ratio to Determine Appropriate Interventions. Free Radical Biology and Medicine, 2017, 112, 135-136.	1.3	0
101	Pregnancy exposure to air pollution and early childhood respiratory health in the Norwegian Mother and Child Cohort Study (MoBa). BMJ Open, 2017, 7, e015796.	0.8	13
102	Epigenome-Wide Meta-Analysis of Methylation in Children Related to Prenatal NO ₂ Air Pollution Exposure. Environmental Health Perspectives, 2017, 125, 104-110.	2.8	176
103	Pesticides are Associated with Allergic and Non-Allergic Wheeze among Male Farmers. Environmental Health Perspectives, 2017, 125, 535-543.	2.8	82
104	House Dust Endotoxin and Peripheral Leukocyte Counts: Results from Two Large Epidemiologic Studies. Environmental Health Perspectives, 2017, 125, 057010.	2.8	7
105	DNA Methylation Score as a Biomarker in Newborns for Sustained Maternal Smoking during Pregnancy. Environmental Health Perspectives, 2017, 125, 760-766.	2.8	86
106	Maternal Age at Delivery Is Associated with an Epigenetic Signature in Both Newborns and Adults. PLoS ONE, 2016, 11, e0156361.	1.1	62
107	Effect of Obesity on Acute Ozone-Induced Changes in Airway Function, Reactivity, and Inflammation in Adult Females. PLoS ONE, 2016, 11, e0160030.	1.1	29
108	Maternal smoking impacts key biological pathways in newborns through epigenetic modification in Utero. BMC Genomics, 2016, 17, 976.	1.2	56

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109	DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium Meta-analysis. American Journal of Human Genetics, 2016, 98, 680-696.	2.6	717
110	Lung function, respiratory symptoms and venous thromboembolism risk: the Atherosclerosis Risk in Communities Study. Journal of Thrombosis and Haemostasis, 2016, 14, 2394-2401.	1.9	32
111	Prediction of gestational age based on genome-wide differentially methylated regions. Genome Biology, 2016, 17, 207.	3.8	132
112	Epigenetic Signatures of Cigarette Smoking. Circulation: Cardiovascular Genetics, 2016, 9, 436-447.	5.1	678
113	Association between pregnancy exposure to air pollution and birth weight in selected areas of Norway. Archives of Public Health, 2016, 74, 26.	1.0	19
114	DNA methylation and smoking in Korean adults: epigenome-wide association study. Clinical Epigenetics, 2016, 8, 103.	1.8	60
115	Maternal plasma total neopterin and kynurenine/tryptophan levels during pregnancy in relation to asthma development in the offspring. Journal of Allergy and Clinical Immunology, 2016, 138, 1319-1325.e4.	1.5	4
116	25-Hydroxyvitamin D in pregnancy and genome wide cord blood DNA methylation in two pregnancy cohorts (MoBa and ALSPAC). Journal of Steroid Biochemistry and Molecular Biology, 2016, 159, 102-109.	1.2	26
117	Lung function decline over 25 years of follow-up among black and white adults in the ARIC study cohort. Respiratory Medicine, 2016, 113, 57-64.	1.3	23
118	Maternal plasma folate impacts differential DNA methylation in an epigenome-wide meta-analysis of newborns. Nature Communications, 2016, 7, 10577.	5.8	219
119	Association of Forced Vital Capacity with the Developmental Gene NCOR2. PLoS ONE, 2016, 11, e0147388.	1.1	17
120	Particulate Matter 2.5 Exposure and Self-Reported Use of Wood Stoves and Other Indoor Combustion Sources in Urban Nonsmoking Homes in Norway. PLoS ONE, 2016, 11, e0166440.	1.1	20
121	Noninvasive Analysis of the Sputum Transcriptome Discriminates Clinical Phenotypes of Asthma. Annals of the American Thoracic Society, 2016, 13, S104-S105.	1.5	10
122	A genome-wide association study of asthma symptoms in Latin American children. BMC Genetics, 2015, 16, 141.	2.7	24
123	Indoor determinants of dustborne allergens in Mexican homes. Allergy and Asthma Proceedings, 2015, 36, 130-137.	1.0	6
124	Genome-wide Meta-analysis on the Sense of Smell Among US Older Adults. Medicine (United States), 2015, 94, e1892.	0.4	12
125	Long-Term Air Pollution Exposure and Blood Pressure in the Sister Study. Environmental Health Perspectives, 2015, 123, 951-958.	2.8	136
126	Peak Weight and Height Velocity to Age 36 Months and Asthma Development: The Norwegian Mother and Child Cohort Study. PLoS ONE, 2015, 10, e0116362.	1.1	17

