Bruno M Ghersi

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Genome Sequences of Five Arenaviruses from Pygmy Mice (Mus minutoides) in Sierra Leone. Microbiology Resource Announcements, 2022, 11, e0009522.	0.6	3
2	Amplification of pathogenic <i>Leptospira</i> infection with greater abundance and coâ€occurrence of rodent hosts across a counterâ€urbanizing landscape. Molecular Ecology, 2021, 30, 2145-2161.	3.9	6
3	Diversity, distribution and natural Leishmania infection of sand flies from communities along the Interoceanic Highway in the Southeastern Peruvian Amazon. PLoS Neglected Tropical Diseases, 2021, 15, e0009000.	3.0	2
4	Bridging the gap: Using reservoir ecology and human serosurveys to estimate Lassa virus spillover in West Africa. PLoS Computational Biology, 2021, 17, e1008811.	3.2	27
5	Rodent Virus Diversity and Differentiation across Post-Katrina New Orleans. Sustainability, 2021, 13, 8034.	3.2	1
6	Flooding and abandonment have shaped rat demography across post-Katrina New Orleans. Landscape and Urban Planning, 2021, 215, 104218.	7.5	2
7	Rodent assemblage structure reflects socioecological mosaics of counter-urbanization across post-Hurricane Katrina New Orleans. Landscape and Urban Planning, 2020, 195, 103710.	7.5	20
8	In the heart of the city: Trypanosoma cruzi infection prevalence in rodents across New Orleans. Parasites and Vectors, 2020, 13, 577.	2.5	10
9	The Expectations and Challenges of Wildlife Disease Research in the Era of Genomics: Forecasting with a Horizon Scan-like Exercise. Journal of Heredity, 2019, 110, 261-274.	2.4	9
10	Urban rat races: spatial population genomics of brown rats (<i>Rattus norvegicus</i>) compared across multiple cities. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20180245.	2.6	48
11	Rodent-Borne Bartonella Infection Varies According to Host Species Within and Among Cities. EcoHealth, 2017, 14, 771-782.	2.0	31
12	Abandonment, Ecological Assembly and Public Health Risks in Counter-Urbanizing Cities. Sustainability, 2016, 8, 491.	3.2	31
13	The Genetic Diversity of Influenza A Viruses in Wild Birds in Peru. PLoS ONE, 2016, 11, e0146059.	2.5	24
14	Disturbance, Reassembly, and Disease Risk in Socioecological Systems. EcoHealth, 2016, 13, 450-455.	2.0	23
15	Prevalencia de Paramixovirus en Murciélagos en Seis Zonas de Madre de Dios y Puno, Perú, con Dos Grados de Perturbación Antropogénica. Revista De Investigaciones Veterinarias Del Peru, 2016, 27, 241.	0.1	Ο
16	Wide distribution and ancient evolutionary history of simian foamy viruses in New World primates. Retrovirology, 2015, 12, 89.	2.0	26
17	Detection of Mycobacterium tuberculosis Complex in New World Monkeys in Peru. EcoHealth, 2015, 12, 288-297.	2.0	16
18	Native Rodent Species Are Unlikely Sources of Infection for Leishmania (Viannia) braziliensis along the Transoceanic Highway in Madre de Dios, Peru. PLoS ONE, 2014, 9, e103358.	2.5	5

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19	Andes Hantavirus Variant in Rodents, Southern Amazon Basin, Peru. Emerging Infectious Diseases, 2014, 20, 257-260.	4.3	9
20	PRESENCIA DEL VIRUS DE INFLUENZA AVIAR EN AVES SILVESTRES DE LOS HUMEDALES DE PUERTO VIEJO, LIMA. Revista De Investigaciones Veterinarias Del Peru, 2013, 24, .	0.1	3
21	AVIAN INFLUENZA INFECTIONS IN NONMIGRANT LAND BIRDS IN ANDEAN PERU. Journal of Wildlife Diseases, 2012, 48, 910-917.	0.8	10
22	Isolation of Low-pathogenic H7N3 Avian Influenza from Wild Birds in Peru. Journal of Wildlife Diseases, 2011, 47, 792-795.	0.8	10
23	Avian Influenza in Wild Birds, Central Coast of Peru. Emerging Infectious Diseases, 2009, 15, 935-938.	4.3	39