## Wouter A A De Steenhuijsen Piters

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2942890/publications.pdf

Version: 2024-02-01

26 papers 3,010 citations

430874 18 h-index 25 g-index

27 all docs

27 docs citations

27 times ranked

4138 citing authors

#	Article	IF	Citations
1	The microbiota of the respiratory tract: gatekeeper to respiratory health. Nature Reviews Microbiology, 2017, 15, 259-270.	28.6	829
2	Nasopharyngeal Microbiota, Host Transcriptome, and Disease Severity in Children with Respiratory Syncytial Virus Infection. American Journal of Respiratory and Critical Care Medicine, 2016, 194, 1104-1115.	5.6	337
3	Maturation of the Infant Respiratory Microbiota, Environmental Drivers, and Health Consequences. A Prospective Cohort Study. American Journal of Respiratory and Critical Care Medicine, 2017, 196, 1582-1590.	5.6	237
4	The role of the local microbial ecosystem in respiratory health and disease. Philosophical Transactions of the Royal Society B: Biological Sciences, 2015, 370, 20140294.	4.0	215
5	Development of Upper Respiratory Tract Microbiota in Infancy is Affected by Mode of Delivery. EBioMedicine, 2016, 9, 336-345.	6.1	194
6	Associations between Pathogens in the Upper Respiratory Tract of Young Children: Interplay between Viruses and Bacteria. PLoS ONE, 2012, 7, e47711.	2.5	177
7	Dysbiosis of upper respiratory tract microbiota in elderly pneumonia patients. ISME Journal, 2016, 10, 97-108.	9.8	166
8	Inflammation induced by influenza virus impairs human innate immune control of pneumococcus. Nature Immunology, 2018, 19, 1299-1308.	14.5	127
9	Development of the Nasopharyngeal Microbiota in Infants with Cystic Fibrosis. American Journal of Respiratory and Critical Care Medicine, 2016, 193, 504-515.	5.6	112
10	Rhinovirus Detection in Symptomatic and Asymptomatic Children: Value of Host Transcriptome Analysis. American Journal of Respiratory and Critical Care Medicine, 2016, 193, 772-782.	5.6	98
11	Comparative gut microbiota and resistome profiling of intensive care patients receiving selective digestive tract decontamination and healthy subjects. Microbiome, 2017, 5, 88.	11.1	90
12	Early Life Microbiota and Respiratory Tract Infections. Cell Host and Microbe, 2020, 28, 223-232.	11.0	61
13	Interaction between the nasal microbiota and S. pneumoniae in the context of live-attenuated influenza vaccine. Nature Communications, 2019, 10, 2981.	12.8	59
14	Concordance between upper and lower airway microbiota in infants with cystic fibrosis. European Respiratory Journal, 2017, 49, 1602235.	6.7	57
15	Loss of Microbial Topography between Oral and Nasopharyngeal Microbiota and Development of Respiratory Infections Early in Life. American Journal of Respiratory and Critical Care Medicine, 2019, 200, 760-770.	5.6	55
16	Seasonal and meteorological determinants of aneurysmal subarachnoid hemorrhage: a systematic review and meta-analysis. Journal of Neurology, 2013, 260, 614-619.	3.6	44
17	Increased risk of pneumonia in residents living near poultry farms: does the upper respiratory tract microbiota play a role?. Pneumonia (Nathan Qld ), 2017, 9, 3.	6.1	40
18	Two Randomized Trials of the Effect of Live Attenuated Influenza Vaccine on Pneumococcal Colonization. American Journal of Respiratory and Critical Care Medicine, 2019, 199, 1160-1163.	5.6	27

#	Article	IF	Citations
19	Early-life viral infections are associated with disadvantageous immune and microbiota profiles and recurrent respiratory infections. Nature Microbiology, 2022, 7, 224-237.	13.3	25
20	Early origins of lung disease: towards an interdisciplinary approach. European Respiratory Review, 2020, 29, 200191.	7.1	21
21	Unraveling the Molecular Mechanisms Underlying the Nasopharyngeal Bacterial Community Structure. MBio, 2016, 7, e00009-16.	4.1	11
22	Benchmarking laboratory processes to characterise low-biomass respiratory microbiota. Scientific Reports, 2021, 11, 17148.	3.3	10
23	Decreased production of epithelial-derived antimicrobial molecules at mucosal barriers during early life. Mucosal Immunology, 2021, 14, 1358-1368.	6.0	9
24	Bacterial and fungal communities in tracheal aspirates of intubated COVID-19 patients: a pilot study. Scientific Reports, 2022, 12, .	3.3	2
25	Bacterial DNA in Fetal Lung Samples May Be Explained by Sample Contamination. American Journal of Respiratory and Critical Care Medicine, 2020, 201, 1310-1311.	5.6	1
26	What is the diagnosis?. Medical Education, 2014, 48, 1114-1115.	2.1	0