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127	A Study on Mediation by Offspring BMI in the Association between Maternal Obesity and Child Respiratory Outcomes in the Amsterdam Born and Their Development Study Cohort. PLoS ONE, 2015, 10, e0140641.	1.1	33
128	Global Analysis of Methylation Profiles From High Resolution CpG Data. Genetic Epidemiology, 2015, 39, 53-64.	0.6	19
129	Stress and Bronchodilator Response in Children with Asthma. American Journal of Respiratory and Critical Care Medicine, 2015, 192, 47-56.	2.5	99
130	Genetic variation in HTR4 and lung function: GWAS followâ€up in mouse. FASEB Journal, 2015, 29, 323-335.	0.2	16
131	Directional dominance on stature and cognition inÂdiverse human populations. Nature, 2015, 523, 459-462.	13.7	173
132	Noninvasive Analysis of the Sputum Transcriptome Discriminates Clinical Phenotypes of Asthma. American Journal of Respiratory and Critical Care Medicine, 2015, 191, 1116-1125.	2.5	86
133	Grandmother's smoking when pregnant with the mother and asthma in the grandchild: the Norwegian Mother and Child Cohort Study. Thorax, 2015, 70, 237-243.	2.7	88
134	Molecular mechanisms underlying variations in lung function: a systems genetics analysis. Lancet Respiratory Medicine,the, 2015, 3, 782-795.	5.2	66
135	Integrative pathway genomics of lung function and airflow obstruction. Human Molecular Genetics, 2015, 24, 6836-6848.	1.4	28
136	Effect of maternal gestational weight gain on offspring DNA methylation: a follow-up to the ALSPAC cohort study. BMC Research Notes, 2015, 8, 321.	0.6	12
137	Ethnic-specific associations of rare and low-frequency DNA sequence variants with asthma. Nature Communications, 2015, 6, 5965.	5.8	66
138	Global, regional, and national comparative risk assessment of 79 behavioural, environmental and occupational, and metabolic risks or clusters of risks in 188 countries, 1990â€“2013: a systematic analysis for the Global Burden of Disease Study 2013. Lancet, The, 2015, 386, 2287-2323.	6.3	2,184
139	Global, regional, and national ageâ€“sex specific all-cause and cause-specific mortality for 240 causes of death, 1990â€“2013: a systematic analysis for the Global Burden of Disease Study 2013. Lancet, The, 2015, 385, 117-171.	6.3	5,847
140	Glucocorticoid Genes and the Developmental Origins of Asthma Susceptibility and Treatment Response. American Journal of Respiratory Cell and Molecular Biology, 2015, 52, 543-553.	1.4	22
141	Prenatal Tobacco Smoke Exposure Is Associated with Childhood DNA CpG Methylation. PLoS ONE, 2014, 9, e99716.	1.1	105
142	Large-Scale Genome-Wide Association Studies and Meta-Analyses of Longitudinal Change in Adult Lung Function. PLoS ONE, 2014, 9, e100776.	1.1	52
143	Respiratory disease in United States farmers. Occupational and Environmental Medicine, 2014, 71, 484-491.	1.3	66
144	ADAM19 and HTR4 Variants and Pulmonary Function. Circulation: Cardiovascular Genetics, 2014, 7, 350-358.	5.1	8

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145	Airflow Obstruction, Lung Function, and Incidence of Atrial Fibrillation. Circulation, 2014, 129, 971-980.	1.6	103
146	Prospective Study of Maternal Alcohol Intake During Pregnancy or Lactation and Risk of Childhood Asthma: The <scp>N</scp>orwegian Mother and Child Cohort Study. Alcoholism: Clinical and Experimental Research, 2014, 38, 1002-1011.	1.4	15
147	APOM and high-density lipoprotein cholesterol are associated with lung function and per cent emphysema. European Respiratory Journal, 2014, 43, 1003-1017.	3.1	37
148	Genome-wide interaction studies reveal sex-specific asthma risk alleles. Human Molecular Genetics, 2014, 23, 5251-5259.	1.4	70
149	Food allergens in mattress dust in <scp>N</scp>orwegian homes â€“ a potentially important source of allergen exposure. Clinical and Experimental Allergy, 2014, 44, 142-149.	1.4	39
150	Neonatal Genome-Wide Methylation Patterns in Relation to Birth Weight in the Norwegian Mother and Child Cohort. American Journal of Epidemiology, 2014, 179, 834-842.	1.6	92
151	Ambient Air Pollution Exposure and Incident Adult Asthma in a Nationwide Cohort of U.S. Women. American Journal of Respiratory and Critical Care Medicine, 2014, 190, 914-921.	2.5	132
