

Phuwanat Sakornsakolpat

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.

25
papers

591
citations

9
h-index

24
g-index

31
ext. papers

1,063
ext. citations

9.3
avg, IF

2.51
L-index

#	Paper	IF	Citations
25	Lung tissue shows divergent gene expression between chronic obstructive pulmonary disease and idiopathic pulmonary fibrosis.. <i>Respiratory Research</i> , 2022 , 23, 97	7.3	0
24	Emphysema Progression and Lung Function Decline Among Angiotensin Converting Enzyme Inhibitors and Angiotensin-Receptor Blockade Users in the COPDGene Cohort. <i>Chest</i> , 2021 , 160, 1245-1254	5.3	0
23	A systematic analysis of protein-altering exonic variants in chronic obstructive pulmonary disease. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2021 , 321, L130-L143	5.8	2
22	Genome-Wide Gene-by-Smoking Interaction Study of Chronic Obstructive Pulmonary Disease. <i>American Journal of Epidemiology</i> , 2021 , 190, 875-885	3.8	9
21	A polygenic risk score for asthma in a large racially diverse population. <i>Clinical and Experimental Allergy</i> , 2021 , 51, 1410-1420	4.1	1
20	Interaction of Cigarette Smoking and Polygenic Risk Score on Reduced Lung Function.. <i>JAMA Network Open</i> , 2021 , 4, e2139525	10.4	3
19	Relative contributions of family history and a polygenic risk score on COPD and related outcomes: COPDGene and ECLIPSE studies. <i>BMJ Open Respiratory Research</i> , 2020 , 7,	5.6	2
18	Adjunctive favipiravir for severe COVID-19: a retrospective observational study of the first 41 patients in Thailand. <i>Asian Biomedicine</i> , 2020 , 14, 107-110	0.4	1
17	Genome-Wide Association Study of Susceptibility to Idiopathic Pulmonary Fibrosis. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020 , 201, 564-574	10.2	81
16	Disease Progression Modeling in Chronic Obstructive Pulmonary Disease. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020 , 201, 294-302	10.2	20
15	Machine Learning Characterization of COPD Subtypes: Insights From the COPDGene Study. <i>Chest</i> , 2020 , 157, 1147-1157	5.3	18
14	Whole genome sequence analysis of pulmonary function and COPD in 19,996 multi-ethnic participants. <i>Nature Communications</i> , 2020 , 11, 5182	17.4	6
13	Chronic obstructive pulmonary disease and related phenotypes: polygenic risk scores in population-based and case-control cohorts. <i>Lancet Respiratory Medicine</i> , 2020 , 8, 696-708	35.1	29
12	Whole Genome Sequencing Identifies CRISPLD2 as a Lung Function Gene in Children With Asthma. <i>Chest</i> , 2019 , 156, 1068-1079	5.3	3
11	Genome-Wide Association Analysis of Single-Breath DL. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2019 , 60, 523-531	5.7	4
10	New genetic signals for lung function highlight pathways and chronic obstructive pulmonary disease associations across multiple ancestries. <i>Nature Genetics</i> , 2019 , 51, 481-493	36.3	156
9	Genetic landscape of chronic obstructive pulmonary disease identifies heterogeneous cell-type and phenotype associations. <i>Nature Genetics</i> , 2019 , 51, 494-505	36.3	119

8	DSP variants may be associated with longitudinal change in quantitative emphysema. <i>Respiratory Research</i> , 2019 , 20, 160	7.3	4
7	Analysis of genetically driven alternative splicing identifies FBXO38 as a novel COPD susceptibility gene. <i>PLoS Genetics</i> , 2019 , 15, e1008229	6	9
6	GWAS and systems biology analysis of depressive symptoms among smokers from the COPDGene cohort. <i>Journal of Affective Disorders</i> , 2019 , 243, 16-22	6.6	7
5	Integrative Genomics Analysis Identifies ACVR1B as a Candidate Causal Gene of Emphysema Distribution. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2019 , 60, 388-398	5.7	9
4	Leveraging lung tissue transcriptome to uncover candidate causal genes in COPD genetic associations. <i>Human Molecular Genetics</i> , 2018 , 27, 1819-1829	5.6	24
3	Integrative genomics identifies new genes associated with severe COPD and emphysema. <i>Respiratory Research</i> , 2018 , 19, 46	7.3	9
2	Longitudinal Phenotypes and Mortality in Preserved Ratio Impaired Spirometry in the COPDGene Study. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2018 , 198, 1397-1405	10.2	59
1	Whole-Genome Sequencing in Severe Chronic Obstructive Pulmonary Disease. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2018 , 59, 614-622	5.7	14