Arnfinn Sundsfjord

List of Publications by Year in descending order

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124 papers 8,025 citations

47 h-index 84 g-index

140 all docs

140 docs citations

140 times ranked

9722 citing authors

#	Article	IF	CITATIONS
1	Structural basis for PoxtA-mediated resistance to phenicol and oxazolidinone antibiotics. Nature Communications, 2022, 13, 1860.	5.8	25
2	A nationwide genomic study of clinical <i>Klebsiella pneumoniae</i> in Norway 2001–15: introduction and spread of ESBLs facilitated by clonal groups CG15 and CG307. Journal of Antimicrobial Chemotherapy, 2022, 77, 665-674.	1.3	16
3	Burden, Antibiotic Resistance, and Clonality of Shigella spp. Implicated in Community-Acquired Acute Diarrhoea in Lilongwe, Malawi. Tropical Medicine and Infectious Disease, 2021, 6, 63.	0.9	12
4	Risk of cervical intraepithelial neoplasia grade 3 or higher (CIN3+) among women with HPV-test in 1990–1992, a 30-year follow-up study. Infectious Agents and Cancer, 2021, 16, 46.	1.2	5
5	Carbapenem Resistance Determinants Acquired through Novel Chromosomal Integrations in Extensively Drug-Resistant Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2021, 65, e0028921.	1.4	6
6	Novel genomic islands and a new vanD-subtype in the first sporadic VanD-type vancomycin resistant enterococci in Norway. PLoS ONE, 2021, 16, e0255187.	1.1	6
7	High prevalence of multidrug resistant ESBL- and plasmid mediated AmpC-producing clinical isolates of Escherichia coli at Maputo Central Hospital, Mozambique. BMC Infectious Diseases, 2021, 21, 16.	1.3	24
8	Gastrointestinal carriage of Klebsiella pneumoniae in a general adult population: a cross-sectional study of risk factors and bacterial genomic diversity. Gut Microbes, 2021, 13, 1939599.	4.3	34
9	Alternative <i>vanHAX</i> promoters and increased <i>vanA</i> plasmid copy number resurrect silenced glycopeptide resistance in <i>Enterococcus faecium</i> Journal of Antimicrobial Chemotherapy, 2021, 76, 876-882.	1.3	11
10	Efficacy of mecillinam against clinical multidrug-resistant Escherichia coli in a murine urinary tract infection model. International Journal of Antimicrobial Agents, 2020, 55, 105851.	1.1	10
11	Antibiotic Sensitivity Screening of Klebsiella spp. and Raoultella spp. Isolated from Marine Bivalve Molluscs Reveal Presence of CTX-M-Producing K. pneumoniae. Microorganisms, 2020, 8, 1909.	1.6	17
12	\hat{l}^2 -lactam and fluoroquinolone resistance in Enterobacteriaceae from imported and locally-produced chicken in Mozambique. Journal of Infection in Developing Countries, 2020, 14, 471-478.	0.5	9
13	Whole-Genome Sequences of Two Multidrug-Resistant Acinetobacter baumannii Strains Isolated from Patients with Urinary Tract Infection in Ghana. Microbiology Resource Announcements, 2019, 8, .	0.3	1
14	Genomic characterization of multidrug-resistant ESBL-producing Klebsiella pneumoniae isolated from a Ghanaian teaching hospital. International Journal of Infectious Diseases, 2019, 85, 117-123.	1.5	28
15	Spread of Plasmid-Encoded NDM-1 and GES-5 Carbapenemases among Extensively Drug-Resistant and Pandrug-Resistant Clinical Enterobacteriaceae in Durban, South Africa. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	65
16	End non-essential use of antimicrobials in livestock. BMJ: British Medical Journal, 2018, 360, k259.	2.4	7
17	Pharmacokinetics and Pharmacodynamics of Fosfomycin and Its Activity against Extended-Spectrum-Î ² -Lactamase-, Plasmid-Mediated AmpC-, and Carbapenemase-Producing Escherichia coli in a Murine Urinary Tract Infection Model. Antimicrobial Agents and Chemotherapy, 2018, 62, .	1.4	31
18	Performance of the EUCAST disc diffusion method and two MIC methods in detection of Enterobacteriaceae with reduced susceptibility to meropenem: the NordicAST CPE study. Journal of Antimicrobial Chemotherapy, 2018, 73, 2738-2747.	1.3	13

