

# David C Coleman

## List of Publications by Year in descending order

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195  
papers

11,975  
citations

23567

58  
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33894

99  
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201  
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201  
docs citations

201  
times ranked

7598  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Field Guide to Pandemic, Epidemic and Sporadic Clones of Methicillin-Resistant <i>Staphylococcus aureus</i> . PLoS ONE, 2011, 6, e17936.	2.5	734
2	<i>Candida dubliniensis</i> sp. nov.: phenotypic and molecular characterization of a novel species associated with oral candidosis in HIV-infected individuals. Microbiology (United Kingdom), 1995, 141, 1507-1521.	1.8	701
3	Classification of Staphylococcal Cassette Chromosome <i>mec</i> (SCC <i>mec</i> ): Guidelines for Reporting Novel SCC <i>mec</i> Elements. Antimicrobial Agents and Chemotherapy, 2009, 53, 4961-4967.	3.2	669
4	<i>Candida dubliniensis</i> : Characteristics and Identification. Journal of Clinical Microbiology, 1998, 36, 329-334.	3.9	341
5	Detection of Staphylococcal Cassette Chromosome <i>mec</i> Type XI Carrying Highly Divergent <i>mecA</i> , <i>mecI</i> , <i>mecR1</i> , <i>blaZ</i> , and <i>ccr</i> Genes in Human Clinical Isolates of Clonal Complex 130 Methicillin-Resistant <i>Staphylococcus aureus</i> . Antimicrobial Agents and Chemotherapy, 2011, 55, 3765-3773.	3.2	336
6	Antifungal drug susceptibilities of oral <i>Candida dubliniensis</i> isolates from human immunodeficiency virus (HIV)-infected and non-HIV-infected subjects and generation of stable fluconazole-resistant derivatives in vitro. Antimicrobial Agents and Chemotherapy, 1997, 41, 617-623.	3.2	263
7	Simple, Inexpensive, Reliable Method for Differentiation of <i>Candida dubliniensis</i> from <i>Candida albicans</i> . Journal of Clinical Microbiology, 1998, 36, 2093-2095.	3.9	209
8	Candidiasis. Aids, 1997, 11, 557-567.	2.2	205
9	Comparative genomics of the fungal pathogens <i>Candida dubliniensis</i> and <i>Candida albicans</i> . Genome Research, 2009, 19, 2231-2244.	5.5	195
10	Identification and Expression of Multidrug Transporters Responsible for Fluconazole Resistance in <i>Candida dubliniensis</i> . Antimicrobial Agents and Chemotherapy, 1998, 42, 1819-1830.	3.2	194
11	Comparison of the epidemiology, drug resistance mechanisms, and virulence of and. FEMS Yeast Research, 2004, 4, 369-376.	2.3	190
12	Widespread geographic distribution of oral <i>Candida dubliniensis</i> strains in human immunodeficiency virus-infected individuals. Journal of Clinical Microbiology, 1997, 35, 960-964.	3.9	178
13	<i>Candida dubliniensis</i> : phylogeny and putative virulence factors. Microbiology (United Kingdom), 1998, 144, 829-838.	1.8	171
14	Seven Novel Variants of the Staphylococcal Chromosomal Cassette <i>mec</i> in Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates from Ireland. Antimicrobial Agents and Chemotherapy, 2005, 49, 2070-2083.	3.2	157
15	The Emergence and Importation of Diverse Genotypes of Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) Harboring the Panton-Valentine Leukocidin Gene ( <i>pvl</i> ) Reveal that <i>pvl</i> Is a Poor Marker for Community-Acquired MRSA Strains in Ireland. Journal of Clinical Microbiology, 2007, 45, 2554-2563.	3.9	154
16	Guidelines for Reporting Novel <i>mecA</i> Gene Homologues. Antimicrobial Agents and Chemotherapy, 2012, 56, 4997-4999.	3.2	144
17	Phylogenetic analysis and rapid identification of <i>Candida dubliniensis</i> based on analysis of ACT1 intron and exon sequences. Microbiology (United Kingdom), 1999, 145, 1871-1882.	1.8	143
18	Oligonucleotide fingerprinting of isolates of <i>Candida</i> species other than <i>C. albicans</i> and of atypical <i>Candida</i> species from human immunodeficiency virus-positive and AIDS patients. Journal of Clinical Microbiology, 1993, 31, 2124-2133.	3.9	139

