William Cookson

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

147	21,993	59	148
papers	citations	h-index	g-index
167 ext. papers	25,684 ext. citations	12.2 avg, IF	6.44 L-index

#	Paper	IF	Citations
147	Evidence of immunometabolic dysregulation and airway dysbiosis in athletes susceptible to respiratory illness <i>EBioMedicine</i> , 2022 , 79, 104024	8.8	1
146	Bacterial Signatures of Paediatric Respiratory Disease: An Individual Participant Data Meta-Analysis <i>Frontiers in Microbiology</i> , 2021 , 12, 711134	5.7	0
145	ORMDL3 regulates poly I:C induced inflammatory responses in airway epithelial cells. <i>BMC Pulmonary Medicine</i> , 2021 , 21, 167	3.5	1
144	Y disruption, autosomal hypomethylation and poor male lung cancer survival. <i>Scientific Reports</i> , 2021 , 11, 12453	4.9	3
143	The fungal airway microbiome in cystic fibrosis and non-cystic fibrosis bronchiectasis. <i>Journal of Cystic Fibrosis</i> , 2021 , 20, 295-302	4.1	16
142	A large-scale genome-wide association analysis of lung function in the Chinese population identifies novel loci and highlights shared genetic aetiology with obesity. <i>European Respiratory Journal</i> , 2021 , 58,	13.6	2
141	Airway microbial communities, smoking and asthma in a general population sample. <i>EBioMedicine</i> , 2021 , 71, 103538	8.8	5
140	Integrated genomics point to immune vulnerabilities in pleural mesothelioma. <i>Scientific Reports</i> , 2021 , 11, 19138	4.9	0
139	Estimating cell-type-specific DNA methylation effects in heterogeneous cellular populations. <i>Epigenomics</i> , 2021 , 13, 87-97	4.4	2
138	Presence of pleomorphic features but not growth patterns improves prognostic stratification of epithelioid malignant pleural mesothelioma by 2-tier nuclear grade. <i>Histopathology</i> , 2020 , 77, 423-436	7.3	5
137	Shared genetic and experimental links between obesity-related traits and asthma subtypes in UK Biobank. <i>Journal of Allergy and Clinical Immunology</i> , 2020 , 145, 537-549	11.5	70
136	Utility of Nuclear Grading System in Epithelioid Malignant Pleural Mesothelioma in Biopsy-heavy Setting: An External Validation Study of 563 Cases. <i>American Journal of Surgical Pathology</i> , 2020 , 44, 347-356	6.7	15
135	The undiagnosed disease burden associated with alpha-1 antitrypsin deficiency genotypes. <i>European Respiratory Journal</i> , 2020 , 56,	13.6	10
134	Metabolomic, transcriptomic and genetic integrative analysis reveals important roles of adenosine diphosphate in haemostasis and platelet activation in non-small-cell lung cancer. <i>Molecular Oncology</i> , 2019 , 13, 2406-2421	7.9	10
133	Inhaled corticosteroid suppression of cathelicidin drives dysbiosis and bacterial infection in chronic obstructive pulmonary disease. <i>Science Translational Medicine</i> , 2019 , 11,	17.5	35
132	Longitudinal development of the airway microbiota in infants with cystic fibrosis. <i>Scientific Reports</i> , 2019 , 9, 5143	4.9	11
131	A Haemophilus sp. dominates the microbiota of sputum from UK adults with non-severe community acquired pneumonia and chronic lung disease. <i>Scientific Reports</i> , 2019 , 9, 2388	4.9	5

(2017-2019)

