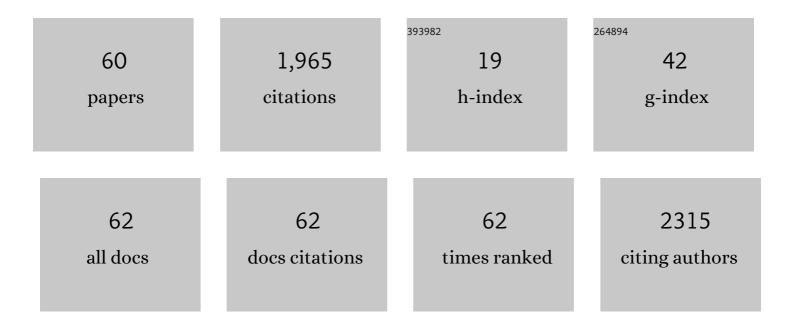
DEnis Piérard

List of Publications by Year in descending order

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ΠΕΝΙς ΡΙΑΩΟΔΑΡΟ

#	Article	IF	CITATIONS
1	Multicenter Evaluation of a Sequence-Based Protocol for Subtyping Shiga Toxins and Standardizing Stx Nomenclature. Journal of Clinical Microbiology, 2012, 50, 2951-2963.	1.8	710
2	Enterohemorrhagic Escherichia coli O26:H11/Hâ^': A New Virulent Clone Emerges in Europe. Clinical Infectious Diseases, 2013, 56, 1373-1381.	2.9	118
3	Isolation and virulence factors of vero cyto toxinproducing Escherichia coli in human stool samples. Clinical Microbiology and Infection, 1997, 3, 531-540.	2.8	95
4	The sink as a potential source of transmission of carbapenemase-producing Enterobacteriaceae in the intensive care unit. Antimicrobial Resistance and Infection Control, 2017, 6, 24.	1.5	79
5	Incidence and Virulence Determinants of Verocytotoxin-Producing Escherichia coli Infections in the Brussels-Capital Region, Belgium, in 2008–2010. Journal of Clinical Microbiology, 2012, 50, 1336-1345.	1.8	63
6	Use of Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry for Identification of Molds of the Fusarium Genus. Journal of Clinical Microbiology, 2015, 53, 465-476.	1.8	63
7	Hearing Loss With Congenital Cytomegalovirus Infection. Pediatrics, 2019, 144, .	1.0	63
8	Pertactin-deficient Bordetella pertussis isolates: evidence of increased circulation in Europe, 1998 to 2015. Eurosurveillance, 2019, 24, .	3.9	59
9	Anaerobic bacteraemia: a 10-year retrospective epidemiological survey. Anaerobe, 2016, 39, 54-59.	1.0	40
10	Ventilator-associated bacterial pneumonia in coronavirus 2019 disease, a retrospective monocentric cohort study. Journal of Infection and Chemotherapy, 2021, 27, 826-833.	0.8	39
11	Twenty-seven years of screening for Shiga toxin-producing Escherichia coli in a university hospital. Brussels, Belgium, 1987-2014. PLoS ONE, 2018, 13, e0199968.	1.1	34
12	Impact of DNA extraction on whole genome sequencing analysis for characterization and relatedness of Shiga toxin-producing Escherichia coli isolates. Scientific Reports, 2020, 10, 14649.	1.6	32
13	Acceptance criteria for identification results of Gram-negative rods by mass spectrometry. Journal of Medical Microbiology, 2011, 60, 684-686.	0.7	30
14	Population structure of Escherichia coli O26 : H11 with recent and repeated stx2 acquisition in multiple lineages. Microbial Genomics, 2017, 3, .	1.0	29
15	Differential dynamics and impacts of prophages and plasmids on the pangenome and virulence factor repertoires of Shiga toxin-producing Escherichia coli O145:H28. Microbial Genomics, 2020, 6, .	1.0	28
16	Surveillance of Circulating Bordetella pertussis Strains in Europe during 1998 to 2015. Journal of Clinical Microbiology, 2018, 56, .	1.8	26
17	Phylogenetic analysis of the Belgian HIV-1 epidemic reveals that local transmission is almost exclusively driven by men having sex with men despite presence of large African migrant communities. Infection, Genetics and Evolution, 2018, 61, 36-44.	1.0	26
18	The Benefits of Whole Genome Sequencing for Foodborne Outbreak Investigation from the Perspective of a National Reference Laboratory in a Smaller Country. Foods, 2020, 9, 1030.	1.9	23