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19	Oral Candida in HIV Infection and AIDS: New Perspectives/New Approaches. Critical Reviews in Microbiology, 1993, 19, 61-82.	6.1	138
20	Staphylococcal cassette chromosome mec: Recent advances and new insights. International Journal of Medical Microbiology, 2013, 303, 350-359.	3.6	135
21	Molecular epidemiology, phylogeny and evolution of Candida albicans. Infection, Genetics and Evolution, 2014, 21, 166-178.	2.3	120
22	Rapid Identification of <i>Candida dubliniensis</i> with Commercial Yeast Identification Systems. Journal of Clinical Microbiology, 1999, 37, 3533-3539.	3.9	111
23	Emerging pathogens. Medical Mycology, 2000, 38, 225-236.	0.7	107
24	Recovery of <i>Candida dubliniensis</i> from Non-Human Immunodeficiency Virus-Infected Patients in Israel. Journal of Clinical Microbiology, 2000, 38, 170-174.	3.9	104
25	<i>Candida albicans</i> versus <i>Candida dubliniensis</i> : Why Is <i>C. albicans</i> More Pathogenic?. International Journal of Microbiology, 2012, 2012, 1-7.	2.3	102
26	Biofilm problems in dental unit water systems and its practical control. Journal of Applied Microbiology, 2009, 106, 1424-1437.	3.1	99
27	Comparative Genomics and the Evolution of Pathogenicity in Human Pathogenic Fungi. Eukaryotic Cell, 2011, 10, 34-42.	3.4	99
28	Candida dubliniensis: Ten years on. FEMS Microbiology Letters, 2005, 253, 9-17.	1.8	97
29	Comparative genomics using Candida albicans DNA microarrays reveals absence and divergence of virulence-associated genes in Candida dubliniensis. Microbiology (United Kingdom), 2004, 150, 3363-3382.	1.8	96
30	Diversity of Staphylococcus aureus Isolates in European Wildlife. PLoS ONE, 2016, 11, e0168433.	2.5	94
31	Identification and Characterization of the Multidrug Resistance Gene <i>cfr</i> in a Pantone-Valentine Leukocidin-Positive Sequence Type 8 Methicillin-Resistant <i>Staphylococcus aureus</i> IVa (USA300) Isolate. Antimicrobial Agents and Chemotherapy, 2010, 54, 4978-4984.	3.2	91
32	Management of dental unit waterline biofilms in the 21st century. Future Microbiology, 2011, 6, 1209-1226.	2.0	90
33	Identification of <i>Candida dubliniensis</i> Based on Temperature and Utilization of Xylose and $\pm$ -Methyl- $\alpha$ -D-Glucoside as Determined with the API 20C AUX and Vitek YBC Systems. Journal of Clinical Microbiology, 1999, 37, 3804-3808.	3.9	87
34	Molecular genetic approaches to identification, epidemiology and taxonomy of non-albicans Candida species. Journal of Medical Microbiology, 1996, 44, 399-408.	1.8	86
35	MDR1 -Mediated Drug Resistance in Candida dubliniensis. Antimicrobial Agents and Chemotherapy, 2001, 45, 3416-3421.	3.2	86
36	High prevalence of non-albicans yeasts and detection of anti-fungal resistance in the oral flora of patients with advanced cancer. Palliative Medicine, 2003, 17, 477-481.	3.1	86

