

# John Kenneth Baillie

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/123901/publications.pdf>

Version: 2024-02-01

135  
papers

35,667  
citations

29994

54  
h-index

12910

131  
g-index

192  
all docs

192  
docs citations

192  
times ranked

61450  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular Patterns in Acute Pancreatitis Reflect Generalizable Endotypes of the Host Response to Systemic Injury in Humans. <i>Annals of Surgery</i> , 2022, 275, e453-e462.	2.1	24
2	Tissue Proteomic Analysis Identifies Mechanisms and Stages of Immunopathology in Fatal COVID-19. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2022, 66, 196-205.	1.4	26
3	SARS-CoV-2 environmental contamination from hospitalised patients with COVID-19 receiving aerosol-generating procedures. <i>Thorax</i> , 2022, 77, 259-267.	2.7	34
4	Multi-ancestry fine mapping implicates OAS1 splicing in risk of severe COVID-19. <i>Nature Genetics</i> , 2022, 54, 125-127.	9.4	75
5	SARS-CoV-2 Omicron-B.1.1.529 leads to widespread escape from neutralizing antibody responses. <i>Cell</i> , 2022, 185, 467-484.e15.	13.5	788
6	Effect of Noninvasive Respiratory Strategies on Intubation or Mortality Among Patients With Acute Hypoxemic Respiratory Failure and COVID-19. <i>JAMA - Journal of the American Medical Association</i> , 2022, 327, 546.	3.8	229
7	A common TMPRSS2 variant has a protective effect against severe COVID-19. <i>Current Research in Translational Medicine</i> , 2022, 70, 103333.	1.2	30
8	Common, low-frequency, rare, and ultra-rare coding variants contribute to COVID-19 severity. <i>Human Genetics</i> , 2022, 141, 147-173.	1.8	22
9	Soluble P-Selectin and von Willebrand Factor Rise in Healthy Volunteers Following Non-exertional Ascent to High Altitude. <i>Frontiers in Physiology</i> , 2022, 13, 825819.	1.3	3
10	Whole-genome sequencing reveals host factors underlying critical COVID-19. <i>Nature</i> , 2022, 607, 97-103.	13.7	174
11	SARS-CoV-2 co-infection with influenza viruses, respiratory syncytial virus, or adenoviruses. <i>Lancet, The</i> , 2022, 399, 1463-1464.	6.3	178
12	Genetic Landscape of the ACE2 Coronavirus Receptor. <i>Circulation</i> , 2022, 145, 1398-1411.	1.6	20
13	Prospective validation of the 4C prognostic models for adults hospitalised with COVID-19 using the ISARIC WHO Clinical Characterisation Protocol. <i>Thorax</i> , 2022, 77, 606-615.	2.7	24
14	Distinct clinical symptom patterns in patients hospitalised with COVID-19 in an analysis of 59,011 patients in the ISARIC-4C study. <i>Scientific Reports</i> , 2022, 12, 6843.	1.6	12
15	Analysis of SARS-CoV-2 in Nasopharyngeal Samples from Patients with COVID-19 Illustrates Population Variation and Diverse Phenotypes, Placing the Growth Properties of Variants of Concern in Context with Other Lineages. <i>MSphere</i> , 2022, 7, e0091321.	1.3	8
16	Procalcitonin Is Not a Reliable Biomarker of Bacterial Coinfection in People With Coronavirus Disease 2019 Undergoing Microbiological Investigation at the Time of Hospital Admission. <i>Open Forum Infectious Diseases</i> , 2022, 9, ofac179.	0.4	10
17	Analysis of SARS-CoV-2 known and novel subgenomic mRNAs in cell culture, animal model, and clinical samples using LeTRS, a bioinformatic tool to identify unique sequence identifiers. <i>GigaScience</i> , 2022, 11, .	3.3	8
18	Fatal COVID-19 outcomes are associated with an antibody response targeting epitopes shared with endemic coronaviruses. <i>JCI Insight</i> , 2022, 7, .	2.3	24

