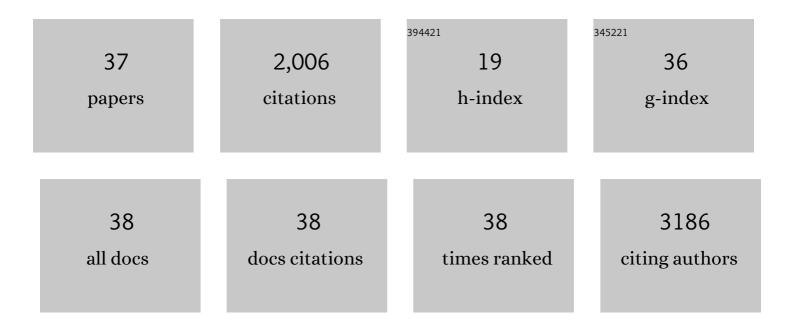
Vanina Guernier

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2854891/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	A genetic variant of <i>Burkholderia mallei</i> detected in Kuwait: Consequences for the PCR diagnosis of glanders. Transboundary and Emerging Diseases, 2021, 68, 960-963.	3.0	12
2	Describing fine spatiotemporal dynamics of rat fleas in an insular ecosystem enlightens abiotic drivers of murine typhus incidence in humans. PLoS Neglected Tropical Diseases, 2021, 15, e0009029.	3.0	4
3	Natural Horizontal Gene Transfer of Antimicrobial Resistance Genes in Campylobacter spp. From Turkeys and Swine. Frontiers in Microbiology, 2021, 12, 732969.	3.5	11
4	Predicting the Presence of Leptospires in Rodents from Environmental Indicators Opens Up Opportunities for Environmental Monitoring of Human Leptospirosis. Remote Sensing, 2021, 13, 325.	4.0	7
5	Leptospirosis infections among hospital patients, Sarawak, Malaysia. Tropical Diseases, Travel Medicine and Vaccines, 2021, 7, 32.	2.2	3
6	Seroprevalence of Coxiella burnetii (Q fever) Exposure in Humans on Reunion Island. Open Forum Infectious Diseases, 2019, 6, ofz227.	0.9	6
7	Seroprevalence of typhus group and spotted fever group Rickettsia exposures on Reunion island. BMC Research Notes, 2019, 12, 387.	1.4	4
8	Diversity of Mycobacterium tuberculosis in the Middle Fly District of Western Province, Papua New Guinea: microbead-based spoligotyping using DNA from Ziehl-Neelsen-stained microscopy preparations. Scientific Reports, 2019, 9, 15549.	3.3	5
9	Molecular Evidence of Drug-Resistant Tuberculosis in the Balimo Region of Papua New Guinea. Tropical Medicine and Infectious Disease, 2019, 4, 33.	2.3	3
10	Low seroprevalence of hepatitis E on Reunion island. One Health, 2019, 8, 100110.	3.4	0
11	Spatial distribution of tuberculosis in a rural region of Western Province, Papua New Guinea. Western Pacific Surveillance and Response Journal: WPSAR, 2019, 10, 31-38.	0.6	4
12	Biogeography of <i>Leptospira</i> in wild animal communities inhabiting the insular ecosystem of the western Indian Ocean islands and neighboring Africa. Emerging Microbes and Infections, 2018, 7, 1-12.	6.5	30
13	Individual and contextual risk factors for chikungunya virus infection: the SEROCHIK cross-sectional population-based study. Epidemiology and Infection, 2018, 146, 1056-1064.	2.1	12
14	Advances and challenges in barcoding pathogenic and environmental <i>Leptospira</i> . Parasitology, 2018, 145, 595-607.	1.5	38
15	Molecular diagnosis of suspected tuberculosis from archived smear slides from the Balimo region, Papua New Guinea. International Journal of Infectious Diseases, 2018, 67, 75-81.	3.3	8
16	A systematic review of human and animal leptospirosis in the Pacific Islands reveals pathogen and reservoir diversity. PLoS Neglected Tropical Diseases, 2018, 12, e0006503.	3.0	61
17	Gut microbiota disturbance during helminth infection: can it affect cognition and behaviour of children?. BMC Infectious Diseases, 2017, 17, 58.	2.9	56
18	The risk of global epidemic replacement with drug-resistant Mycobacterium tuberculosis strains. International Journal of Infectious Diseases, 2017, 56, 14-20.	3.3	67