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37	Molecular and phenotypic analysis of <i>Candida dubliniensis</i> : A recently identified species linked with oral candidosis in HIV-infected and AIDS patients. <i>Oral Diseases</i> , 1997, 3, S96-101.	3.0	84
38	Characterization of a Novel Arginine Catabolic Mobile Element (ACME) and Staphylococcal Chromosomal Cassette <i>mec</i> Composite Island with Significant Homology to <i>Staphylococcus epidermidis</i> ACME Type II in Methicillin-Resistant <i>Staphylococcus aureus</i> Genotype ST22-MRSA-IV. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 1896-1905.	3.2	83
39	Novel multiresistance <i>cfr</i> plasmids in linezolid-resistant methicillin-resistant <i>Staphylococcus epidermidis</i> and vancomycin-resistant <i>Enterococcus faecium</i> (VRE) from a hospital outbreak: co-location of <i>cfr</i> and <i>optrA</i> in VRE. <i>Journal of Antimicrobial Chemotherapy</i> , 2017, 72, 3252-3257.	3.0	80
40	Comparative Transcript Profiling of <i>Candida albicans</i> and <i>Candida dubliniensis</i> Identifies <i>SFL2</i> , a <i>C. albicans</i> Gene Required for Virulence in a Reconstituted Epithelial Infection Model. <i>Eukaryotic Cell</i> , 2010, 9, 251-265.	3.4	78
41	Analysis of the reduction in expression of tetracycline resistance determined by transposon Tn10 in the multicopy state. <i>Molecular Genetics and Genomics</i> , 1981, 182, 171-177.	2.4	77
42	Identification of Four Distinct Genotypes of <i>Candida dubliniensis</i> and Detection of Microevolution In Vitro and In Vivo. <i>Journal of Clinical Microbiology</i> , 2002, 40, 556-574.	3.9	77
43	Cloning and expression in <i>Escherichia coli</i> and <i>Staphylococcus aureus</i> of the beta-lysin determinant from <i>Staphylococcus aureus</i> : evidence that bacteriophage conversion of beta-lysin activity is caused by insertional inactivation of the beta-lysin determinant. <i>Microbial Pathogenesis</i> , 1986, 1, 549-564.	2.9	76
44	Rapid PCR Test for Discriminating between <i>Candida albicans</i> and <i>Candida dubliniensis</i> Isolates Using Primers Derived from the pH-Regulated PHR1 and PHR2 Genes of <i>C. albicans</i> . <i>Journal of Clinical Microbiology</i> , 1999, 37, 1587-1590.	3.9	75
45	Insertional inactivation of the <i>Staphylococcus aureus</i> $\beta$ -toxin by bacteriophage $\phi$ 13 occurs by site- and orientation-specific integration of the $\phi$ 13 genome. <i>Molecular Microbiology</i> , 1991, 5, 933-939.	2.5	74
46	Effective control of dental chair unit waterline biofilm and marked reduction of bacterial contamination of output water using two peroxide-based disinfectants. <i>Journal of Hospital Infection</i> , 2002, 52, 192-205.	2.9	74
47	Detection of <i>mecC</i> -Positive <i>Staphylococcus aureus</i> (CC130-MRSA-XI) in Diseased European Hedgehogs ( <i>Erinaceus europaeus</i> ) in Sweden. <i>PLoS ONE</i> , 2013, 8, e66166.	2.5	74
48	Lower filamentation rates of <i>Candida dubliniensis</i> contribute to its lower virulence in comparison with <i>Candida albicans</i> . <i>Fungal Genetics and Biology</i> , 2007, 44, 920-931.	2.1	73
49	Differentiation of <i>Candida dubliniensis</i> from <i>Candida albicans</i> on Pal's Agar. <i>Journal of Clinical Microbiology</i> , 2003, 41, 4787-4789.	3.9	72
50	Emergence of Sequence Type 779 Methicillin-Resistant <i>Staphylococcus aureus</i> Harboring a Novel Pseudo Staphylococcal Cassette Chromosome <i>mec</i> ( <i>mec</i> )-SCC-SCC <sub>CRISPR</sub> Composite Element in Irish Hospitals. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 524-531.	3.2	72
51	Differentiation of <i>Candida dubliniensis</i> from <i>Candida albicans</i> on Staib Agar and Caffeic Acid-Ferric Citrate Agar. <i>Journal of Clinical Microbiology</i> , 2001, 39, 323-327.	3.9	71
52	Panton-Valentine Leukocidin-Positive <i>Staphylococcus aureus</i> in Ireland from 2002 to 2011: 21 Clones, Frequent Importation of Clones, Temporal Shifts of Predominant Methicillin-Resistant <i>S. aureus</i> Clones, and Increasing Multiresistance. <i>Journal of Clinical Microbiology</i> , 2014, 52, 859-870.	3.9	68
53	Emergence of Hospital- and Community-Associated Panton-Valentine Leukocidin-Positive Methicillin-Resistant <i>Staphylococcus aureus</i> Genotype ST772-MRSA-V in Ireland and Detailed Investigation of an ST772-MRSA-V Cluster in a Neonatal Intensive Care Unit. <i>Journal of Clinical Microbiology</i> , 2012, 50, 841-847.	3.9	67
54	Isolation of <i>C. dubliniensis</i> from insulin-using diabetes mellitus patients. <i>Journal of Oral Pathology and Medicine</i> , 2000, 29, 86-90.	2.7	66