130	Genome-wide interaction study of early-life smoking exposure on time-to-asthma onset in childhood. <i>Clinical and Experimental Allergy</i> , 2019 , 49, 1342-1351	4.1	4
129	EGF receptor (EGFR) inhibition promotes a slow-twitch oxidative, over a fast-twitch, muscle phenotype. <i>Scientific Reports</i> , 2019 , 9, 9218	4.9	4
128	Viral respiratory infections and the oropharyngeal bacterial microbiota in acutely wheezing children. <i>PLoS ONE</i> , 2019 , 14, e0223990	3.7	6
127	The ORMDL3 Asthma Gene Regulates ICAM1 and Has Multiple Effects on Cellular Inflammation. American Journal of Respiratory and Critical Care Medicine, 2019, 199, 478-488	10.2	45
126	DNA methylation in childhood asthma: an epigenome-wide meta-analysis. <i>Lancet Respiratory Medicine,the</i> , 2018 , 6, 379-388	35.1	119
125	Multiancestry association study identifies new asthma risk loci that colocalize with immune-cell enhancer marks. <i>Nature Genetics</i> , 2018 , 50, 42-53	36.3	246
124	Role of airway glucose in bacterial infections in patients with chronic obstructive pulmonary disease. <i>Journal of Allergy and Clinical Immunology</i> , 2018 , 142, 815-823.e6	11.5	42
123	New opportunities for managing acute and chronic lung infections. <i>Nature Reviews Microbiology</i> , 2018 , 16, 111-120	22.2	46
122	A novel role for ciliary function in atopy: ADGRV1 and DNAH5 interactions. <i>Journal of Allergy and Clinical Immunology</i> , 2018 , 141, 1659-1667.e11	11.5	6
121	Metal worker's lung: spatial association with. <i>Thorax</i> , 2018 , 73, 151-156	7.3	11
121	Metal worker's lung: spatial association with. <i>Thorax</i> , 2018 , 73, 151-156 COPD is accompanied by co-ordinated transcriptional perturbation in the quadriceps affecting the mitochondria and extracellular matrix. <i>Scientific Reports</i> , 2018 , 8, 12165	7·3 4·9	17
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120	COPD is accompanied by co-ordinated transcriptional perturbation in the quadriceps affecting the mitochondria and extracellular matrix. <i>Scientific Reports</i> , 2018 , 8, 12165 Comparison of the upper and lower airway microbiota in children with chronic lung diseases. <i>PLoS ONE</i> , 2018 , 13, e0201156 Corticosteroid suppression of antiviral immunity increases bacterial loads and mucus production in	4.9	17
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120 119 118	COPD is accompanied by co-ordinated transcriptional perturbation in the quadriceps affecting the mitochondria and extracellular matrix. <i>Scientific Reports</i> , 2018 , 8, 12165 Comparison of the upper and lower airway microbiota in children with chronic lung diseases. <i>PLoS ONE</i> , 2018 , 13, e0201156 Corticosteroid suppression of antiviral immunity increases bacterial loads and mucus production in COPD exacerbations. <i>Nature Communications</i> , 2018 , 9, 2229 Manipulation of dipeptidylpeptidase 10 in mouse and human and models indicates a protective role in asthma. <i>DMM Disease Models and Mechanisms</i> , 2018 , 11, Whole-Blood Gene Expression in Pulmonary Nontuberculous Mycobacterial Infection. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2018 , 58, 510-518	4·9 3·7 17·4 4·1	17 13 100
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112	Host-Microbial Interactions in Idiopathic Pulmonary Fibrosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2017 , 195, 1640-1650	10.2	114
111	Changes in the respiratory microbiome during acute exacerbations of idiopathic pulmonary fibrosis. <i>Respiratory Research</i> , 2017 , 18, 29	7.3	108
110	Network-assisted analysis of GWAS data identifies a functionally-relevant gene module for childhood-onset asthma. <i>Scientific Reports</i> , 2017 , 7, 938	4.9	11
109	Addressing unmet needs in understanding asthma mechanisms: From the European Asthma Research and Innovation Partnership (EARIP) Work Package (WP)2 collaborators. <i>European Respiratory Journal</i> , 2017 , 49,	13.6	31
108	A mechanistic target of rapamycin complex 1/2 (mTORC1)/V-Akt murine thymoma viral oncogene homolog 1 (AKT1)/cathepsin H axis controls filaggrin expression and processing in skin, a novel mechanism for skin barrier disruption in patients with atopic dermatitis. <i>Journal of Allergy and</i>	11.5	20
107	Clinical Immunology, 2017, 139, 1228-1241 The impact of persistent bacterial bronchitis on the pulmonary microbiome of children. <i>PLoS ONE</i> , 2017, 12, e0190075	3.7	19
106	Vitamin D levels and susceptibility to asthma, elevated immunoglobulin E levels, and atopic dermatitis: A Mendelian randomization study. <i>PLoS Medicine</i> , 2017 , 14, e1002294	11.6	47
105	Validation of a 52-gene risk profile for outcome prediction in patients with idiopathic pulmonary fibrosis: an international, multicentre, cohort study. <i>Lancet Respiratory Medicine,the</i> , 2017 , 5, 857-868	35.1	59
104	Bacterial microbiota of the upper respiratory tract and childhood asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2017 , 139, 826-834.e13	11.5	117
103	Pulmonary ORMDL3 is critical for induction of Alternaria-induced allergic airways disease. <i>Journal of Allergy and Clinical Immunology</i> , 2017 , 139, 1496-1507.e3	11.5	61
102	Longitudinal assessment of sputum microbiome by sequencing of the 16S rRNA gene in non-cystic fibrosis bronchiectasis patients. <i>PLoS ONE</i> , 2017 , 12, e0170622	3.7	70
101	Identification of a new locus at 16q12 associated with time to asthma onset. <i>Journal of Allergy and Clinical Immunology</i> , 2016 , 138, 1071-1080	11.5	21
100	Airway Microbiota in Severe Asthma and Relationship to Asthma Severity and Phenotypes. <i>PLoS ONE</i> , 2016 , 11, e0152724	3.7	104
99	Global gene regulation during activation of immunoglobulin class switching in human B cells. <i>Scientific Reports</i> , 2016 , 6, 37988	4.9	15
98	DNA methylation within melatonin receptor 1A (MTNR1A) mediates paternally transmitted genetic variant effect on asthma plus rhinitis. <i>Journal of Allergy and Clinical Immunology</i> , 2016 , 138, 748-753	11.5	17
97	eQTL mapping identifies insertion- and deletion-specific eQTLs in multiple tissues. <i>Nature Communications</i> , 2015 , 6, 6821	17.4	11
96	Effects of different antibiotic classes on airway bacteria in stable COPD using culture and molecular techniques: a randomised controlled trial. <i>Thorax</i> , 2015 , 70, 930-8	7-3	45
95	Imputation of KIR Types from SNP Variation Data. <i>American Journal of Human Genetics</i> , 2015 , 97, 593-6	071	44

