## Noel D Mccarthy

List of Publications by Year in descending order

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70961 60497 7,014 103 41 81 citations h-index g-index papers 109 109 109 7725 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	CD8+ T-cell responses to different HIV proteins have discordant associations with viral load. Nature Medicine, 2007, 13, 46-53.	15.2	910
2	MLST revisited: the gene-by-gene approach to bacterial genomics. Nature Reviews Microbiology, 2013, 11, 728-736.	13.6	590
3	<i>Campylobacter</i> Genotyping to Determine the Source of Human Infection. Clinical Infectious Diseases, 2009, 48, 1072-1078.	2.9	358
4	Olfactory detection of human bladder cancer by dogs: proof of principle study. BMJ: British Medical Journal, 2004, 329, 712.	2.4	318
5	An Outbreak of Rift Valley Fever in Northeastern Kenya, 1997-98. Emerging Infectious Diseases, 2002, 8, 138-144.	2.0	263
6	Convergence of <i>Campylobacter</i> Species: Implications for Bacterial Evolution. Science, 2008, 320, 237-239.	6.0	231
7	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
8	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
9	Genetic Analysis of Meningococci Carried by Children and Young Adults. Journal of Infectious Diseases, 2005, 191, 1263-1271.	1.9	178
10	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
11	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
12	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
13	Host-associated Genetic Import in <i>Campylobacter jejuni</i> . Emerging Infectious Diseases, 2007, 13, 267-272.	2.0	134
14	Progressive genomeâ€wide introgression in agricultural <i>Campylobacter coli</i> . Molecular Ecology, 2013, 22, 1051-1064.	2.0	128
15	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
16	Host Association of <i>Campylobacter</i> Genotypes Transcends Geographic Variation. Applied and Environmental Microbiology, 2010, 76, 5269-5277.	1.4	116
17	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
18	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

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19	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
20	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
21	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
22	<i>Campylobacter</i> infection of broiler chickens in a freeâ€range environment. Environmental Microbiology, 2008, 10, 2042-2050.	1.8	89
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	Early virological suppression with three-class antiretroviral therapy in HIV-infected African infants. Aids, 2008, 22, 1333-1343.	1.0	83
25	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
26	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
27	Cross-sectional survey of users of Internet depression communities. BMC Psychiatry, 2003, 3, 19.	1.1	73
28	Extended Sequence Typing of <i>Campylobacter </i> spp., United Kingdom. Emerging Infectious Diseases, 2008, 14, 1620-1622.	2.0	73
29	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
30	Genetic Diversity and Carriage Dynamics of Neisseria lactamica in Infants. Infection and Immunity, 2005, 73, 2424-2432.	1.0	70
31	<i>Campylobacter</i> populations in wild and domesticated Mallard ducks ( <i>Anas) Tj ETQq1 1 0.784314 rgBT</i>	/Overlock 1.0	10 Tf 50 262
32	Dynamics of <i>Campylobacter</i> colonization of a natural host, <i>Sturnus vulgaris</i> (European) Tj ETQq0 C	0 rgBT /O	verlock 10 Tr
33	Association of a Bacteriophage with Meningococcal Disease in Young Adults. PLoS ONE, 2008, 3, e3885.	1.1	62
34	Wild birdâ€essociated <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
35	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
36	How to conduct systematic reviews more expeditiously?. Systematic Reviews, 2015, 4, 160.	2.5	53

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37	A systematic review of source attribution of human campylobacteriosis using multilocus sequence typing. Eurosurveillance, 2019, 24, .	3.9	50
38	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
39	Introgression in the genus Campylobacter: generation and spread of mosaic alleles. Microbiology (United Kingdom), 2011, 157, 1066-1074.	0.7	47
40	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
41	MLST clustering of <i>Campylobacter jejuni</i> isolates from patients with gastroenteritis, reactive arthritis and Guillain–BarrÃf© syndrome. Journal of Applied Microbiology, 2010, 108, 591-599.	1.4	45
42	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
43	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
44	Molecular epidemiology of humanCampylobacter jejunishows association between seasonal and international patterns of disease. Epidemiology and Infection, 2012, 140, 2247-2255.	1.0	38
45	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
46	Demographic risk factors for classical and atypical scrapie in Great Britain. Journal of General Virology, 2007, 88, 3486-3492.	1.3	35
47	Genomic Epidemiology Analysis of Infectious Disease Outbreaks Using TransPhylo. Current Protocols, 2021, 1, e60.	1.3	34
48	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
49	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
50	Campylobacter genotypes from poultry transportation crates indicate a source of contamination and transmission. Journal of Applied Microbiology, 2011, 110, 266-276.	1.4	30
51	Factors Associated with Sequelae of Campylobacter and Non-typhoidal Salmonella Infections: A Systematic Review. EBioMedicine, 2017, 15, 100-111.	2.7	30
52	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
53	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
54	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