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55	Comparative Genotypes, Staphylococcal Cassette Chromosome <i>mec</i> (SCC <i>mec</i> ) Genes and Antimicrobial Resistance amongst <i>Staphylococcus epidermidis</i> and <i>Staphylococcus haemolyticus</i> Isolates from Infections in Humans and Companion Animals. <i>PLoS ONE</i> , 2015, 10, e0138079.	2.5	66
56	Detection of Staphylococcal Cassette Chromosome <i>mec</i> -Associated DNA Segments in Multiresistant Methicillin-Susceptible <i>Staphylococcus aureus</i> (MSSA) and Identification of <i>Staphylococcus epidermidis</i> <i>ccrAB4</i> in both Methicillin-Resistant <i>S. aureus</i> and MSSA. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 4407-4419.	3.2	65
57	Novel organization of the site-specific integration and excision recombination functions of the <i>Staphylococcus aureus</i> serotype F virulence-converting phages $\phi$ 13 and $\phi$ 42. <i>Molecular Microbiology</i> , 1995, 16, 877-893.	2.5	64
58	Rapid Identification of <i>Candida dubliniensis</i> by Indirect Immunofluorescence Based on Differential Localization of Antigens on <i>C. dubliniensis</i> Blastospores and <i>Candida albicans</i> Germ Tubes. <i>Journal of Clinical Microbiology</i> , 1998, 36, 2428-2433.	3.9	62
59	Molecular Mechanisms of Itraconazole Resistance in <i>Candida dubliniensis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2003, 47, 2424-2437.	3.2	61
60	Enterotoxin production by <i>Staphylococcus aureus</i> isolates from cases of septicaemia and from healthy carriers. <i>Journal of Medical Microbiology</i> , 1989, 28, 163-172.	1.8	60
61	A polymeric system for the intra-oral delivery of an anti-fungal agent. <i>Biomaterials</i> , 2001, 22, 2319-2324.	11.4	60
62	Distinctive Carbohydrate Assimilation Profiles Used To Identify the First Clinical Isolates of <i>Candida dubliniensis</i> Recovered in the United States. <i>Journal of Clinical Microbiology</i> , 1998, 36, 1467-1467.	3.9	60
63	Persistence, replacement, and microevolution of <i>Cryptococcus neoformans</i> strains in recurrent meningitis in AIDS patients. <i>Journal of Clinical Microbiology</i> , 1996, 34, 1739-1744.	3.9	59
64	Multilocus Sequence Typing Reveals that the Population Structure of <i>Candida dubliniensis</i> Is Significantly Less Divergent than That of <i>Candida albicans</i> . <i>Journal of Clinical Microbiology</i> , 2008, 46, 652-664.	3.9	57
65	<i>Candida dubliniensis</i> : an emerging opportunistic pathogen. <i>Current Topics in Medical Mycology</i> , 1997, 8, 15-25.	0.8	56
66	Genomic DNA fingerprinting of clinical isolates of <i>Helicobacter pylori</i> using short oligonucleotide probes containing repetitive sequences. <i>Journal of Applied Bacteriology</i> , 1996, 81, 509-517.	1.1	55
67	Enhanced Discrimination of Highly Clonal ST22-Methicillin-Resistant <i>Staphylococcus aureus</i> IV Isolates Achieved by Combining <i>spa</i> , <i>dru</i> , and Pulsed-Field Gel Electrophoresis Typing Data. <i>Journal of Clinical Microbiology</i> , 2010, 48, 1839-1852.	3.9	55
68	Differential Filamentation of <i>Candida albicans</i> and <i>Candida dubliniensis</i> Is Governed by Nutrient Regulation of <i>UME6</i> Expression. <i>Eukaryotic Cell</i> , 2010, 9, 1383-1397.	3.4	55
69	Differential regulation of the transcriptional repressor <i>NRG1</i> accounts for altered host-cell interactions in <i>Candida albicans</i> and <i>Candida dubliniensis</i> . <i>Molecular Microbiology</i> , 2007, 66, 915-929.	2.5	50
70	Evolution and Global Transmission of a Multidrug-Resistant, Community-Associated Methicillin-Resistant <i>Staphylococcus aureus</i> Lineage from the Indian Subcontinent. <i>MBio</i> , 2019, 10, .	4.1	50
71	Linezolid resistance in <i>Enterococcus faecium</i> and <i>Enterococcus faecalis</i> from hospitalized patients in Ireland: high prevalence of the MDR genes <i>optrA</i> and <i>poxxA</i> in isolates with diverse genetic backgrounds. <i>Journal of Antimicrobial Chemotherapy</i> , 2020, 75, 1704-1711.	3.0	48
72	Sau421, a Bcgl-like restriction modification system encoded by the <i>Staphylococcus aureus</i> quadruple-converting phage $\phi$ 42. <i>Microbiology (United Kingdom)</i> , 2005, 151, 1301-1311.	1.8	47

