Noel D Mccarthy

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7713862/publications.pdf

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103 papers 7,014 citations

71061 41 h-index 81 g-index

109 all docs

109 docs citations

109 times ranked 7725 citing authors

#	Article	IF	CITATIONS
1	Cost-effectiveness of testing for latent tuberculosis infection in people with HIV. Aids, 2022, 36, 1-9.	1.0	4
2	Genomic Epidemiology Analysis of Infectious Disease Outbreaks Using TransPhylo. Current Protocols, 2021, 1, e60.	1.3	34
3	Association between vitamin D supplementation or serum vitamin D level and susceptibility to SARS-CoV-2 infection or COVID-19 including clinical course, morbidity and mortality outcomes? A systematic review. BMJ Open, 2021, 11, e043737.	0.8	18
4	Contrasting factors associated with COVID-19-related ICU admission and death outcomes in hospitalised patients by means of Shapley values. PLoS Computational Biology, 2021, 17, e1009121.	1.5	10
5	Evaluating multi-purpose syndromic surveillance systems – a complex problem. Online Journal of Public Health Informatics, 2021, 13, E15.	0.4	4
6	Two centuries of immunisation in the UK (part II). Archives of Disease in Childhood, 2020, 105, 216-222.	1.0	5
7	Incidence, risk factors, and health service burden of sequelae of campylobacter and non-typhoidal salmonella infections in England, 2000–2015: A retrospective cohort study using linked electronic health records. Journal of Infection, 2020, 81, 221-230.	1.7	14
8	Outbreaks of Shiga Toxin–Producing Escherichia coli Linked to Sprouted Seeds, Salad, and Leafy Greens: A Systematic Review. Journal of Food Protection, 2019, 82, 1950-1958.	0.8	46
9	Pneumococcal Disease: A Systematic Review of Health Utilities, Resource Use, Costs, and Economic Evaluations of Interventions. Value in Health, 2019, 22, 1329-1344.	0.1	30
10	The impact of childhood pneumococcal vaccination on hospital admissions in England: a whole population observational study. BMC Infectious Diseases, 2019, 19, 510.	1.3	11
11	Comparison of statistical algorithms for daily syndromic surveillance aberration detection. Bioinformatics, 2019, 35, 3110-3118.	1.8	20
12	A systematic review of source attribution of human campylobacteriosis using multilocus sequence typing. Eurosurveillance, 2019, 24, .	3.9	50
13	Comparative efficacy of drugs for treating giardiasis: a systematic update of the literature and network meta-analysis of randomized clinical trials. Journal of Antimicrobial Chemotherapy, 2018, 73, 596-606.	1.3	25
14	Bacterial Load and Molecular Markers Associated With Early-onset Group B Streptococcus. Pediatric Infectious Disease Journal, 2018, 37, e306-e314.	1.1	11
15	Novel application of the matched case–control design to compare food supply chains during an Escherichia coli O157 outbreak, United Kingdom, 2016. Eurosurveillance, 2018, 23, .	3.9	5
16	Campylobacter. , 2017, , 127-143.		0
17	Local genes for local bacteria: Evidence of allopatry in the genomes of transatlantic <i>Campylobacter</i> populations. Molecular Ecology, 2017, 26, 4497-4508.	2.0	36
18	Core Genome Multilocus Sequence Typing Scheme for Stable, Comparative Analyses of Campylobacter jejuni and C. coli Human Disease Isolates. Journal of Clinical Microbiology, 2017, 55, 2086-2097.	1.8	105

