Kim de Jong

List of Publications by Year
in descending order

40 papers	1,612 citations	394286 19 h-index	330025 37 g-index
40 all docs	40 docs citations	40 times ranked	3727 citing authors

#	Article	IF	Citations
1	Embryologically Based Classification Specifies Gender Differences in the Prevalence of Orofacial Cleft Subphenotypes. Cleft Palate-Craniofacial Journal, 2021, 58, 54-60.	0.5	12
2	Seasonal Influence on the Numbers of Gender-Related Orofacial Cleft Conceptions in the Netherlands. Cleft Palate-Craniofacial Journal, 2021, 58, 1422-1429.	0.5	1
3	Targeting dynamic hyperinflation in moderate-to-severe asthma: a randomised controlled trial. ERJ Open Research, 2021, 7, 00738-2020.	1.1	4
4	A prognostic model for the preoperative identification of patients at risk for receiving transfusion of packed red blood cells in cardiac surgery. Transfusion, 2021, 61, 2336-2346.	0.8	3
5	Occupational exposures and genetic susceptibility to occupational exposures are related to sickness absence in the Lifelines cohort study. Scientific Reports, 2020, 10, 12963.	1.6	3
6	An age-adjusted instruction video enhances the correct use of nasal corticosteroid sprays in children. Allergologia Et Immunopathologia, 2020, 48, 465-468.	1.0	3
7	Analysis of genetically driven alternative splicing identifies FBXO38 as a novel COPD susceptibility gene. PLoS Genetics, 2019, 15, e1008229.	1.5	17
8	Epigenome-wide association study of lung function level and its change. European Respiratory Journal, 2019, 54, 1900457.	3.1	49
9	Dynamic hyperinflation impairs daily lifeÂactivity in asthma. European Respiratory Journal, 2019, 53, 1801500.	3.1	21
10	Occupational exposure to gases/fumes and mineral dust affect DNA methylation levels of genes regulating expression. Human Molecular Genetics, 2019, 28, 2477-2485.	1.4	9
11	Limited overlap in significant hits between genome-wide association studies on two airflow obstruction definitions in the same population. BMC Pulmonary Medicine, 2019, 19, 58.	0.8	4
12	Genetic landscape of chronic obstructive pulmonary disease identifies heterogeneous cell-type and phenotype associations. Nature Genetics, 2019, 51, 494-505.	9.4	257
13	The relationship between burnout, personality traits, and medical specialty. A national study among Dutch residents. Medical Teacher, 2019, 41, 584-590.	1.0	37
14	Genome-wide interaction study of gene-by-occupational exposures on respiratory symptoms. Environment International, 2019, 122, 263-269.	4.8	17
15	Corticosteroid phobia (corticophobia) in parents of young children with atopic dermatitis and their health care providers. Pediatric Dermatology, 2019, 36, 100-104.	0.5	43
16	COPD GWAS variant at 19q13.2 in relation with DNA methylation and gene expression. Human Molecular Genetics, 2018, 27, 396-405.	1.4	24
17	Understanding the role of the chromosome 15q25.1 in COPD through epigenetics and transcriptomics. European Journal of Human Genetics, 2018, 26, 709-722.	1.4	21
18	Leveraging lung tissue transcriptome to uncover candidate causal genes in COPD genetic associations. Human Molecular Genetics, 2018, 27, 1819-1829.	1.4	37

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19	Whole specimen intraoperative frozen section analysis. Experience with 1082 basal cell carcinomas. European Journal of Surgical Oncology, 2018, 44, 157-162.	0.5	4
20	Long-term Air Pollution Exposure, Genome-wide DNA Methylation and Lung Function in the LifeLines Cohort Study. Environmental Health Perspectives, 2018, 126, 027004.	2.8	71
21	Occupational exposure to pesticides is associated with differential DNA methylation. Occupational and Environmental Medicine, 2018, 75, 427-435.	1.3	61
22	Multiethnic meta-analysis identifies ancestry-specific and cross-ancestry loci for pulmonary function. Nature Communications, 2018, 9, 2976.	5.8	85
23	Evidence for large-scale gene-by-smoking interaction effects on pulmonary function. International Journal of Epidemiology, 2017, 46, dyw318.	0.9	36
24	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
25	Sulfatase modifying factor 1 (SUMF1) is associated with Chronic Obstructive Pulmonary Disease. Respiratory Research, 2017, 18, 77.	1.4	9
26	Surfactant protein D is a causal risk factor for COPD: results of Mendelian randomisation. European Respiratory Journal, 2017, 50, 1700657.	3.1	45
27	No convincing association between genetic markers and respiratory symptoms: results of a GWA study. Respiratory Research, 2017, 18, 11.	1.4	5
28	Genome-wide association study on the FEV 1 /FVC ratio in never-smokers identifies HHIP and FAM13A. Journal of Allergy and Clinical Immunology, 2017, 139, 533-540.	1.5	45
29	Genes and pathways underlying susceptibility to impaired lung function in the context of environmental tobacco smoke exposure. Respiratory Research, 2017, 18, 142.	1.4	16
30	Air pollution exposure is associated with restrictive ventilatory patterns. European Respiratory Journal, 2016, 48, 1221-1224.	3.1	19
31	Novel Genetic Susceptibility Loci for FEV₁ in the Context of Occupational Exposure in Never-Smokers. American Journal of Respiratory and Critical Care Medicine, 2016, 194, 769-772.	2.5	1
32	The Well-Known Gene HHIP and Novel Gene MECR Are Implicated in Small Airway Obstruction. American Journal of Respiratory and Critical Care Medicine, 2016, 194, 1299-1302.	2.5	11
33	Genome-wide interaction study of gene-by-occupational exposure and effects on FEV1 levels. Journal of Allergy and Clinical Immunology, 2015, 136, 1664-1672.e14.	1.5	34
34	Association of Occupational Pesticide Exposure With Accelerated Longitudinal Decline in Lung Function. American Journal of Epidemiology, 2014, 179, 1323-1330.	1.6	45
35	Occupational Exposure to Vapors, Gases, Dusts, and Fumes Is Associated with Small Airways Obstruction. American Journal of Respiratory and Critical Care Medicine, 2014, 189, 487-490.	2.5	21
36	Risk factors for chronic mucus hypersecretion in individuals with and without COPD: influence of smoking and job exposure on CMH. Occupational and Environmental Medicine, 2014, 71, 346-352.	1.3	17

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37	Genome-wide association analysis identifies six new loci associated with forced vital capacity. Nature Genetics, 2014, 46, 669-677.	9.4	131
38	GST-omega genes interact with environmental tobacco smoke on adult level of lung function. Respiratory Research, 2013, 14, 83.	1.4	16
39	Area-aggregated assessments of perceived environmental attributes may overcome single-source bias in studies of green environments and health: results from a cross-sectional survey in southern Sweden. Environmental Health, 2011, 10, 4.	1.7	49
40	Mapping Snakebite Epidemiology in Nicaragua â€“ Pitfalls and Possible Solutions. PLoS Neglected Tropical Diseases, 2010, 4, e896.	1.3	23