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73	The recent emergence in hospitals of multidrug-resistant community-associated sequence type 1 and spa type t127 methicillin-resistant <i>Staphylococcus aureus</i> investigated by whole-genome sequencing: Implications for screening. <i>PLoS ONE</i> , 2017, 12, e0175542.	2.5	45
74	Molecular Typing of ST239-MRSA-III From Diverse Geographic Locations and the Evolution of the SCCmec III Element During Its Intercontinental Spread. <i>Frontiers in Microbiology</i> , 2018, 9, 1436.	3.5	45
75	Cloning and characterisation of the serC and aroA genes of <i>Yersinia enterocolitica</i> , and construction of an aroA mutant. <i>Gene</i> , 1989, 84, 23-30.	2.2	44
76	Isogenic Strain Construction and Gene Targeting in <i>Candida dubliniensis</i> . <i>Journal of Bacteriology</i> , 2001, 183, 2859-2865.	2.2	44
77	Casein Agar: a Useful Medium for Differentiating <i>Candida dubliniensis</i> from <i>Candida albicans</i> . <i>Journal of Clinical Microbiology</i> , 2003, 41, 1259-1262.	3.9	44
78	SUSCEPTIBILITY TO ANTIMICROBIAL AGENTS AND ANALYSIS OF PLASMIDS IN GENTAMICIN- AND METHICILLIN-RESISTANT STAPHYLOCOCCUS AUREUS FROM DUBLIN HOSPITALS. <i>Journal of Medical Microbiology</i> , 1985, 20, 157-167.	1.8	43
79	Molecular typing of nasal carriage isolates of <i>Staphylococcus aureus</i> from an Irish university student population based on toxin gene PCR, agr locus types and multiple locus, variable number tandem repeat analysis. <i>Journal of Medical Microbiology</i> , 2008, 57, 348-358.	1.8	43
80	Disinfection procedures: Their efficacy and effect on dimensional accuracy and surface quality of an irreversible hydrocolloid impression material. <i>Journal of Dentistry</i> , 2011, 39, 133-140.	4.1	43
81	Staphylococcal toxins in human disease. <i>Journal of Applied Bacteriology</i> , 1990, 69, 101S-107S.	1.1	42
82	Range Expansion and the Origin of USA300 North American Epidemic Methicillin-Resistant <i>Staphylococcus aureus</i> . <i>MBio</i> , 2018, 9, .	4.1	42
83	Significant Enrichment and Diversity of the Staphylococcal Arginine Catabolic Mobile Element ACME in <i>Staphylococcus epidermidis</i> Isolates From Subgingival Peri-implantitis Sites and Periodontal Pockets. <i>Frontiers in Microbiology</i> , 2018, 9, 1558.	3.5	42
84	The <i>Candida dubliniensis</i> CdCDR1 Gene Is Not Essential for Fluconazole Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2002, 46, 2829-2841.	3.2	41
85	Bacterial contamination of dental chair units in a modern dental hospital caused by leakage from suction system hoses containing extensive biofilm. <i>Journal of Hospital Infection</i> , 2005, 59, 348-360.	2.9	41
86	The role of manufacturers in reducing biofilms in dental chair waterlines. <i>Journal of Dentistry</i> , 2007, 35, 701-711.	4.1	41
87	When are the hands of healthcare workers positive for methicillin-resistant <i>Staphylococcus aureus</i> ? <i>Journal of Hospital Infection</i> , 2010, 75, 107-111.	2.9	41
88	Contribution of whole-genome sequencing to understanding of the epidemiology and control of methicillin-resistant <i>Staphylococcus aureus</i> . <i>Journal of Hospital Infection</i> , 2019, 102, 189-199.	2.9	40
89	Analysis of tetracycline resistance encoded by transposon Tn10: deletion mapping of tetracycline-sensitive point mutations and identification of two structural genes. <i>Journal of Bacteriology</i> , 1983, 153, 921-929.	2.2	40
90	Functional analysis of the phospholipase C gene CaPLC1 and two unusual phospholipase C genes, CaPLC2 and CaPLC3, of <i>Candida albicans</i> . <i>Microbiology (United Kingdom)</i> , 2005, 151, 3381-3394.	1.8	39