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19	Comparing interferon-gamma release assays with tuberculin skin test for identifying latent tuberculosis infection that progresses to active tuberculosis: systematic review and meta-analysis. BMC Infectious Diseases, 2017, 17, 200.	1.3	106
20	Indirect effects of childhood pneumococcal conjugate vaccination on invasive pneumococcal disease: a systematic review and meta-analysis. The Lancet Global Health, 2017, 5, e51-e59.	2.9	144
21	Factors Associated with Sequelae of Campylobacter and Non-typhoidal Salmonella Infections: A Systematic Review. EBioMedicine, 2017, 15, 100-111.	2.7	30
22	A seroprevalence study to determine the frequency of hantavirus infection in people exposed to wild and pet fancy rats in England. Epidemiology and Infection, 2017, 145, 2458-2465.	1.0	17
23	Genomeâ€wide association of functional traits linked with <scp><i>C</i></scp> <i>ampylobacter jejuni</i> survival from farm to fork. Environmental Microbiology, 2017, 19, 361-380.	1.8	88
24	Agreement between gastrointestinal panel testing and standard microbiology methods for detecting pathogens in suspected infectious gastroenteritis: Test evaluation and meta-analysis in the absence of a reference standard. PLoS ONE, 2017, 12, e0173196.	1.1	30
25	Recurrent seasonal outbreak of an emerging serotype of Shiga toxin-producing Escherichia coli (STEC) Tj ETQq1	1 0.7843	14 rgBT /Ove
26	Multiplex tests to identify gastrointestinal bacteria, viruses and parasites in people with suspected infectious gastroenteritis: a systematic review and economic analysis. Health Technology Assessment, 2017, 21, 1-188.	1.3	72
27	Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Outbreak of <i>Streptococcus pyogenes </i>). Emerging Infectious Diseases, 2016, 22, 973-980.	2.0	18
28	Community-onset sepsis and its public health burden: a systematic review. Systematic Reviews, 2016, 5, 81.	2.5	23
29	Pertussis vaccination for healthcare workers: staff attitudes and perceptions associated with high coverage vaccination programmes in England. Public Health, 2016, 137, 196-199.	1.4	9
30	A quantitative review of healthcare professionals' questions to a local immunization advice service: 4299 enquiries from 3 years. Journal of Public Health, 2016, 38, 578-584.	1.0	1
31	Vaccination of chemotherapy patients—effect of guideline implementation. Supportive Care in Cancer, 2016, 24, 2317-2321.	1.0	17
32	Genetic Diversity of Campylobacter jejuni and Campylobacter coli Isolates from Conventional Broiler Flocks and the Impacts of Sampling Strategy and Laboratory Method. Applied and Environmental Microbiology, 2016, 82, 2347-2355.	1.4	33
33	Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Outbreak of <i>Streptococcus pyogenes </i> Outbreak of <i>Streptococcus pyogenes</i>	2.0	2
34	Taste and Safety: Is the Exceptional Cuisine Offered by High End Restaurants Paralleled by High Standards of Food Safety?. PLOS Currents, 2016, 8, .	1.4	0
35	Wild birdâ€associated <scp><i>C</i></scp> <i>ampylobacter jejuni</i> isolates are a consistent source of human disease, in <scp>O</scp> xfordshire, <scp>U</scp> nited <scp>K</scp> ingdom. Environmental Microbiology Reports, 2015, 7, 782-788.	1.0	61
36	How to conduct systematic reviews more expeditiously?. Systematic Reviews, 2015, 4, 160.	2.5	53

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37	The longâ€term dynamics of <scp><i>C</i></scp> <i>ampylobacter</i> colonizing a freeâ€range broiler breeder flock: an observational study. Environmental Microbiology, 2015, 17, 938-946.	1.8	24
38	Variation in incidence and notification of Campylobacter and Salmonella by general practice in the Thames Valley area. Public Health, 2015, 129, 258-265.	1.4	2
39	Community-onset sepsis and its public health burden: protocol of a systematic review. Systematic Reviews, 2015, 4, 119.	2.5	10
40	Partial Failure of Milk Pasteurization as a Risk for the Transmission of <i>Campylobacter </i> From Cattle to Humans. Clinical Infectious Diseases, 2015, 61, 903-909.	2.9	41
41	Approaches to the detection of very small, common, and easily missed outbreaks that together contribute substantially to human <i>Cryptosporidium</i> infection. Epidemiology and Infection, 2014, 142, 1869-1876.	1.0	20
42	The Authors Reply. American Journal of Epidemiology, 2014, 179, 262-263.	1.6	3
43	McCarthy et al. Respond to "Evaluating Case-Chaos for Outbreaks Investigations". American Journal of Epidemiology, 2014, 180, 412-413.	1.6	0
44	Assessment of Mycobacterium tuberculosis transmission in Oxfordshire, UK, 2007–12, with whole pathogen genome sequences: an observational study. Lancet Respiratory Medicine,the, 2014, 2, 285-292.	5.2	199
45	Molecular Epidemiology of Campylobacter Species. , 2014, , 191-211.		9
46	Widespread acquisition of antimicrobial resistance among Campylobacter isolates from UK retail poultry and evidence for clonal expansion of resistant lineages. BMC Microbiology, 2013, 13, 160.	1.3	57
47	MLST revisited: the gene-by-gene approach to bacterial genomics. Nature Reviews Microbiology, 2013, 11, 728-736.	13.6	590
48	Self-reported adverse events in adolescents aged 13–18 years after mass vaccination with pertussis-containing vaccine, following a school outbreak. Public Health, 2013, 127, 1133-1136.	1.4	1
49	An epidemiological view of microbial genomic data. Lancet Infectious Diseases, The, 2013, 13, 104-105.	4.6	5
50	UK hantavirus, renal failure, and pet rats. Lancet, The, 2013, 381, 1070.	6.3	21
51	Factors affecting delay in initiation of treatment of tuberculosis in the Thames Valley, UK. Public Health, 2013, 127, 171-177.	1.4	25
52	Marked host specificity and lack of phylogeographic population structure of <i>Campylobacter jejuni</i> in wild birds. Molecular Ecology, 2013, 22, 1463-1472.	2.0	96
53	Real-Time Genomic Epidemiological Evaluation of Human Campylobacter Isolates by Use of Whole-Genome Multilocus Sequence Typing. Journal of Clinical Microbiology, 2013, 51, 2526-2534.	1.8	124
54	Three Authors Rreply. American Journal of Epidemiology, 2013, 177, 1022-1022.	1.6	3