#	ARTICLE	IF	CITATIONS
19	Hypoxia shapes the immune landscape in lung injury and promotes the persistence of inflammation. <i>Nature Immunology</i> , 2022, 23, 927-939.	7.0	21
20	Multiomic analysis reveals cell-type-specific molecular determinants of COVID-19 severity. <i>Cell Systems</i> , 2022, 13, 598-614.e6.	2.9	10
21	Mapping of SARS-CoV-2 IgM and IgG in gingival crevicular fluid: Antibody dynamics and linkage to severity of COVID-19 in hospital inpatients. <i>Journal of Infection</i> , 2022, 85, 152-160.	1.7	6
22	Redefining critical illness. <i>Nature Medicine</i> , 2022, 28, 1141-1148.	15.2	136
23	Clonal hematopoiesis is not significantly associated with COVID-19 disease severity. <i>Blood</i> , 2022, 140, 1650-1655.	0.6	10
24	Dexamethasone in Hospitalized Patients with Covid-19. <i>New England Journal of Medicine</i> , 2021, 384, 693-704.	13.9	8,063
25	Tissue-Specific Immunopathology in Fatal COVID-19. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021, 203, 192-201.	2.5	243
26	Sepsis Subclasses: A Framework for Development and Interpretation*. <i>Critical Care Medicine</i> , 2021, 49, 748-759.	0.4	81
27	A haemagglutination test for rapid detection of antibodies to SARS-CoV-2. <i>Nature Communications</i> , 2021, 12, 1951.	5.8	54
28	Inflammatory profiles across the spectrum of disease reveal a distinct role for GM-CSF in severe COVID-19. <i>Science Immunology</i> , 2021, 6, .	5.6	161
29	Obesity, Ethnicity, and Risk of Critical Care, Mechanical Ventilation, and Mortality in Patients Admitted to Hospital with COVID-19: Analysis of the ISARIC CCP-UK Cohort. <i>Obesity</i> , 2021, 29, 1223-1230.	1.5	34
30	Two Key Takeaways From a Year of Pandemic Research. <i>Critical Care Medicine</i> , 2021, Publish Ahead of Print, 1223-1226.	0.4	1
31	Development and validation of the ISARIC 4C Deterioration model for adults hospitalised with COVID-19: a prospective cohort study. <i>Lancet Respiratory Medicine</i> , 2021, 9, 349-359.	5.2	161
32	COVID-19 symptoms at hospital admission vary with age and sex: results from the ISARIC prospective multinational observational study. <i>Infection</i> , 2021, 49, 889-905.	2.3	62
33	Discovery of widespread transcription initiation at microsatellites predictable by sequence-based deep neural network. <i>Nature Communications</i> , 2021, 12, 3297.	5.8	11
34	COVID-19 pneumothorax in the UK: a prospective observational study using the ISARIC WHO clinical characterisation protocol. <i>European Respiratory Journal</i> , 2021, 58, 2100929.	3.1	21
35	Changes in in-hospital mortality in the first wave of COVID-19: a multicentre prospective observational cohort study using the WHO Clinical Characterisation Protocol UK. <i>Lancet Respiratory Medicine</i> , 2021, 9, 773-785.	5.2	78
36	Characterisation of in-hospital complications associated with COVID-19 using the ISARIC WHO Clinical Characterisation Protocol UK: a prospective, multicentre cohort study. <i>Lancet</i> , 2021, 398, 223-237.	6.3	110