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91	Air and surface contamination patterns of methicillin-resistant <i>Staphylococcus aureus</i> on eight acute hospital wards. <i>Journal of Hospital Infection</i> , 2014, 86, 201-208.	2.9	39
92	Lack of a relationship between Lewis antigen expression and <i>cagA</i> , <i>CagA</i> , <i>vacA</i> and <i>VacA</i> status of Irish <i>Helicobacter pylori</i> isolates. <i>FEMS Immunology and Medical Microbiology</i> , 1999, 24, 79-90.	2.7	38
93	Molecular cloning and characterization of the genetic determinant encoding CS3 fimbriae of enterotoxigenic <i>Escherichia coli</i> . <i>Microbial Pathogenesis</i> , 1987, 2, 195-209.	2.9	37
94	Molecular analysis of <i>Helicobacter pylori</i> populations in antral biopsies from individual patients using randomly amplified polymorphic DNA (RAPD) fingerprinting. <i>FEMS Immunology and Medical Microbiology</i> , 1995, 10, 317-324.	2.7	37
95	Reduced Azole Susceptibility in Genotype 3 <i>Candida dubliniensis</i> Isolates Associated with Increased Cd CDR1 and Cd CDR2 Expression. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 1312-1318.	3.2	37
96	A novel automated waterline cleaning system that facilitates effective and consistent control of microbial biofilm contamination of dental chair unit waterlines: A one-year study. <i>Journal of Dentistry</i> , 2006, 34, 648-661.	4.1	37
97	Genome-wide gene expression profiling and a forward genetic screen show that differential expression of the sodium ion transporter <i>Ena21</i> contributes to the differential tolerance of <i>Candida albicans</i> and <i>Candida dubliniensis</i> to osmotic stress. <i>Molecular Microbiology</i> , 2009, 72, 216-228.	2.5	37
98	Enrichment of Multilocus Sequence Typing Clade 1 with Oral <i>Candida albicans</i> Isolates in Patients with Untreated Periodontitis. <i>Journal of Clinical Microbiology</i> , 2012, 50, 3335-3344.	3.9	37
99	Extensive Genetic Diversity Identified among Sporadic Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates Recovered in Irish Hospitals between 2000 and 2012. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 1907-1917.	3.2	37
100	First Report of <i>ccr</i> -Carrying Plasmids in the Pandemic Sequence Type 22 Methicillin-Resistant <i>Staphylococcus aureus</i> Staphylococcal Cassette Chromosome <i>mec</i> Type IV Clone. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 3007-3015.	3.2	37
101	Genetic characterization of a phospholipase C gene from <i>Candida albicans</i> : presence of homologous sequences in <i>Candida</i> species other than <i>Candida albicans</i> . <i>Microbiology (United Kingdom)</i> , 1998, 144, 55-72.	1.8	37
102	Characterisation of MRSA from Malta and the description of a Maltese epidemic MRSA strain. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2010, 29, 163-170.	2.9	36
103	Genotyping <i>Candida albicans</i> from <i>Candida</i> Leukoplakia and Non- <i>Candida</i> Leukoplakia Shows No Enrichment of Multilocus Sequence Typing Clades but Enrichment of ABC Genotype C in <i>Candida</i> Leukoplakia. <i>PLoS ONE</i> , 2013, 8, e73738.	2.5	36
104	Multicenter prospective surveillance of oral <i>Candida dubliniensis</i> among adult Brazilian human immunodeficiency virus-positive and AIDS patients. <i>Diagnostic Microbiology and Infectious Disease</i> , 2001, 41, 29-35.	1.8	34
105	Epidemiological typing of MRSA isolates from blood cultures taken in Irish hospitals participating in the European Antimicrobial Resistance Surveillance System (1999-2003). <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2006, 25, 79-89.	2.9	34
106	The Effect of Rapid Screening for Methicillin-Resistant <i>Staphylococcus aureus</i> (MRSA) on the Identification and Earlier Isolation of MRSA-Positive Patients. <i>Infection Control and Hospital Epidemiology</i> , 2010, 31, 374-381.	1.8	34
107	Molecular diagnosis and epidemiology of fungal infections. <i>Medical Mycology</i> , 1998, 36 Suppl 1, 249-57.	0.7	34
108	In situ hybridization and the polymerase chain reaction (PCR) in the analysis of biopsies and exfoliative cytology specimens for definitive diagnosis of oral hairy leukoplakia (DHL). <i>Journal of Oral Pathology and Medicine</i> , 1994, 23, 302-308.	2.7	33