(2013-2015)

94	Dysbiosis anticipating necrotizing enterocolitis in very premature infants. <i>Clinical Infectious Diseases</i> , 2015 , 60, 389-97	11.6	118
93	iGWAS: Integrative Genome-Wide Association Studies of Genetic and Genomic Data for Disease Susceptibility Using Mediation Analysis. <i>Genetic Epidemiology</i> , 2015 , 39, 347-56	2.6	32
92	Late-Onset Bloodstream Infection and Perturbed Maturation of the Gastrointestinal Microbiota in Premature Infants. <i>PLoS ONE</i> , 2015 , 10, e0132923	3.7	56
91	An epigenome-wide association study of total serum immunoglobulin E concentration. <i>Nature</i> , 2015 , 520, 670-674	50.4	162
90	The role of bacteria in the pathogenesis and progression of idiopathic pulmonary fibrosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2014 , 190, 906-13	10.2	320
89	Novel childhood asthma genes interact with in utero and early-life tobacco smoke exposure. Journal of Allergy and Clinical Immunology, 2014 , 133, 885-8	11.5	36
88	Functional analysis of a novel ENU-induced PHD finger 11 (Phf11) mouse mutant. <i>Mammalian Genome</i> , 2014 , 25, 573-82	3.2	6
87	Reagent and laboratory contamination can critically impact sequence-based microbiome analyses. <i>BMC Biology</i> , 2014 , 12, 87	7.3	1745
86	Grasping nettles: cellular heterogeneity and other confounders in epigenome-wide association studies. <i>Human Molecular Genetics</i> , 2014 , 23, R83-8	5.6	36
85	Predicting DNA methylation level across human tissues. <i>Nucleic Acids Research</i> , 2014 , 42, 3515-28	20.1	81
84	Fraction of exhaled nitric oxide values in childhood are associated with 17q11.2-q12 and 17q12-q21 variants. <i>Journal of Allergy and Clinical Immunology</i> , 2014 , 134, 46-55	11.5	27
83	A functional IL-6 receptor (IL6R) variant is a risk factor for persistent atopic dermatitis. <i>Journal of Allergy and Clinical Immunology</i> , 2013 , 132, 371-7	11.5	65
82	Meta-analysis of gene-level associations for rare variants based on single-variant statistics. American Journal of Human Genetics, 2013 , 93, 236-48	11	49
81	Sequencing the human microbiome in health and disease. <i>Human Molecular Genetics</i> , 2013 , 22, R88-94	5.6	89
80	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. <i>Nature Genetics</i> , 2013 , 45, 501-12	36.3	437
79	A molecular comparison of microbial communities in bronchiectasis and cystic fibrosis. <i>European Respiratory Journal</i> , 2013 , 41, 991-3	13.6	12
78	Outgrowth of the bacterial airway microbiome after rhinovirus exacerbation of chronic obstructive pulmonary disease. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2013 , 188, 1224-31	10.2	262
77	A cross-platform analysis of 14,177 expression quantitative trait loci derived from lymphoblastoid cell lines. <i>Genome Research</i> , 2013 , 23, 716-26	9.7	117

