

Carina Brehony

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3816413/publications.pdf>

Version: 2024-02-01

27
papers

1,548
citations

567144

15
h-index

610775

24
g-index

28
all docs

28
docs citations

28
times ranked

1932
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Establishment of sentinel surveillance of human clinical campylobacteriosis in Ireland. <i>Zoonoses and Public Health</i> , 2021, 68, 121-130. | 0.9 | 4 |
| 2 | Evaluation of molecular testing for <i>Mycoplasma genitalium</i> for symptomatic women. <i>Irish Journal of Medical Science</i> , 2021, , 1. | 0.8 | 0 |
| 3 | Molecular epidemiology of an extended multiple-species OXA-48 CPE outbreak in a hospital ward in Ireland, 2018â€“2019. <i>Antimicrobial Stewardship & Healthcare Epidemiology</i> , 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. <i>Journal of Clinical Virology</i> , 2020, 132, 104653. | 1.6 | 0 |
| 5 | Detection of OXA-48-like-producing Enterobacterales in Irish recreational water. <i>Science of the Total Environment</i> , 2019, 690, 1-6. | 3.9 | 25 |
| 6 | An MLST approach to support tracking of plasmids carrying OXA-48-like carbapenemase. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 1856-1862. | 1.3 | 16 |
| 7 | Hospital effluent: A reservoir for carbapenemase-producing Enterobacterales?. <i>Science of the Total Environment</i> , 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. <i>Eurosurveillance</i> , 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. <i>Lancet, The</i> , 2017, 389, S85. | 6.3 | 3 |
| 10 | Indistinguishable NDM-producing <i>Escherichia coli</i> isolated from recreational waters, sewage, and a clinical specimen in Ireland, 2016 to 2017. <i>Eurosurveillance</i> , 2017, 22, . | 3.9 | 43 |
| 11 | Resolution of a Protracted Serogroup B Meningococcal Outbreak with Whole-Genome Sequencing Shows Interspecies Genetic Transfer. <i>Journal of Clinical Microbiology</i> , 2016, 54, 2891-2899. | 1.8 | 16 |
| 12 | Distribution of Bexsero® Antigen Sequence Types (BASTs) in invasive meningococcal disease isolates: Implications for immunisation. <i>Vaccine</i> , 2016, 34, 4690-4697. | 1.7 | 63 |
| 13 | Authorsâ€™ response: Meningococcal vaccine antigen diversity in global databases. <i>Eurosurveillance</i> , 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. <i>PLoS ONE</i> , 2015, 10, e0134353. | 1.1 | 9 |
| 15 | Genomic epidemiology of age-associated meningococcal lineages in national surveillance: an observational cohort study. <i>Lancet Infectious Diseases, The</i> , 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. <i>Journal of Infection</i> , 2015, 71, 326-337. | 1.7 | 40 |
| 17 | Meningococcal vaccine antigen diversity in global databases. <i>Eurosurveillance</i> , 2015, 20, . | 3.9 | 15 |
| 18 | Neisseria Adhesin A Variation and Revised Nomenclature Scheme. <i>Vaccine Journal</i> , 2014, 21, 966-971. | 3.2 | 54 |

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