#	ARTICLE	IF	CITATIONS
37	Global infectious disease research collaborations in crises: building capacity and inclusivity through cooperation. <i>Globalization and Health</i> , 2021, 17, 84.	2.4	11
38	Obesity, chronic disease, age, and in-hospital mortality in patients with covid-19: analysis of ISARIC clinical characterisation protocol UK cohort. <i>BMC Infectious Diseases</i> , 2021, 21, 717.	1.3	19
39	Pan-ancestry exome-wide association analyses of COVID-19 outcomes in 586,157 individuals. <i>American Journal of Human Genetics</i> , 2021, 108, 1350-1355.	2.6	72
40	Mapping the human genetic architecture of COVID-19. <i>Nature</i> , 2021, 600, 472-477.	13.7	640
41	Endomembrane targeting of human OAS1 p46 augments antiviral activity. <i>ELife</i> , 2021, 10, .	2.8	41
42	Hospital-acquired SARS-CoV-2 infection in the UK's first COVID-19 pandemic wave. <i>Lancet, The</i> , 2021, 398, 1037-1038.	6.3	75
43	Understanding the burden of interstitial lung disease post-COVID-19: the UK Interstitial Lung Disease-Long COVID Study (UKILD-Long COVID). <i>BMJ Open Respiratory Research</i> , 2021, 8, e001049.	1.2	28
44	A prenylated dsRNA sensor protects against severe COVID-19. <i>Science</i> , 2021, 374, eabj3624.	6.0	124
45	Genetic mechanisms of critical illness in COVID-19. <i>Nature</i> , 2021, 591, 92-98.	13.7	1,014
46	Histological Evidence of Pulmonary Microthrombosis and Vasculitis in Life-Threatening Respiratory Virus Diseases. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofaa640.	0.4	6
47	Subtractive CRISPR screen identifies the ATG16L1/vacuolar ATPase axis as required for non-canonical LC3 lipidation. <i>Cell Reports</i> , 2021, 37, 109899.	2.9	33
48	Vitamin D insufficiency in COVID-19 and influenza A, and critical illness survivors: a cross-sectional study. <i>BMJ Open</i> , 2021, 11, e055435.	0.8	10
49	The value of open-source clinical science in pandemic response: lessons from ISARIC. <i>Lancet Infectious Diseases, The</i> , 2021, 21, 1623-1624.	4.6	21
50	Genome-wide CRISPR screen identifies host dependency factors for influenza A virus infection. <i>Nature Communications</i> , 2020, 11, 164.	5.8	136
51	Hypoxia Modulates Platelet Purinergic Signalling Pathways. <i>Thrombosis and Haemostasis</i> , 2020, 120, 253-261.	1.8	12
52	Outcome of Hospitalization for COVID-19 in Patients with Interstitial Lung Disease. An International Multicenter Study. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 202, 1656-1665.	2.5	171
53	Effect of Hydroxychloroquine in Hospitalized Patients with Covid-19. <i>New England Journal of Medicine</i> , 2020, 383, 2030-2040.	13.9	1,013
54	RECOVERY- Respiratory Support: Respiratory Strategies for patients with suspected or proven COVID-19 respiratory failure; Continuous Positive Airway Pressure, High-flow Nasal Oxygen, and standard care: A structured summary of a study protocol for a randomised controlled trial. <i>Trials</i> , 2020, 21, 687.	0.7	28

#	ARTICLE	IF	CITATIONS
55	Functional annotation of human long noncoding RNAs via molecular phenotyping. <i>Genome Research</i> , 2020, 30, 1060-1072.	2.4	109
56	Amplicon-Based Detection and Sequencing of SARS-CoV-2 in Nasopharyngeal Swabs from Patients With COVID-19 and Identification of Deletions in the Viral Genome That Encode Proteins Involved in Interferon Antagonism. <i>Viruses</i> , 2020, 12, 1164.	1.5	51
57	Effect of Hydrocortisone on Mortality and Organ Support in Patients With Severe COVID-19. <i>JAMA - Journal of the American Medical Association</i> , 2020, 324, 1317.	3.8	671
58	Broad and strong memory CD4+ and CD8+ T cells induced by SARS-CoV-2 in UK convalescent individuals following COVID-19. <i>Nature Immunology</i> , 2020, 21, 1336-1345.	7.0	1,066
59	Performance characteristics of five immunoassays for SARS-CoV-2: a head-to-head benchmark comparison. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 1390-1400.	4.6	336
60	Risk stratification of patients admitted to hospital with covid-19 using the ISARIC WHO Clinical Characterisation Protocol: development and validation of the 4C Mortality Score. <i>BMJ</i> , The, 2020, 370, m3339.	3.0	779
61	Comprehensive Characterization of Transcriptional Activity during Influenza A Virus Infection Reveals Biases in Cap-Snatching of Host RNA Sequences. <i>Journal of Virology</i> , 2020, 94, .	1.5	14
62	Hybrid Gene Origination Creates Human-Virus Chimeric Proteins during Infection. <i>Cell</i> , 2020, 181, 1502-1517.e23.	13.5	33
63	Global outbreak research: harmony not hegemony. <i>Lancet Infectious Diseases</i> , The, 2020, 20, 770-772.	4.6	40
64	Developing Novel Host-Based Therapies Targeting Microbicidal Responses in Macrophages and Neutrophils to Combat Bacterial Antimicrobial Resistance. <i>Frontiers in Immunology</i> , 2020, 11, 786.	2.2	10
65	Using imaging to combat a pandemic: rationale for developing the UK National COVID-19 Chest Imaging Database. <i>European Respiratory Journal</i> , 2020, 56, 2001809.	3.1	24
66	Linking protein to phenotype with Mendelian Randomization detects 38 proteins with causal roles in human diseases and traits. <i>PLoS Genetics</i> , 2020, 16, e1008785.	1.5	29
67	Clinical evidence does not support corticosteroid treatment for 2019-nCoV lung injury. <i>Lancet</i> , The, 2020, 395, 473-475.	6.3	1,644
68	Dynamic data-driven meta-analysis for prioritisation of host genes implicated in COVID-19. <i>Scientific Reports</i> , 2020, 10, 22303.	1.6	31
69	Features of 20,133 UK patients in hospital with covid-19 using the ISARIC WHO Clinical Characterisation Protocol: prospective observational cohort study. <i>BMJ</i> , The, 2020, 369, m1985.	3.0	2,474
70	Antibody testing for COVID-19: A report from the National COVID Scientific Advisory Panel. <i>Wellcome Open Research</i> , 2020, 5, 139.	0.9	179
71	An evidence-based framework for priority clinical research questions for COVID-19. <i>Journal of Global Health</i> , 2020, 10, .	1.2	22
72	Letter to the Editor: Response to Luke Fletcher and Philip Peyton regarding "Predictive validity of a novel non-invasive estimation of effective shunt fraction in critically ill patients". <i>Intensive Care Medicine Experimental</i> , 2020, 8, 57.	0.9	0