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19	Undetectable viral RNA in follicular fluid, cumulus cells, and endometrial tissue samples in SARS-CoV-2–positive women. Fertility and Sterility, 2022, 117, 771-780.	0.5	23
20	Global spatial dynamics and vaccine-induced fitness changes of <i>Bordetella pertussis</i> . Science Translational Medicine, 2022, 14, eabn3253.	5.8	22
21	Respiratory Bacterial Culture Sampling in Expectorating and Non-expectorating Patients With Cystic Fibrosis. Frontiers in Pediatrics, 2018, 6, 403.	0.9	20
22	Validation strategy of a bioinformatics whole genome sequencing workflow for Shiga toxin-producing Escherichia coli using a reference collection extensively characterized with conventional methods. Microbial Genomics, 2021, 7, .	1.0	20
23	Banana infecting fungus,Fusarium musae, is also an opportunistic human pathogen: Are bananas potential carriers and source of fusariosis?. Mycologia, 2015, 107, 46-53.	0.8	19
24	Towards Real-Time and Affordable Strain-Level Metagenomics-Based Foodborne Outbreak Investigations Using Oxford Nanopore Sequencing Technologies. Frontiers in Microbiology, 2021, 12, 738284.	1.5	19
25	A Belgian Serosurveillance/Seroprevalence Study of Diphtheria, Tetanus and Pertussis Using a Luminex xMAP Technology-Based Pentaplex. Vaccines, 2016, 4, 16.	2.1	18
26	Diphtheria in Belgium: 2010–2017. Journal of Medical Microbiology, 2019, 68, 1517-1525.	0.7	17
27	Staphylococcus jettensis sp. nov., a coagulase-negative staphylococcal species isolated from human clinical specimens. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3250-3256.	0.8	15
28	Reclassification of Staphylococcus jettensis De Bel et al. 2013 as Staphylococcus petrasii subsp. jettensis subsp. nov. and emended description of Staphylococcus petrasii Pantucek et al. 2013. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 4198-4201.	0.8	15
29	Improving specificity of Bordetella pertussis detection using a four target real-time PCR. PLoS ONE, 2017, 12, e0175587.	1.1	15
30	Strain-Level Metagenomic Data Analysis of Enriched In Vitro and In Silico Spiked Food Samples: Paving the Way towards a Culture-Free Foodborne Outbreak Investigation Using STEC as a Case Study. International Journal of Molecular Sciences, 2020, 21, 5688.	1.8	14
31	A Practical Method to Implement Strain-Level Metagenomics-Based Foodborne Outbreak Investigation and Source Tracking in Routine. Microorganisms, 2020, 8, 1191.	1.6	14
32	Lessons learned from a textbook outbreak: EHEC-O157:H7 infections associated with the consumption of raw meat products, June 2012, Limburg, Belgium. Archives of Public Health, 2014, 72, 44.	1.0	13
33	Evaluation of the Alere SHIGA TOXIN QUIK CHEKâ,,¢ in comparison to multiplex Shiga toxin PCR. Diagnostic Microbiology and Infectious Disease, 2016, 86, 35-39.	0.8	11
34	<i>Fusarium musae</i> infected banana fruits as potential source of human fusariosis: May occur more frequently than we might think and hypotheses about infection. Communicative and Integrative Biology, 2016, 9, e1162934.	0.6	11
35	Detection of beta-lactamase-negative ampicillin resistance in Haemophilus influenzae in Belgium. Diagnostic Microbiology and Infectious Disease, 2019, 93, 243-249.	0.8	11
36	Genomics of an endemic cystic fibrosis Burkholderia multivorans strain reveals low within-patient evolution but high between-patient diversity. PLoS Pathogens, 2021, 17, e1009418.	2.1	11