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55	Factors affecting delay in initiation of treatment of tuberculosis in the Thames Valley, UK. Public Health, 2013, 127, 171-177.	1.4	25
56	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
57	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
58	The longâ€ŧerm dynamics of <scp><i>C</i></scp> <i>ampylobacter</i> colonizing a freeâ€ŧange broiler breeder flock: an observational study. Environmental Microbiology, 2015, 17, 938-946.	1.8	24
59	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
60	Community-onset sepsis and its public health burden: a systematic review. Systematic Reviews, 2016, 5, 81.	2.5	23
61	UK hantavirus, renal failure, and pet rats. Lancet, The, 2013, 381, 1070.	6.3	21
62	Recurrent seasonal outbreak of an emerging serotype of Shiga toxin-producing Escherichia coli (STEC) Tj ETQq0	0	Overlock 10 T
63	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
64	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
65	Comparison of statistical algorithms for daily syndromic surveillance aberration detection. Bioinformatics, 2019, 35, 3110-3118.	1.8	20
66	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
67	Duck Liver–associated Outbreak of Campylobacteriosis among Humans, United Kingdom, 2011. Emerging Infectious Diseases, 2013, 19, 1310-1313.	2.0	18
68	Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Outbreak of <i>Streptococcus pyogenes </i>	2.0	18
69	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
70	Effectiveness Analyses May Underestimate Protection of Infants after Group C Meningococcal Immunization. Journal of Infectious Diseases, 2006, 194, 231-237.	1.9	17
71	Vaccination of chemotherapy patientsâ€"effect of guideline implementation. Supportive Care in Cancer, 2016, 24, 2317-2321.	1.0	17
72	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

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73	Opa Protein Repertoires of Disease-Causing and Carried Meningococci. Journal of Clinical Microbiology, 2008, 46, 3033-3041.	1.8	15
74	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
75	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
76	Bacterial Load and Molecular Markers Associated With Early-onset Group B Streptococcus. Pediatric Infectious Disease Journal, 2018, 37, e306-e314.	1.1	11
77	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
78	Community-onset sepsis and its public health burden: protocol of a systematic review. Systematic Reviews, 2015, 4, 119.	2.5	10
79	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
80	Resource Allocation during an Influenza Pandemic. Emerging Infectious Diseases, 2008, 14, 520-522.	2.0	9
81	Transmission and control in an institutional pandemic influenza A(H1N1) 2009 outbreak. Epidemiology and Infection, 2012, 140, 1102-1110.	1.0	9
82	Molecular Epidemiology of Campylobacter Species. , 2014, , 191-211.		9
83	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
84	Outbreak of waterborne cryptosporidiosis associated with low oocyst concentrations. Epidemiology and Infection, 2007, 135, 1159-1164.	1.0	8
85	Primary Peritonitis Due to Nonenteric Salmonellae. Clinical Infectious Diseases, 1999, 29, 211-212.	2.9	7
86	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
87	An epidemiological view of microbial genomic data. Lancet Infectious Diseases, The, 2013, 13, 104-105.	4.6	5
88	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
89	Two centuries of immunisation in the UK (part II). Archives of Disease in Childhood, 2020, 105, 216-222.	1.0	5
90	Cost-effectiveness of testing for latent tuberculosis infection in people with HIV. Aids, 2022, 36, 1-9.	1.0	4

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91	Evaluating multi-purpose syndromic surveillance systems – a complex problem. Online Journal of Public Health Informatics, 2021, 13, E15.	0.4	4
92	Three Authors Rreply. American Journal of Epidemiology, 2013, 177, 1022-1022.	1.6	3
93	The Authors Reply. American Journal of Epidemiology, 2014, 179, 262-263.	1.6	3
94	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
95	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
96	Integration of Genomic and Other Epidemiologic Data to Investigate and Control a Cross-Institutional Outbreak of <i>Streptococcus pyogenes </i> <institution contr<="" control="" of="" td=""><td>2.0</td><td>2</td></institution>	2.0	2
97	A world free from polio?. Lancet, The, 1997, 349, 956.	6.3	1
98	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
99	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
100	Resource Allocation during an Influenza Pandemic. Emerging Infectious Diseases, 2008, 14, 1676b-1677.	2.0	1
101	McCarthy et al. Respond to "Evaluating Case-Chaos for Outbreaks Investigations". American Journal of Epidemiology, 2014, 180, 412-413.	1.6	0
102	Campylobacter. , 2017, , 127-143.		0
103	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	О