#	ARTICLE	IF	CITATIONS
73	An evidence-based framework for priority clinical research questions for COVID-19. <i>Journal of Global Health</i> , 2020, 10, 011001.	1.2	11
74	Predictive validity of a novel non-invasive estimation of effective shunt fraction in critically ill patients. <i>Intensive Care Medicine Experimental</i> , 2019, 7, 49.	0.9	6
75	Host susceptibility to severe influenza A virus infection. <i>Critical Care</i> , 2019, 23, 303.	2.5	73
76	The utility of peripheral blood leucocyte ratios as biomarkers in infectious diseases: A systematic review and meta-analysis. <i>Journal of Infection</i> , 2019, 78, 339-348.	1.7	131
77	The effect of dexmedetomidine and clonidine on the inflammatory response in critical illness: a systematic review of animal and human studies. <i>Critical Care</i> , 2019, 23, 402.	2.5	42
78	The Search for Efficacious New Therapies in Sepsis Needs to Embrace Heterogeneity. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019, 199, 936-938.	2.5	17
79	Thromboelastometry and Platelet Function during Acclimatization to High Altitude. <i>Thrombosis and Haemostasis</i> , 2018, 118, 063-071.	1.8	30
80	The 2018 Lake Louise Acute Mountain Sickness Score. <i>High Altitude Medicine and Biology</i> , 2018, 19, 4-6.	0.5	324
81	Shared activity patterns arising at genetic susceptibility loci reveal underlying genomic and cellular architecture of human disease. <i>PLoS Computational Biology</i> , 2018, 14, e1005934.	1.5	17
82	Middle East Respiratory Syndrome. <i>New England Journal of Medicine</i> , 2017, 376, 584-594.	13.9	351
83	Treatable traits and therapeutic targets: Goals for systems biology in infectious disease. <i>Current Opinion in Systems Biology</i> , 2017, 2, 140-146.	1.3	37
84	FANTOM5 CAGE profiles of human and mouse samples. <i>Scientific Data</i> , 2017, 4, 170112.	2.4	195
85	Analysis of the human monocyte-derived macrophage transcriptome and response to lipopolysaccharide provides new insights into genetic aetiology of inflammatory bowel disease. <i>PLoS Genetics</i> , 2017, 13, e1006641.	1.5	161
86	Microglial brain region-dependent diversity and selective regional sensitivities to aging. <i>Nature Neuroscience</i> , 2016, 19, 504-516.	7.1	919
87	Transcribed enhancers lead waves of coordinated transcription in transitioning mammalian cells. <i>Science</i> , 2015, 347, 1010-1014.	6.0	517
88	Gateways to the FANTOM5 promoter level mammalian expression atlas. <i>Genome Biology</i> , 2015, 16, 22.	3.8	687
89	The Effectiveness of Convalescent Plasma and Hyperimmune Immunoglobulin for the Treatment of Severe Acute Respiratory Infections of Viral Etiology: A Systematic Review and Exploratory Meta-analysis. <i>Journal of Infectious Diseases</i> , 2015, 211, 80-90.	1.9	841
90	Open source clinical science for emerging infections. <i>Lancet Infectious Diseases</i> , The, 2014, 14, 8-9.	4.6	82