76	A genome-wide association study of atopic dermatitis identifies loci with overlapping effects on asthma and psoriasis. <i>Human Molecular Genetics</i> , 2013 , 22, 4841-56	5.6	140
75	Analgesia and central side-effects: two separate dimensions of morphine response. <i>British Journal of Clinical Pharmacology</i> , 2013 , 75, 1340-50	3.8	31
74	A polymorphism affecting MYB binding within the promoter of the PDCD4 gene is associated with severe asthma in children. <i>Human Mutation</i> , 2013 , 34, 1131-9	4.7	16
73	Seventy-five genetic loci influencing the human red blood cell. <i>Nature</i> , 2012 , 492, 369-75	50.4	257
72	Impact of collection and storage of lung tumor tissue on whole genome expression profiling. <i>Journal of Molecular Diagnostics</i> , 2012 , 14, 140-8	5.1	30
71	Improved detection of bifidobacteria with optimised 16S rRNA-gene based pyrosequencing. <i>PLoS ONE</i> , 2012 , 7, e32543	3.7	143
70	The origin, global distribution, and functional impact of the human 8p23 inversion polymorphism. <i>Genome Research</i> , 2012 , 22, 1144-53	9.7	48
69	Integrating pathway analysis and genetics of gene expression for genome-wide association study of basal cell carcinoma. <i>Human Genetics</i> , 2012 , 131, 615-23	6.3	26
68	Copy number variation leads to considerable diversity for B but not A haplotypes of the human KIR genes encoding NK cell receptors. <i>Genome Research</i> , 2012 , 22, 1845-54	9.7	127
67	Significance of the microbiome in obstructive lung disease. <i>Thorax</i> , 2012 , 67, 456-63	7.3	161
66	Genetic and genomic approaches to asthma: new insights for the origins. <i>Current Opinion in Pulmonary Medicine</i> , 2012 , 18, 6-13	3	75
65	Genetic risks and childhood-onset asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2011 , 128, 266-70; quiz 271-2	11.5	22
64	Gene-environment interaction in chronic disease: a European Science Foundation Forward Look. <i>Journal of Allergy and Clinical Immunology</i> , 2011 , 128, S27-49	11.5	25
63	Genome-wide association study identifies loci influencing concentrations of liver enzymes in plasma. <i>Nature Genetics</i> , 2011 , 43, 1131-8	36.3	415
62	Gene-environment interactions in chronic inflammatory disease. <i>Nature Immunology</i> , 2011 , 12, 273-7	19.1	124
61	Genetics of complex airway disease. <i>Proceedings of the American Thoracic Society</i> , 2011 , 8, 149-53		19
60	Exposure to environmental microorganisms and childhood asthma. <i>New England Journal of Medicine</i> , 2011 , 364, 701-9	59.2	1060
59	Chromosome 17q21 SNP and severe asthma. <i>Journal of Human Genetics</i> , 2011 , 56, 97-8	4.3	40

(2007-2011)

