

# Ronnie G Gavilan

## List of Publications by Year in descending order

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Version: 2024-02-01

21  
papers

3,548  
citations

759055  
12  
h-index

477173  
29  
g-index

34  
all docs

34  
docs citations

34  
times ranked

5947  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cholera dynamics: lessons from an epidemic. <i>Journal of Medical Microbiology</i> , 2021, 70, .	0.7	14
2	Whole genome analysis of extensively drug resistant <i>Mycobacterium tuberculosis</i> strains in Peru. <i>Scientific Reports</i> , 2021, 11, 9493.	1.6	9
3	Phylogenomics reveals multiple introductions and early spread of SARS-CoV-2 into Peru. <i>Journal of Medical Virology</i> , 2021, 93, 5961-5968.	2.5	15
4	Emergence of ciprofloxacin-resistant <i>Neisseria meningitidis</i> B from asymptomatic carriers during an outbreak in Peru, 2017. <i>Journal of Medical Microbiology</i> , 2021, 70, .	0.7	0
5	Large Outbreak of Guillain-Barré Syndrome, Peru, 2019. <i>Emerging Infectious Diseases</i> , 2020, 26, 2778-2780.	2.0	5
6	Hidden biodiversity in Neotropical streams: DNA barcoding uncovers high endemicity of freshwater macroinvertebrates at small spatial scales. <i>PLoS ONE</i> , 2020, 15, e0231683.	1.1	11
7	Global Expansion of Pacific Northwest <i>Vibrio parahaemolyticus</i> Sequence Type 36. <i>Emerging Infectious Diseases</i> , 2020, 26, 323-326.	2.0	24
8	Phylogenetic structure of <i>Salmonella Enteritidis</i> provides context for a foodborne outbreak in Peru. <i>Scientific Reports</i> , 2020, 10, 22080.	1.6	5
9	Antimicrobial-producing <i>Pseudoalteromonas</i> from the marine environment of Panama shows a high phylogenetic diversity and clonal structure. <i>Journal of Basic Microbiology</i> , 2018, 58, 747-769.	1.8	24
10	Multiplex PCR assay for genotyping of <i>Mycobacterium tuberculosis</i> in Lima, Peru. <i>Revista Argentina De Microbiologia</i> , 2017, 49, 298-300.	0.4	0
11	Outbreak of <i>Vibrio parahaemolyticus</i> Sequence Type 120, Peru, 2009. <i>Emerging Infectious Diseases</i> , 2016, 22, 1235-1237.	2.0	26
12	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. <i>Nature Biotechnology</i> , 2016, 34, 828-837.	9.4	2,802
13	Transoceanic Spreading of Pathogenic Strains of <i>Vibrio parahaemolyticus</i> with Distinctive Genetic Signatures in the recA Gene. <i>PLoS ONE</i> , 2015, 10, e0117485.	1.1	32
14	Microbiota of Healthy Corals Are Active against Fungi in a Light-Dependent Manner. <i>ACS Chemical Biology</i> , 2014, 9, 2300-2308.	1.6	58
15	Imaging Mass Spectrometry of a Coral Microbe Interaction with Fungi. <i>Journal of Chemical Ecology</i> , 2013, 39, 1045-1054.	0.9	53
16	High clustering rates of multidrug-resistant <i>Mycobacterium tuberculosis</i> genotypes in Panama. <i>BMC Infectious Diseases</i> , 2013, 13, 442.	1.3	8
17	Molecular Epidemiology and Genetic Variation of Pathogenic <i>Vibrio parahaemolyticus</i> in Peru. <i>PLoS Neglected Tropical Diseases</i> , 2013, 7, e2210.	1.3	45
18	MS/MS networking guided analysis of molecule and gene cluster families. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E2611-20.	3.3	250

#	ARTICLE	IF	CITATIONS
19	Microevolution of Pandemic <i>Vibrio parahaemolyticus</i> Assessed by the Number of Repeat Units in Short Sequence Tandem Repeat Regions. PLoS ONE, 2012, 7, e30823.	1.1	11
20	Origins and colonization history of pandemic < i>Vibrio parahaemolyticus</i> in South America. Molecular Ecology, 2010, 19, 3924-3937.	2.0	20
21	Emergence of Asiatic Vibrio Diseases in South America in Phase With El Niño. Epidemiology, 2008, 19, 829-837.	1.2	91