#	ARTICLE	IF	CITATIONS
91	A promoter-level mammalian expression atlas. <i>Nature</i> , 2014, 507, 462-470.	13.7	1,838
92	An atlas of active enhancers across human cell types and tissues. <i>Nature</i> , 2014, 507, 455-461.	13.7	2,269
93	Targeting the host immune response to fight infection. <i>Science</i> , 2014, 344, 807-808.	6.0	30
94	Antiviral combinations for severe influenza. <i>Lancet Infectious Diseases</i> , The, 2014, 14, 1259-1270.	4.6	159
95	Stratified infection medicine: a call to arms. <i>Lancet Infectious Diseases</i> , The, 2014, 14, 451.	4.6	2
96	Transcriptional profiling of the human fibrillin/LTBP gene family, key regulators of mesenchymal cell functions. <i>Molecular Genetics and Metabolism</i> , 2014, 112, 73-83.	0.5	39
97	Network Analysis Reveals Distinct Clinical Syndromes Underlying Acute Mountain Sickness. <i>PLoS ONE</i> , 2014, 9, e81229.	1.1	48
98	Influenza – Time to Target the Host?. <i>New England Journal of Medicine</i> , 2013, 369, 191-193.	13.9	50
99	Complete genome sequence of methicillin-sensitive <i>Staphylococcus aureus</i> containing a heterogeneous staphylococcal cassette chromosome element. <i>Science China Life Sciences</i> , 2013, 56, 268-274.	2.3	0
100	An expression atlas of human primary cells: inference of gene function from coexpression networks. <i>BMC Genomics</i> , 2013, 14, 632.	1.2	347
101	Wide variation and systematic bias in expert clinicians' perceptions of prognosis following brain injury. <i>British Journal of Neurosurgery</i> , 2013, 27, 340-343.	0.4	25
102	Endogenous Retrotransposition Activates Oncogenic Pathways in Hepatocellular Carcinoma. <i>Cell</i> , 2013, 153, 101-111.	13.5	352
103	Deciding authorship order. <i>BMJ</i> , The, 2013, 347, f7182-f7182.	3.0	6
104	An updated systematic review of the role of host genetics in susceptibility to influenza. <i>Influenza and Other Respiratory Viruses</i> , 2013, 7, 37-41.	1.5	39
105	Modelling mutational and selection pressures on dinucleotides in eukaryotic phyla – selection against CpG and UpA in cytoplasmically expressed RNA and in RNA viruses. <i>BMC Genomics</i> , 2013, 14, 610.	1.2	71
106	Identification of Novel Genes Selectively Expressed in the Follicle-Associated Epithelium from the Meta-Analysis of Transcriptomics Data from Multiple Mouse Cell and Tissue Populations. <i>DNA Research</i> , 2012, 19, 407-422.	1.5	17
107	Retinal Vessel Tortuosity in Response to Hypobaric Hypoxia. <i>High Altitude Medicine and Biology</i> , 2012, 13, 263-268.	0.5	12
108	IFITM3 restricts the morbidity and mortality associated with influenza. <i>Nature</i> , 2012, 484, 519-523.	13.7	668

