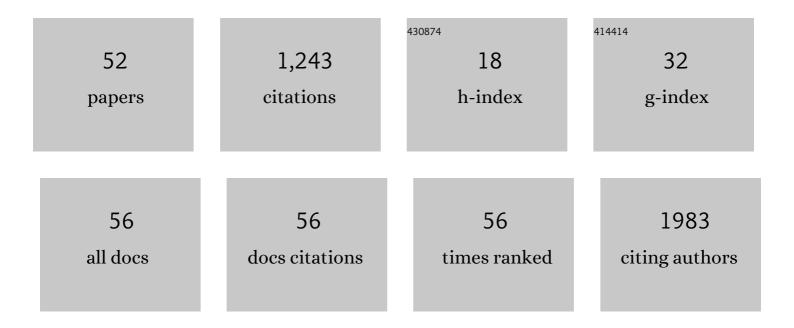
Antoine M Snijders

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5787019/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Influence of early life exposure, host genetics and diet on the mouse gut microbiome and metabolome. Nature Microbiology, 2017, 2, 16221.	13.3	138
2	Drosophila Histone Demethylase KDM5 Regulates Social Behavior through Immune Control and Gut Microbiota Maintenance. Cell Host and Microbe, 2019, 25, 537-552.e8.	11.0	125
3	Resveratrol Modulates the Gut Microbiota and Inflammation to Protect Against Diabetic Nephropathy in Mice. Frontiers in Pharmacology, 2020, 11, 1249.	3.5	106
4	<scp>FAM</scp> 83 family oncogenes are broadly involved in human cancers: an integrative multiâ€omics approach. Molecular Oncology, 2017, 11, 167-179.	4.6	102
5	Genetic and metabolic links between the murine microbiome and memory. Microbiome, 2020, 8, 53.	11.1	56
6	Identification of genetic factors that modify motor performance and body weight using Collaborative Cross mice. Scientific Reports, 2015, 5, 16247.	3.3	47
7	Systematic Analysis of Impact of Sampling Regions and Storage Methods on Fecal Gut Microbiome and Metabolome Profiles. MSphere, 2020, 5, .	2.9	37
8	FAM83D promotes epithelial-mesenchymal transition, invasion and cisplatin resistance through regulating the AKT/mTOR pathway in non-small-cell lung cancer. Cellular Oncology (Dordrecht), 2020, 43, 395-407.	4.4	31
9	Early exposure to thirdhand cigarette smoke affects body mass and the development of immunity in mice. Scientific Reports, 2017, 7, 41915.	3.3	30
10	Short-term early exposure to thirdhand cigarette smoke increases lung cancer incidence in mice. Clinical Science, 2018, 132, 475-488.	4.3	30
11	Adverse Health Effects of Thirdhand Smoke: From Cell to Animal Models. International Journal of Molecular Sciences, 2017, 18, 932.	4.1	29
12	A Multi-institutional Comparative Analysis of Proton and Photon Therapy-Induced Hematologic Toxicity in Patients With Medulloblastoma. International Journal of Radiation Oncology Biology Physics, 2021, 109, 726-735.	0.8	29
13	Genetic Differences in Transcript Responses to Low-Dose Ionizing Radiation Identify Tissue Functions Associated with Breast Cancer Susceptibility. PLoS ONE, 2012, 7, e45394.	2.5	28
14	Contribution of trace element exposure to gestational diabetes mellitus through disturbing the gut microbiome. Environment International, 2021, 153, 106520.	10.0	28
15	Light-Stress Influences the Composition of the Murine Gut Microbiome, Memory Function, and Plasma Metabolome. Frontiers in Molecular Biosciences, 2019, 6, 108.	3.5	26
16	<scp>FBXW</scp> 7 deletion contributes to lung tumor development and confers resistance to gefitinib therapy. Molecular Oncology, 2018, 12, 883-895.	4.6	25
17	Diverse tumour susceptibility in Collaborative Cross mice: identification of a new mouse model for human gastric tumourigenesis. Gut, 2019, 68, 1942-1952.	12.1	24
18	Prospective study reveals a microbiome signature that predicts the occurrence of post-operative enterocolitis in Hirschsprung disease (HSCR) patients. Gut Microbes, 2020, 11, 842-854.	9.8	24

