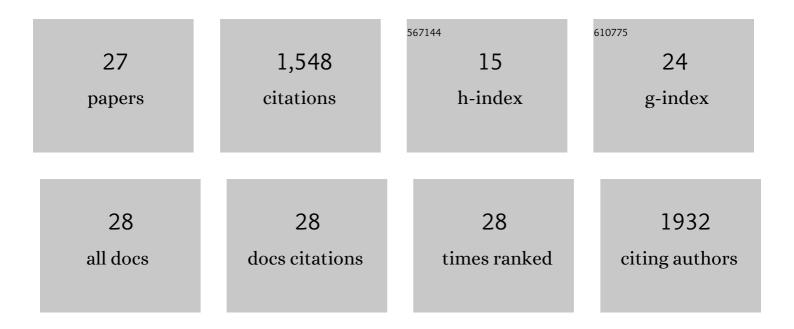
Carina Brehony

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3816413/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Establishment of sentinel surveillance of human clinical campylobacteriosis in Ireland. Zoonoses and Public Health, 2021, 68, 121-130.	0.9	4
2	Evaluation of molecular testing for Mycoplasma genitalium for symptomatic women. Irish Journal of Medical Science, 2021, , 1.	0.8	0
3	Molecular epidemiology of an extended multiple-species OXA-48 CPE outbreak in a hospital ward in Ireland, 2018–2019. Antimicrobial Stewardship & Healthcare Epidemiology, 2021, 1, .	0.2	3
4	Neuraminidase characterisation reveals very low levels of antiviral resistance and the presence of mutations associated with reduced antibody effectiveness in the Irish influenza 2018/2019 season. Journal of Clinical Virology, 2020, 132, 104653.	1.6	0
5	Detection of OXA-48-like-producing Enterobacterales in Irish recreational water. Science of the Total Environment, 2019, 690, 1-6.	3.9	25
6	An MLST approach to support tracking of plasmids carrying OXA-48-like carbapenemase. Journal of Antimicrobial Chemotherapy, 2019, 74, 1856-1862.	1.3	16
7	Hospital effluent: A reservoir for carbapenemase-producing Enterobacterales?. Science of the Total Environment, 2019, 672, 618-624.	3.9	83
8	Establishment of the European meningococcal strain collection genome library (EMSC-GL) for the 2011 to 2012 epidemiological year. Eurosurveillance, 2018, 23, .	3.9	8
9	Genomic surveillance and meningococcal group B vaccine coverage estimates after introduction of the vaccine into the national immunisation programme in the UK. Lancet, The, 2017, 389, S85.	6.3	3
10	Indistinguishable NDM-producing Escherichia coli isolated from recreational waters, sewage, and a clinical specimen in Ireland, 2016 to 2017. Eurosurveillance, 2017, 22, .	3.9	43
11	Resolution of a Protracted Serogroup B Meningococcal Outbreak with Whole-Genome Sequencing Shows Interspecies Genetic Transfer. Journal of Clinical Microbiology, 2016, 54, 2891-2899.	1.8	16
12	Distribution of Bexsero® Antigen Sequence Types (BASTs) in invasive meningococcal disease isolates: Implications for immunisation. Vaccine, 2016, 34, 4690-4697.	1.7	63
13	Authors' response: Meningococcal vaccine antigen diversity in global databases. Eurosurveillance, 2016, 21, .	3.9	0
14	An OMV Vaccine Derived from a Capsular Group B Meningococcus with Constitutive FetA Expression: Preclinical Evaluation of Immunogenicity and Toxicity. PLoS ONE, 2015, 10, e0134353.	1.1	9
15	Genomic epidemiology of age-associated meningococcal lineages in national surveillance: an observational cohort study. Lancet Infectious Diseases, The, 2015, 15, 1420-1428.	4.6	63
16	A novel meningococcal outer membrane vesicle vaccine with constitutive expression of FetA: A phase I clinical trial. Journal of Infection, 2015, 71, 326-337.	1.7	40
17	Meningococcal vaccine antigen diversity in global databases. Eurosurveillance, 2015, 20, .	3.9	15
18	Neisseria Adhesin A Variation and Revised Nomenclature Scheme. Vaccine Journal, 2014, 21, 966-971.	3.2	54

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#	Article	IF	CITATIONS
19	Implications of Differential Age Distribution of Disease-Associated Meningococcal Lineages for Vaccine Development. Vaccine Journal, 2014, 21, 847-853.	3.2	19
20	The Genus Neisseria. , 2014, , 881-900.		12
21	Ribosomal multilocus sequence typing: universal characterization of bacteria from domain to strain. Microbiology (United Kingdom), 2012, 158, 1005-1015.	0.7	497
22	The effect of iron availability on transcription of the Neisseria meningitidis fHbp gene varies among clonal complexes. Microbiology (United Kingdom), 2012, 158, 869-876.	0.7	20
23	Population structure of the <i>Yersinia pseudotuberculosis</i> complex according to multilocus sequence typing. Environmental Microbiology, 2011, 13, 3114-3127.	1.8	84
24	Variation of the factor H-binding protein of Neisseria meningitidis. Microbiology (United Kingdom), 2009, 155, 4155-4169.	0.7	79
25	Multilocus sequence typing for global surveillance of meningococcal disease. FEMS Microbiology Reviews, 2007, 31, 15-26.	3.9	105
26	Molecular typing of meningococci: recommendations for target choice and nomenclature. FEMS Microbiology Reviews, 2007, 31, 89-96.	3.9	150
27	A surveillance network for meningococcal disease in Europe. FEMS Microbiology Reviews, 2007, 31, 27-36.	3.9	134