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55	Duck Liver–associated Outbreak of Campylobacteriosis among Humans, United Kingdom, 2011. Emerging Infectious Diseases, 2013, 19, 1310-1313.	2.0	18
56	The burden and impact of measles among the Gypsy-Traveller communities, Thames Valley, 2006-09. Journal of Public Health, 2013, 35, 27-31.	1.0	24
57	Progressive genomeâ€wide introgression in agricultural <i>Campylobacter coli</i> . Molecular Ecology, 2013, 22, 1051-1064.	2.0	128
58	The "Case-Chaos Study―as an Adjunct or Alternative to Conventional Case-Control Study Methodology. American Journal of Epidemiology, 2012, 176, 497-505.	1.6	13
59	A Longitudinal 6-Year Study of the Molecular Epidemiology of Clinical Campylobacter Isolates in Oxfordshire, United Kingdom. Journal of Clinical Microbiology, 2012, 50, 3193-3201.	1.8	79
60	Transmission and control in an institutional pandemic influenza A(H1N1) 2009 outbreak. Epidemiology and Infection, 2012, 140, 1102-1110.	1.0	9
61	A large foodborne outbreak of norovirus in diners at a restaurant in England between January and February 2009. Epidemiology and Infection, 2012, 140, 1695-1701.	1.0	28
62	Molecular epidemiology of humanCampylobacter jejunishows association between seasonal and international patterns of disease. Epidemiology and Infection, 2012, 140, 2247-2255.	1.0	38
63	Extensively drug-resistant tuberculosis case in the Thames Valley, UK and public health interventions. Journal of Infection and Public Health, 2011, 4, 207-210.	1.9	2
64	<i>Campylobacter</i> populations in wild and domesticated Mallard ducks (<i>Anas) Tj ETQq0 0 0 rgBT /Overlo</i>	ck 10 Tf 5	0 382 Td (plat
65	The Prevalence of Campylobacter amongst a Free-Range Broiler Breeder Flock Was Primarily Affected by Flock Age. PLoS ONE, 2011, 6, e22825.	1.1	20
66	Campylobacter genotypes from poultry transportation crates indicate a source of contamination and transmission. Journal of Applied Microbiology, 2011, 110, 266-276.	1.4	30
67	Niche segregation and genetic structure of <i>Campylobacter jejuni</i> populations from wild and agricultural host species. Molecular Ecology, 2011, 20, 3484-3490.	2.0	105
68	Changes in Serogroup and Genotype Prevalence Among Carried Meningococci in the United Kingdom During Vaccine Implementation. Journal of Infectious Diseases, 2011, 204, 1046-1053.	1.9	44
69	An outbreak of norovirus infection linked to oyster consumption at a UK restaurant, February 2010. Journal of Public Health, 2011, 33, 205-211.	1.0	49
70	Introgression in the genus Campylobacter: generation and spread of mosaic alleles. Microbiology (United Kingdom), 2011, 157, 1066-1074.	0.7	47
71	Comparison of Campylobacter populations isolated from a free-range broiler flock before and after slaughter. International Journal of Food Microbiology, 2010, 137, 259-264.	2.1	32
72	MLST clustering of <i>Campylobacter jejuni</i> isolates from patients with gastroenteritis, reactive arthritis and Guillainâ€ÓBarré syndrome. Journal of Applied Microbiology, 2010, 108, 591-599.	1.4	45