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37	SARS-CoV-2 RNA and antibodies in tear fluid. BMJ Open Ophthalmology, 2021, 6, e000733.	0.8	10
38	Evaluation of four laboratory-based SARS-CoV-2 IgG antibody immunoassays. Diagnostic Microbiology and Infectious Disease, 2021, 100, 115313.	0.8	10
39	Performance of BacT/Alert resin-based FN plus bottles compared with BacT/Alert charcoal-based FN bottles for the detection of anaerobes in experimentally seeded blood cultures. Anaerobe, 2015, 35, 92-95.	1.0	9
40	Multicenter evaluation of the cobas® HIV-1 quantitative nucleic acid test for use on the cobas® 4800 system for the quantification of HIV-1 plasma viral load. Journal of Clinical Virology, 2019, 114, 43-49.	1.6	9
41	Comparison of Two Commercial Colorimetric Broth Microdilution Tests for Candida Susceptibility Testing: Sensititre YeastOne versus MICRONAUT-AM. Journal of Fungi (Basel, Switzerland), 2021, 7, 356.	1.5	9
42	The global population structure and evolutionary history of the acquisition of major virulence factor-encoding genetic elements in Shiga toxin-producing Escherichia coli O121:H19. Microbial Genomics, 2021, 7, .	1.0	9
43	Antibiotic Prescriptions Targeting Bacterial Respiratory Infections in Admitted Patients with COVID-19: A Prospective Observational Study. Infectious Diseases and Therapy, 2021, 10, 2575-2591.	1.8	7
44	Diagnostic accuracy of screening tests for patients suspected of COVID-19, a retrospective cohort study. Infectious Diseases, 2021, 53, 855-864.	1.4	6
45	Presumed Urinary Tract Infection in Patients Admitted with COVID-19: Are We Treating Too Much?. Antibiotics, 2021, 10, 1493.	1.5	6
46	Validation of the ColibrÃ-Instrument for Automated Preparation of MALDI-TOF MS Targets for Yeast Identification. Journal of Clinical Microbiology, 2022, 60, .	1.8	6
47	The burden of legionnaires' disease in Belgium, 2013 to 2017. Archives of Public Health, 2020, 78, 92.	1.0	5
48	Microbiology of the middle meatus in children requiring adenotonsillectomy. Journal of Laryngology and Otology, 1999, 113, 24-27.	0.4	4
49	Serodiagnosis of whooping cough in Belgium: results of the National Reference Centre for <i>Bordetella pertussis</i> anno 2013. Acta Clinica Belgica, 2016, 71, 86-91.	0.5	4
50	Matrix-Assisted Laser Desorption Ionization–Time of Flight Mass Spectrometry for Rapid Detection of Isolates Belonging to the Epidemic Clones Achromobacter xylosoxidans ST137 and Achromobacter ruhlandii DES from Cystic Fibrosis Patients. Journal of Clinical Microbiology, 2021, 59, e0094621.	1.8	4
51	Unique Phylogenetic Lineage Found in the <i>Fusarium</i> -like Clade after Re-examining BCCM/IHEM Fungal Culture Collection Material. Mycobiology, 2016, 44, 121-130.	0.6	3
52	Antimicrobial susceptibility testing of Eggerthella lenta blood culture isolates at a university hospital in Belgium from 2004 to 2018. Anaerobe, 2021, 69, 102348.	1.0	3
53	Fear for CoViD-19 and reluctance to work among health care workers during the epidemic, a prospective monocentric cohort study. American Journal of Infection Control, 2022, 50, 312-318.	1.1	3
54	Validation of Rapid Antimicrobial Susceptibility Testing directly from blood cultures using WASPLab®, including ColibrĂâ"¢ and Radian® in-Line Carousel. European Journal of Clinical Microbiology and Infectious Diseases, 2022, 41, 733-739.	1.3	3

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55	Enteric infections due to Escherichia coli. Clinical Microbiology and Infection, 1998, 4, 414-415.	2.8	2
56	Nontuberculous mycobacteria among pulmonary tuberculosis patients: a retrospective Belgian multicenter study. Acta Clinica Belgica, 2017, 72, 45-48.	0.5	2
57	SARS-CoV-2 seroprevalence among employees of a university hospital in Belgium during the 2020 COVID-19 outbreak (COVEMUZ-study). Epidemiology and Infection, 2021, 149, 1-24.	1.0	2
58	Symptomatic severe acute respiratory syndrome coronavirusÂ2 reinfection in a lupus patient treated with hydroxychloroquine: a case report. Journal of Medical Case Reports, 2021, 15, 572.	0.4	2
59	Proof-of-concept study of a new LC-ESI-MS/MS-based assay to identify Aspergillus spp. in artificially mixed samples using species/genus-specific proteotypic peptides. Mycological Progress, 2017, 16, 231-246.	0.5	1
60	Evaluation of the Copan Myco-TB kit for the decontamination of respiratory samples for the detection of Mycobacteria. European Journal of Clinical Microbiology and Infectious Diseases, 2018, 37, 711-714.	1.3	1