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109	Purification and germination of <i>Candida albicans</i> and <i>Candida dubliniensis</i> chlamydo spores cultured in liquid media. <i>FEMS Yeast Research</i> , 2009, 9, 1051-1060.	2.3	33
110	CYP56 (Dit2p) in <i>Candida albicans</i> : Characterization and Investigation of Its Role in Growth and Antifungal Drug Susceptibility. <i>Antimicrobial Agents and Chemotherapy</i> , 2008, 52, 3718-3724.	3.2	32
111	Triclosan Antagonizes Fluconazole Activity against <i>Candida albicans</i> . <i>Journal of Dental Research</i> , 2012, 91, 65-70.	5.2	32
112	Novel 5-Flucytosine-Resistant Clade of <i>Candida dubliniensis</i> from Saudi Arabia and Egypt Identified by Cd25 Fingerprinting. <i>Journal of Clinical Microbiology</i> , 2005, 43, 4026-4036.	3.9	31
113	Optimisation of the long-term efficacy of dental chair waterline disinfection by the identification and rectification of factors associated with waterline disinfection failure. <i>Journal of Dentistry</i> , 2007, 35, 438-451.	4.1	29
114	DNA Microarray Profiling of a Diverse Collection of Nosocomial Methicillin-Resistant <i>Staphylococcus aureus</i> Isolates Assigns the Majority to the Correct Sequence Type and Staphylococcal Cassette Chromosome <i>mec</i> (SCC <i>mec</i> ) Type and Results in the Subsequent Identification and Characterization of Novel SCC <i>mec</i> -SCC <sub>M1</sub> Composite Islands. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 5340-5355.	3.2	29
115	Hospital outbreak of linezolid-resistant and vancomycin-resistant ST80 <i>Enterococcus faecium</i> harbouring an <i>optrA</i> -encoding conjugative plasmid investigated by whole-genome sequencing. <i>Journal of Hospital Infection</i> , 2020, 105, 726-735.	2.9	28
116	A centralised, automated dental hospital water quality and biofilm management system using neutral Ecasol, maintains dental unit waterline output at better than potable quality: A 2-year longitudinal study. <i>Journal of Dentistry</i> , 2009, 37, 748-762.	4.1	26
117	Intra-Hospital, Inter-Hospital and Intercontinental Spread of ST78 MRSA From Two Neonatal Intensive Care Unit Outbreaks Established Using Whole-Genome Sequencing. <i>Frontiers in Microbiology</i> , 2018, 9, 1485.	3.5	26
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