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73	Evolution of an Agriculture-Associated Disease Causing Campylobacter coli Clade: Evidence from National Surveillance Data in Scotland. PLoS ONE, 2010, 5, e15708.	1.1	75
74	Host Association of <i>Campylobacter</i> Genotypes Transcends Geographic Variation. Applied and Environmental Microbiology, 2010, 76, 5269-5277.	1.4	116
75	Campylobacter genotypes from food animals, environmental sources and clinical disease in Scotland 2005/6. International Journal of Food Microbiology, 2009, 134, 96-103.	2.1	158
76	Dynamics of <i>Campylobacter</i> colonization of a natural host, <i>Sturnus vulgaris</i> (European) Tj ETQq0 (0 0 rgBT /0	Overlock 10 T
77	Invasive meningococcal disease: Completeness and timeliness of reporting of confirmed cases in Thames Valley, 2006–2007. Public Health, 2009, 123, 805-808.	1.4	7
78	<i>Campylobacter</i> Genotyping to Determine the Source of Human Infection. Clinical Infectious Diseases, 2009, 48, 1072-1078.	2.9	358
79	<i>Campylobacter</i> infection of broiler chickens in a freeâ€range environment. Environmental Microbiology, 2008, 10, 2042-2050.	1.8	89
80	Convergence of <i>Campylobacter</i> Species: Implications for Bacterial Evolution. Science, 2008, 320, 237-239.	6.0	231
81	Opa Protein Repertoires of Disease-Causing and Carried Meningococci. Journal of Clinical Microbiology, 2008, 46, 3033-3041.	1.8	15
82	Extended Sequence Typing of <i>Campylobacter </i> spp., United Kingdom. Emerging Infectious Diseases, 2008, 14, 1620-1622.	2.0	73
83	Early virological suppression with three-class antiretroviral therapy in HIV-infected African infants. Aids, 2008, 22, 1333-1343.	1.0	83
84	Resource Allocation during an Influenza Pandemic. Emerging Infectious Diseases, 2008, 14, 520-522.	2.0	9
85	Association of a Bacteriophage with Meningococcal Disease in Young Adults. PLoS ONE, 2008, 3, e3885.	1.1	62
86	Resource Allocation during an Influenza Pandemic. Emerging Infectious Diseases, 2008, 14, 1676b-1677.	2.0	1
87	Control of Human Immunodeficiency Virus Type 1 Is Associated with HLA-B*13 and Targeting of Multiple Gag-Specific CD8 + T-Cell Epitopes. Journal of Virology, 2007, 81, 3667-3672.	1.5	138
88	High frequency of rapid immunological progression in African infants infected in the era of perinatal HIV prophylaxis. Aids, 2007, 21, 1253-1261.	1.0	91
89	Outbreak of waterborne cryptosporidiosis associated with low oocyst concentrations. Epidemiology and Infection, 2007, 135, 1159-1164.	1.0	8
90	Host-associated Genetic Import in <i>Campylobacter jejuni </i> . Emerging Infectious Diseases, 2007, 13, 267-272.	2.0	134

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91	CD8+ T-cell responses to different HIV proteins have discordant associations with viral load. Nature Medicine, 2007, 13, 46-53.	15.2	910
92	Demographic risk factors for classical and atypical scrapie in Great Britain. Journal of General Virology, 2007, 88, 3486-3492.	1.3	35
93	Effectiveness Analyses May Underestimate Protection of Infants after Group C Meningococcal Immunization. Journal of Infectious Diseases, 2006, 194, 231-237.	1.9	17
94	Passive Sexual Transmission of Human Immunodeficiency Virus Type 1 Variants and Adaptation in New Hosts. Journal of Virology, 2006, 80, 7226-7234.	1.5	23
95	Genetic Analysis of Meningococci Carried by Children and Young Adults. Journal of Infectious Diseases, 2005, 191, 1263-1271.	1.9	178
96	Genetic Diversity and Carriage Dynamics of Neisseria lactamica in Infants. Infection and Immunity, 2005, 73, 2424-2432.	1.0	70
97	Multilocus Sequence Typing Directly on DNA from Clinical Samples and a Cultured Isolate To Investigate Linked Fatal Pneumococcal Disease in Residents of a Shelter for Homeless Men. Journal of Clinical Microbiology, 2005, 43, 2004-2008.	1.8	18
98	Distribution of Serogroups and Genotypes among Disease-Associated and Carried Isolates of Neisseria meningitidis from the Czech Republic, Greece, and Norway. Journal of Clinical Microbiology, 2004, 42, 5146-5153.	1.8	222
99	Olfactory detection of human bladder cancer by dogs: proof of principle study. BMJ: British Medical Journal, 2004, 329, 712.	2.4	318
100	Cross-sectional survey of users of Internet depression communities. BMC Psychiatry, 2003, 3, 19.	1.1	73
101	An Outbreak of Rift Valley Fever in Northeastern Kenya, 1997-98. Emerging Infectious Diseases, 2002, 8, 138-144.	2.0	263
102	Primary Peritonitis Due to Nonenteric Salmonellae. Clinical Infectious Diseases, 1999, 29, 211-212.	2.9	7
103	A world free from polio?. Lancet, The, 1997, 349, 956.	6.3	1