58	Opportunities and challenges in the genetics of COPD 2010: an International COPD Genetics Conference report. <i>COPD: Journal of Chronic Obstructive Pulmonary Disease</i> , 2011 , 8, 121-35	2	38
57	Meta-analysis of 20 genome-wide linkage studies evidenced new regions linked to asthma and atopy. <i>European Journal of Human Genetics</i> , 2010 , 18, 700-6	5.3	44
56	Polymorphisms of PHF11 and DPP10 are associated with asthma and related traits in a Chinese population. <i>Respiration</i> , 2010 , 79, 17-24	3.7	19
55	A genome-wide association study on African-ancestry populations for asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2010 , 125, 336-346.e4	11.5	179
54	PDE11A associations with asthma: results of a genome-wide association scan. <i>Journal of Allergy and Clinical Immunology</i> , 2010 , 126, 871-873.e9	11.5	39
53	Variants of DENND1B associated with asthma in children. <i>New England Journal of Medicine</i> , 2010 , 362, 36-44	59.2	261
52	A large-scale, consortium-based genomewide association study of asthma. <i>New England Journal of Medicine</i> , 2010 , 363, 1211-1221	59.2	1431
51	Gene expression in skin and lymphoblastoid cells: Refined statistical method reveals extensive overlap in cis-eQTL signals. <i>American Journal of Human Genetics</i> , 2010 , 87, 779-89	11	144
50	Disordered microbial communities in asthmatic airways. <i>PLoS ONE</i> , 2010 , 5, e8578	3.7	1085
49	Dynamic and physical clustering of gene expression during epidermal barrier formation in differentiating keratinocytes. <i>PLoS ONE</i> , 2009 , 4, e7651	3.7	23
48	Genome-wide association studies in the genetics of asthma. <i>Current Allergy and Asthma Reports</i> , 2009 , 9, 3-9	5.6	19
47	Mapping complex disease traits with global gene expression. <i>Nature Reviews Genetics</i> , 2009 , 10, 184-94	30.1	658
46	Genome-wide association analysis identifies PDE4D as an asthma-susceptibility gene. <i>American Journal of Human Genetics</i> , 2009 , 84, 581-93	11	264
45	ENU mutagenesis as a tool for understanding lung development and disease. <i>Biochemical Society Transactions</i> , 2009 , 37, 838-42	5.1	10
44	Genome-wide scan on total serum IgE levels identifies FCER1A as novel susceptibility locus. <i>PLoS Genetics</i> , 2008 , 4, e1000166	6	218
43	A genome-wide association study of global gene expression. <i>Nature Genetics</i> , 2007 , 39, 1202-7	36.3	801
42	Filaggrin mutations in children with severe atopic dermatitis. <i>Journal of Investigative Dermatology</i> , 2007 , 127, 1667-72	4.3	160
41	Genetic variants regulating ORMDL3 expression contribute to the risk of childhood asthma. <i>Nature</i> , 2007 , 448, 470-3	50.4	1201

40	Atopic sensitization and the international variation of asthma symptom prevalence in children. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2007 , 176, 565-74	10.2	223
39	Genetic and environmental effects on complex traits in mice. <i>Genetics</i> , 2006 , 174, 959-84	4	134
38	The genetics of atopic dermatitis. <i>Journal of Allergy and Clinical Immunology</i> , 2006 , 118, 24-34; quiz 35-6	511.5	194
37	Genetic variation in the beta subunit of the high affinity IgE receptor and atopy and asthma. <i>Clinical and Experimental Allergy</i> , 2006 , 36, 855-7	4.1	5
36	Genome-wide genetic association of complex traits in heterogeneous stock mice. <i>Nature Genetics</i> , 2006 , 38, 879-87	36.3	442
35	Investigation of the chromosome 17q25 PSORS2 locus in atopic dermatitis. <i>Journal of Investigative Dermatology</i> , 2006 , 126, 603-6	4.3	13
34	A protocol for high-throughput phenotyping, suitable for quantitative trait analysis in mice. <i>Mammalian Genome</i> , 2006 , 17, 129-46	3.2	88
33	Association between a complex insertion/deletion polymorphism in NOD1 (CARD4) and susceptibility to inflammatory bowel disease. <i>Human Molecular Genetics</i> , 2005 , 14, 1245-50	5.6	260
32	Haplotypes and asthma. American Journal of Respiratory and Critical Care Medicine, 2005, 171, 1066-7	10.2	1
31	NOD1 variation, immunoglobulin E and asthma. <i>Human Molecular Genetics</i> , 2005 , 14, 935-41	5.6	219
30	The immunogenetics of asthma and eczema: a new focus on the epithelium. <i>Nature Reviews Immunology</i> , 2004 , 4, 978-88	36.5	309
29	LD mapping of maternally and non-maternally derived alleles and atopy in FcepsilonRI-beta. <i>Human Molecular Genetics</i> , 2003 , 12, 2577-85	5.6	40
28	Benign asbestos pleural diseases. Current Opinion in Pulmonary Medicine, 2003, 9, 266-71	3	46
27	A new gene for asthma: would you ADAM and Eve it?. <i>Trends in Genetics</i> , 2003 , 19, 169-72	8.5	23
26	Positional cloning of a quantitative trait locus on chromosome 13q14 that influences immunoglobulin E levels and asthma. <i>Nature Genetics</i> , 2003 , 34, 181-6	36.3	263
25	Positional cloning of a novel gene influencing asthma from chromosome 2q14. <i>Nature Genetics</i> , 2003 , 35, 258-63	36.3	283
24	Competing functions encoded in the allergy-associated F(c)epsilonRIbeta gene. <i>Immunity</i> , 2003 , 18, 665	- 74 3	59
23	Atopy, respiratory function and HLA-DR in Aboriginal Australians. <i>Human Molecular Genetics</i> , 2003 , 12, 625-30	5.6	5