#	ARTICLE	IF	CITATIONS
109	A gene expression atlas of the domestic pig. BMC Biology, 2012, 10, 90.	1.7	199
110	Conservation and divergence in Toll-like receptor 4-regulated gene expression in primary human versus mouse macrophages. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, E944-53.	3.3	332
111	High altitude adaptation in Daghestani populations from the Caucasus. Human Genetics, 2012, 131, 423-433.	1.8	31
112	The Role of Host Genetics in Susceptibility to Influenza: A Systematic Review. PLoS ONE, 2012, 7, e33180.	1.1	98
113	Sildenafil Citrate for the Prevention of High Altitude Hypoxic Pulmonary Hypertension: Double Blind, Randomized, Placebo-Controlled Trial. High Altitude Medicine and Biology, 2011, 12, 207-214.	0.5	32
114	Expression of mesenchyme-specific gene signatures by follicular dendritic cells: insights from the meta-analysis of microarray data from multiple mouse cell populations. Immunology, 2011, 133, 482-498.	2.0	50
115	Somatic retrotransposition alters the genetic landscape of the human brain. Nature, 2011, 479, 534-537.	13.7	621
116	High Altitude Pulmonary Oedema. Journal of the Royal Army Medical Corps, 2011, 157, 68-72.	0.8	25
117	Is somatic retrotransposition a parasitic or symbiotic phenomenon?. Mobile Genetic Elements, 2011, 1, 279-328.	1.8	14
118	Passage of pathogenic microorganisms through breathing system filters used in anaesthesia and intensive care. Anaesthesia, 2010, 65, 670-673.	1.8	10
119	Meta-analysis of lineage-specific gene expression signatures in mouse leukocyte populations. Immunobiology, 2010, 215, 724-736.	0.8	81
120	Functional clustering and lineage markers: Insights into cellular differentiation and gene function from large-scale microarray studies of purified primary cell populations. Genomics, 2010, 95, 328-338.	1.3	112
121	Incidence and predictors of acute mountain sickness among trekkers on Mount Kilimanjaro. High Altitude Medicine and Biology, 2010, 11, 217-222.	0.5	58
122	Oral antioxidant supplementation does not prevent acute mountain sickness: double blind, randomized placebo-controlled trial. QJM - Monthly Journal of the Association of Physicians, 2009, 102, 341-348.	0.2	85
123	The citric acid cough threshold and the ventilatory response to carbon dioxide on ascent to high altitude. Respiratory Medicine, 2009, 103, 1182-1188.	1.3	5
124	Measurement errors caused by incompatibility of apparatus. European Journal of Anaesthesiology, 2009, 26, 977-980.	0.7	0
125	Simple, easily memorised "rules of thumb" for the rapid assessment of physiological compensation for respiratory acid-base disorders. Thorax, 2008, 63, 289-290.	2.7	6
126	NT-proBNP Does Not Rise on Acute Ascent to High Altitude. High Altitude Medicine and Biology, 2008, 9, 307-310.	0.5	20

#	ARTICLE	IF	CITATIONS
127	Endogenous Urate Production Augments Plasma Antioxidant Capacity in Healthy Lowland Subjects Exposed to High Altitude. <i>Chest</i> , 2007, 131, 1473-1478.	0.4	52
128	Change in plasma vascular endothelial growth factor during onset and recovery from acute mountain sickness. <i>Respiratory Medicine</i> , 2007, 101, 587-594.	1.3	39
129	Contamination of anaesthetic machines with pathogenic organisms*. <i>Anaesthesia</i> , 2007, 62, 1257-1261.	1.8	20
130	Activated protein C: controversy and hope in the treatment of sepsis. <i>Current Opinion in Investigational Drugs</i> , 2007, 8, 933-8.	2.3	3
131	Endothelial Nitric Oxide Synthase Polymorphisms Do Not Influence Pulmonary Artery Systolic Pressure at Altitude. <i>High Altitude Medicine and Biology</i> , 2006, 7, 221-227.	0.5	19
132	The mechanism of action of gabapentin in neuropathic pain. <i>Current Opinion in Investigational Drugs</i> , 2006, 7, 33-9.	2.3	26
133	Pericardial effusions in healthy lowlanders after acute ascent to high altitude. <i>Heart</i> , 2005, 92, 539-540.	1.2	6
134	Morphine, Gabapentin, or Their Combination for Neuropathic Pain. <i>New England Journal of Medicine</i> , 2005, 352, 2650-2651.	13.9	20
135	A Common <i>TMPRSS2</i> Variant Protects Against Severe COVID-19. <i>SSRN Electronic Journal</i> , 0, , .	0.4	2