152	A systematic assessment of normalization approaches for the Infinium 450K methylation platform. Epigenetics, 2014, 9, 318-329.	1.3	61
153	Exacerbation of symptoms in agricultural pesticide applicators with asthma. International Archives of Occupational and Environmental Health, 2014, 87, 423-432.	1.1	45
154	Probiotic milk consumption in pregnancy and infancy and subsequent childhood allergic diseases. Journal of Allergy and Clinical Immunology, 2014, 133, 165-171.e8.	1.5	105
155	Reliability of triclosan measures in repeated urine samples from Norwegian pregnant women. Journal of Exposure Science and Environmental Epidemiology, 2014, 24, 517-521.	1.8	48
156	A genome-wide survey of CD4+ lymphocyte regulatory genetic variants identifies novel asthma genes. Journal of Allergy and Clinical Immunology, 2014, 134, 1153-1162.	1.5	46
157	Gender difference in interactions between <scp>MAOA</scp> promoter <scp>uVNTR</scp> polymorphism and negative familial stressors on body mass index among <scp>C</scp>hinese adolescents. Pediatric Obesity, 2014, 9, e80-90.	1.4	7
158	Genome-wide association analysis identifies six new loci associated with forced vital capacity. Nature Genetics, 2014, 46, 669-677.	9.4	131
159	Maternal Smoking and DNA Methylation in Newborns: In Utero Effect or Epigenetic Inheritance?. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1007-1017.	1.1	108
160	Integrated genome-wide association, coexpression network, and expression single nucleotide polymorphism analysis identifies novel pathway in allergic rhinitis. BMC Medical Genomics, 2014, 7, 48.	0.7	63
161	Common genes underlying asthma and COPD? Genome-wide analysis on the Dutch hypothesis. European Respiratory Journal, 2014, 44, 860-872.	3.1	49
162	Global, regional, and national levels and causes of maternal mortality during 1990â€“2013: a systematic analysis for the Global Burden of Disease Study 2013. Lancet, The, 2014, 384, 980-1004.	6.3	1,230

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163	Genome-wide association study and admixture mapping identify different asthma-associated loci in Latinos: The Genes-environments & Admixture in Latino Americans study. Journal of Allergy and Clinical Immunology, 2014, 134, 295-305.	1.5	106
164	The genetics of Mexico recapitulates Native American substructure and affects biomedical traits. Science, 2014, 344, 1280-1285.	6.0	420
165	Accurate construction of long range haplotype In unrelated individuals. Statistica Sinica, 2014, , .	0.2	0
166	The State of US Health, 1990-2010. JAMA - Journal of the American Medical Association, 2013, 310, 591.	3.8	2,070
167	Ozone exposure, vitamin C intake, and genetic susceptibility of asthmatic children in Mexico City: a cohort study. Respiratory Research, 2013, 14, 14.	1.4	33
168	Replication and fine mapping of asthma-associated loci in individuals of African ancestry. Human Genetics, 2013, 132, 1039-1047.	1.8	12
169	Nitrogen dioxide and allergic sensitization in the 2005â€“2006 National Health and Nutrition Examination Survey. Respiratory Medicine, 2013, 107, 1763-1772.	1.3	28
170	Maternal Vitamin D Status During Pregnancy and Asthma in the Offspring Among Participants in the Norwegian Mother and Child Cohort Study. Journal of Allergy and Clinical Immunology, 2013, 131, AB128.	1.5	0
171	A meta-analysis of genome-wide association studies for serum total IgE in diverse study populations. Journal of Allergy and Clinical Immunology, 2013, 131, 1176-1184.	1.5	58
172	Triclosan exposure and allergic sensitization in <scp>N</scp>orwegian children. Allergy: European Journal of Allergy and Clinical Immunology, 2013, 68, 84-91.	2.7	85
173	Phthalate Exposure and Allergy in the U.S. Population: Results from NHANES 2005â€“2006. Environmental Health Perspectives, 2013, 121, 1129-1134.	2.8	113
174	Prospective Study of Maternal Midâ€pregnancy 25â€hydroxyvitamin <scp>D</scp> Level and Early Childhood Respiratory Disorders. Paediatric and Perinatal Epidemiology, 2013, 27, 532-541.	0.8	53
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