22	Genetics and genomics of asthma and allergic diseases. <i>Immunological Reviews</i> , 2002 , 190, 195-206	11.3	89
21	Merlinrapid analysis of dense genetic maps using sparse gene flow trees. <i>Nature Genetics</i> , 2002 , 30, 97-101	36.3	2878
20	Positive association to IgE levels and a physical map of the 13q14 atopy locus. <i>European Journal of Human Genetics</i> , 2002 , 10, 266-70	5.3	25
19	A detailed genetic map of the chromosome 7 bronchial hyper-responsiveness locus. <i>European Journal of Human Genetics</i> , 2002 , 10, 177-82	5.3	16
18	Genetic and perinatal risk factors for asthma onset and severity: a review and theoretical analysis. <i>Epidemiologic Reviews</i> , 2002 , 24, 176-89	4.1	58
17	Allergy to dermatophagoides in a group of Spanish gypsies: genetic restrictions. <i>International Archives of Allergy and Immunology</i> , 2001 , 125, 297-306	3.7	17
16	Allergy-associated polymorphisms of the Fc epsilon RI beta subunit do not impact its two amplification functions. <i>Journal of Immunology</i> , 2000 , 165, 3917-22	5.3	53
15	A genome-wide screen for asthma-associated quantitative trait loci in a mouse model of allergic asthma. <i>Human Molecular Genetics</i> , 1999 , 8, 601-5	5.6	59
14	The alliance of genes and environment in asthma and allergy. <i>Nature</i> , 1999 , 402, B5-11	50.4	249
13	Germline TCR-A restriction of immunoglobulin E responses to allergen. <i>Immunogenetics</i> , 1997 , 46, 226-	·3 9 .2	42
13	Germline TCR-A restriction of immunoglobulin E responses to allergen. <i>Immunogenetics</i> , 1997 , 46, 226-Naked DNA: new shots for allergy?. <i>Nature Medicine</i> , 1996 , 2, 515-6	-3 9 .2 50.5	1
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12	Naked DNA: new shots for allergy?. <i>Nature Medicine</i> , 1996 , 2, 515-6	50.5	1
12	Naked DNA: new shots for allergy?. <i>Nature Medicine</i> , 1996 , 2, 515-6 A genome-wide search for quantitative trait loci underlying asthma. <i>Nature</i> , 1996 , 383, 247-50	50.5	1 676
12 11 10	Naked DNA: new shots for allergy?. <i>Nature Medicine</i> , 1996 , 2, 515-6 A genome-wide search for quantitative trait loci underlying asthma. <i>Nature</i> , 1996 , 383, 247-50 Reply to Atopy in Australia <i>Nature Genetics</i> , 1995 , 10, 260-260 Detection of a recessive major gene for high IgE levels acting independently of specific response to	50.5	1 676 5
12 11 10	Naked DNA: new shots for allergy?. <i>Nature Medicine</i> , 1996 , 2, 515-6 A genome-wide search for quantitative trait loci underlying asthma. <i>Nature</i> , 1996 , 383, 247-50 Reply to Atopy in Australia <i>Nature Genetics</i> , 1995 , 10, 260-260 Detection of a recessive major gene for high IgE levels acting independently of specific response to allergens. <i>Genetic Epidemiology</i> , 1995 , 12, 93-105	50.5 50.4 36.3 2.6	1 676 5 42 7
12 11 10 9 8	Naked DNA: new shots for allergy?. <i>Nature Medicine</i> , 1996 , 2, 515-6 A genome-wide search for quantitative trait loci underlying asthma. <i>Nature</i> , 1996 , 383, 247-50 Reply to Atopy in Australia <i>Nature Genetics</i> , 1995 , 10, 260-260 Detection of a recessive major gene for high IgE levels acting independently of specific response to allergens. <i>Genetic Epidemiology</i> , 1995 , 12, 93-105 Atopy: a complex genetic disease. <i>Annals of Medicine</i> , 1994 , 26, 351-3	50.5 50.4 36.3 2.6	1 676 5 42 7

4	The natural history of asbestosis in former crocidolite workers of Wittenoom Gorge. <i>The American Review of Respiratory Disease</i> , 1986 , 133, 994-8	32
3	CTAB extraction of DNA and RNA of respiratory samples for microbial work v1	1
2	Longitudinal assessment of sputum microbiome by sequencing of the 16S rRNA gene in non-CF bronchiectasis patients	2
1	MUC5AC drives COPD exacerbation severity through amplification of virus-induced airway inflammation	3