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19	A new platform for ultra-high dose rate radiobiological research using the BELLA PW laser proton beamline. Scientific Reports, 2022, 12, 1484.	3.3	23
20	Revisiting the impact of age and molecular subtype on overall survival after radiotherapy in breast cancer patients. Scientific Reports, 2017, 7, 12587.	3.3	19
21	Genome-wide screen identifies a novel prognostic signature for breast cancer survival. Oncotarget, 2017, 8, 14003-14016.	1.8	19
22	An interferon signature identified by RNA-sequencing of mammary tissues varies across the estrous cycle and is predictive of metastasis-free survival. Oncotarget, 2014, 5, 4011-4025.	1.8	19
23	Overcoming the challenges of cancer drug resistance through bacterialâ€mediated therapy. Chronic Diseases and Translational Medicine, 2019, 5, 258-266.	1.2	18
24	Inactivation of Human Coronavirus by FATHHOME's Dry Sanitizer Device: Rapid and Eco-Friendly Ozone-Based Disinfection of SARS-CoV-2. Pathogens, 2021, 10, 339.	2.8	18
25	Clinical significance and molecular annotation of cellular morphometric subtypes in lower-grade gliomas discovered by machine learning. Neuro-Oncology, 2023, 25, 68-81.	1.2	18
26	Identification of genetic loci that control mammary tumor susceptibility through the host microenvironment. Scientific Reports, 2015, 5, 8919.	3.3	16
27	Acquired genomic aberrations associated with methotrexate resistance vary with background genomic instability. Genes Chromosomes and Cancer, 2008, 47, 71-83.	2.8	15
28	Integrative analysis of multi-omics data reveals distinct impacts of DDB1-CUL4 associated factors in human lung adenocarcinomas. Scientific Reports, 2017, 7, 333.	3.3	15
29	53BP1 Repair Kinetics for Prediction of In Vivo Radiation Susceptibility in 15 Mouse Strains. Radiation Research, 2020, 194, 485-499.	1.5	15
30	Human X chromosome exome sequencing identifies <i>BCORL1</i> as contributor to spermatogenesis. Journal of Medical Genetics, 2021, 58, 56-65.	3.2	13
31	Thirdhand smoke: Genotoxicity and carcinogenic potential. Chronic Diseases and Translational Medicine, 2020, 6, 27-34.	1.2	12
32	Simulated space radiation-induced mutants in the mouse kidney display widespread genomic change. PLoS ONE, 2017, 12, e0180412.	2.5	12
33	Thirdhand smoke exposure causes replication stress and impaired transcription in human lung cells. Environmental and Molecular Mutagenesis, 2020, 61, 635-646.	2.2	10
34	An integrated host-microbiome response to atrazine exposure mediates toxicity in Drosophila. Communications Biology, 2021, 4, 1324.	4.4	10
35	Genetic Susceptibility to Thirdhand-Smoke-Induced Lung Cancer Development. Nicotine and Tobacco Research, 2019, 21, 1294-1296.	2.6	8
36	<i>In utero</i> and early-life exposure to thirdhand smoke causes profound changes to the immune system. Clinical Science, 2021, 135, 1053-1063.	4.3	8

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37	A robust gene expression-based prognostic risk score predicts overall survival of lung adenocarcinoma patients. Oncotarget, 2018, 9, 6862-6871.	1.8	8
38	Micronucleus formation in human keratinocytes is dependent on radiation quality and tissue architecture. Environmental and Molecular Mutagenesis, 2015, 56, 22-31.	2.2	6
39	A Robust Gene Expression Prognostic Signature for Overall Survival in High-Grade Serous Ovarian Cancer. Journal of Oncology, 2019, 2019, 1-12.	1.3	6
40	Genetic background influences the effect of thirdhand smoke exposure on anxiety and memory in Collaborative Cross mice. Scientific Reports, 2021, 11, 13285.	3.3	6
41	Host genetics and gut microbiota cooperatively contribute to azoxymethane-induced acute toxicity in Collaborative Cross mice. Archives of Toxicology, 2021, 95, 949-958.	4.2	6
42	Multi-omics approach to infer cancer therapeutic targets on chromosome 20q across tumor types. Advances in Modern Oncology Research, 2016, 2, 215.	0.1	6
43	From Mouse to Human: Cellular Morphometric Subtype Learned From Mouse Mammary Tumors Provides Prognostic Value in Human Breast Cancer. Frontiers in Oncology, 2021, 11, 819565.	2.8	5
44	Genetic Background Modulates IncRNA-Coordinated Tissue Response to Low Dose Ionizing Radiation. International Journal of Genomics, 2015, 2015, 1-7.	1.6	4
45	Thirdhand cigarette smoke leads to ageâ€dependent and persistent alterations in the cecal microbiome of mice. MicrobiologyOpen, 2021, 10, e1198.	3.0	3
46	No difference in 4â€nitroquinoline induced tumorigenesis between germâ€free and colonized mice. Molecular Carcinogenesis, 2019, 58, 627-632.	2.7	2
47	Expression profiling reveals transcriptional regulation by Fbxw7/mTOR pathway in radiation-induced mouse thymic lymphomas. Oncotarget, 2015, 6, 44794-44805.	1.8	2
48	Dose Fractionation During Puberty Is More Detrimental to Mammary Gland Development Than an Equivalent Acute Dose of Radiation Exposure. International Journal of Radiation Oncology Biology Physics, 2021, 109, 1521-1532.	0.8	1
49	Identification of a novel 15â€gene expression signature predicting overall survival of human colorectal cancer. Clinical and Translational Medicine, 2020, 10, e258.	4.0	1
50	Low-dose radiation cancer susceptibility models. Aging, 2015, 7, 352-353.	3.1	0
51	Genetic background influences loss of heterozygosity patterns in radiation-induced mouse thymic lymphoma. Journal of Nature and Science, 2015, 1, e96.	1.1	0
52	Co-Expression Network Analysis of Fbxw7-Associated LncRNAs Reveals Their Functions in Radiation-Induced Thymic Lymphoma. , 2016, 1, 1-5.		0