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19	Human leptospirosis in Seychelles: A prospective study confirms the heavy burden of the disease but suggests that rats are not the main reservoir. PLoS Neglected Tropical Diseases, 2017, 11, e0005831.	3.0	30
20	Leptospira diversity in animals and humans in Tahiti, French Polynesia. PLoS Neglected Tropical Diseases, 2017, 11, e0005676.	3.0	27
21	Use of big data in the surveillance of veterinary diseases: early detection of tick paralysis in companion animals. Parasites and Vectors, 2016, 9, 303.	2.5	21
22	Investigation of a leptospirosis outbreak in triathlon participants, Réunion Island, 2013. Epidemiology and Infection, 2016, 144, 661-669.	2.1	20
23	Human Leptospirosis on Reunion Island, Indian Ocean: Are Rodents the (Only) Ones to Blame?. PLoS Neglected Tropical Diseases, 2016, 10, e0004733.	3.0	51
24	Rickettsia and Bartonella Species in Fleas from Reunion Island. American Journal of Tropical Medicine and Hygiene, 2015, 92, 617-619.	1.4	8
25	Utilisation de la trÃ [°] s haute résolution spatiale pour la caractérisation des habitats de rongeurs, vecteurs de zoonoses à la Réunion. Revue Francaise De Photogrammetrie Et De Teledetection, 2015, , 65-71.	0.2	4
26	Fleas of Small Mammals on Reunion Island: Diversity, Distribution and Epidemiological Consequences. PLoS Neglected Tropical Diseases, 2014, 8, e3129.	3.0	23
27	Pathogenic <i>Leptospira</i> spp. in Bats, Madagascar and Union of the Comoros. Emerging Infectious Diseases, 2012, 18, 1696-1698.	4.3	54
28	Deciphering arboviral emergence within insular ecosystems. Infection, Genetics and Evolution, 2012, 12, 1333-1339.	2.3	31
29	Pandemic Influenza Due to pH1N1/2009 Virus: Estimation of Infection Burden in Reunion Island through a Prospective Serosurvey, Austral Winter 2009. PLoS ONE, 2011, 6, e25738.	2.5	22
30	May Rapoport's Rule Apply to Human Associated Pathogens?. EcoHealth, 2009, 6, 509-521.	2.0	15
31	Use of cluster-graphs from spoligotyping data to study genotype similarities and a comparison of three indices to quantify recent tuberculosis transmission among culture positive cases in French Guiana during a eight year period. BMC Infectious Diseases, 2008, 8, 46.	2.9	19
32	Estimating Chikungunya prevalence in La Réunion Island outbreak by serosurveys: Two methods for two critical times of the epidemic. BMC Infectious Diseases, 2008, 8, 99.	2.9	199
33	GLOBALIZATION OF HUMAN INFECTIOUS DISEASE. Ecology, 2007, 88, 1903-1910.	3.2	120
34	An evaluation of the actual incidence of tuberculosis in French Guiana using a capture-recapture model. Microbes and Infection, 2006, 8, 721-727.	1.9	15
35	Long-Term Population-Based Genotyping Study of Mycobacterium tuberculosis Complex Isolates in the French Departments of the Americas. Journal of Clinical Microbiology, 2006, 44, 183-191.	3.9	41
36	Pathogen-Driven Selection and Worldwide HLA Class I Diversity. Current Biology, 2005, 15, 1022-1027.	3.9	449

#	Article	IF	CITATIONS
37	Ecology Drives the Worldwide Distribution of Human Diseases. PLoS Biology, 2004, 2, e141.	5.6	525