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19	First environmental sample containing plasmidâ€mediated colistinâ€resistant ESBLâ€producing <i>Escherichia coli</i> detected in Norway. Apmis, 2017, 125, 822-825.	0.9	28
20	Fecal carriage of extended spectrum β-lactamase producing Escherichia coli and Klebsiella pneumoniae after urinary tract infection – A three year prospective cohort study. PLoS ONE, 2017, 12, e0173510.	1.1	38
21	Molecular and epidemiological characterization of carbapenemase-producing Enterobacteriaceae in Norway, 2007 to 2014. PLoS ONE, 2017, 12, e0187832.	1.1	53
22	A comparison of extended spectrum \hat{l}^2 -lactamase producing Escherichia coli from clinical, recreational water and wastewater samples associated in time and location. PLoS ONE, 2017, 12, e0186576.	1.1	70
23	Identification of a novel IMI carbapenemase variant (IMI-9) in Enterobacter cloacae complex. International Journal of Antimicrobial Agents, 2016, 48, 764-765.	1.1	6
24	Multilevel population genetic analysis of <i>vanA</i> and <i>vanB Enterococcus faecium</i> causing nosocomial outbreaks in 27 countries (1986–2012). Journal of Antimicrobial Chemotherapy, 2016, 71, 3351-3366.	1.3	129
25	Understanding the mechanisms and drivers of antimicrobial resistance. Lancet, The, 2016, 387, 176-187.	6.3	1,633
26	The antimicrobial activity of mecillinam, nitrofurantoin, temocillin and fosfomycin and comparative analysis of resistance patterns in a nationwide collection of ESBL-producing ⟨i⟩Escherichia coli⟨/i⟩ in Norway 2010–2011. Infectious Diseases, 2016, 48, 99-107.	1.4	39
27	En kvinne med sepsis etter brannskade i Pakistan. Tidsskrift for Den Norske Laegeforening, 2016, 136, 1228-1232.	0.2	2
28	Identification of VIM-2-Producing Pseudomonas aeruginosa from Tanzania Is Associated with Sequence Types 244 and 640 and the Location of <i>bla</i> _{VIM-2} in a TniC Integron. Antimicrobial Agents and Chemotherapy, 2015, 59, 682-685.	1.4	26
29	Investigating the mobilome in clinically important lineages of Enterococcus faecium and Enterococcus faecalis. BMC Genomics, 2015, 16, 282.	1.2	82
30	Extended-spectrum β-lactamase-producing Enterobacteriaceae among pregnant women in Norway: prevalence and maternal–neonatal transmission. Journal of Perinatology, 2015, 35, 907-912.	0.9	54
31	Persistence of a pKPN3-Like CTX-M-15-Encoding IncFIIK Plasmid in a Klebsiella pneumonia ST17 Host during Two Years of Intestinal Colonization. PLoS ONE, 2015, 10, e0116516.	1.1	31
32	High Rate of Per Oral Mecillinam Treatment Failure in Community-Acquired Urinary Tract Infections Caused by ESBL-Producing Escherichia coli. PLoS ONE, 2014, 9, e85889.	1.1	55
33	Emergence of mutationâ€based linezolidâ€resistant invasive <i>Enterococcus faecalis</i> in a haemodialysis patient in Norway. Apmis, 2014, 122, 83-84.	0.9	4
34	Within-Population Distribution of Trimethoprim Resistance in Escherichia coli before and after a Community-Wide Intervention on Trimethoprim Use. Antimicrobial Agents and Chemotherapy, 2014, 58, 7492-7500.	1.4	8
35	Increased prevalence of aminoglycoside resistance in clinical isolates of Escherichia coli and Klebsiella spp. in Norway is associated with the acquisition of AAC(3)-II and AAC(6′)-Ib. Diagnostic Microbiology and Infectious Disease, 2014, 78, 66-69.	0.8	46
36	High prevalence of faecal carriage of ESBL-producing Enterobacteriaceae in Norwegian patients with gastroenteritis. Scandinavian Journal of Infectious Diseases, 2014, 46, 462-465.	1.5	15

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37	Performance of the EUCAST Disk Diffusion Method, the CLSI Agar Screen Method, and the Vitek 2 Automated Antimicrobial Susceptibility Testing System for Detection of Clinical Isolates of Enterococci with Low- and Medium-Level VanB-Type Vancomycin Resistance: a Multicenter Study. Journal of Clinical Microbiology, 2014, 52, 1582-1589.	1.8	33
38	Multilocus sequence typing and ftsI sequencing: a powerful tool for surveillance of penicillin-binding protein 3-mediated beta-lactam resistance in nontypeable Haemophilus influenzae. BMC Microbiology, 2014, 14, 131.	1.3	49
39	Evaluation of the total MBL confirm kit (ROSCO) for detection of metallo-β-lactamases in Pseudomonas aeruginosa and Acinetobacter baumannii. Diagnostic Microbiology and Infectious Disease, 2014, 79, 486-488.	0.8	8
40	Extended-spectrum beta-lactamase-producing bacteria are not detected in supragingival plaque samples from human fecal carriers of ESBL-producing <i>Enterobacteriaceae</i> Journal of Oral Microbiology, 2014, 6, 24026.	1.2	5
41	A Multicentre Hospital Outbreak in Sweden Caused by Introduction of a vanB2 Transposon into a Stably Maintained pRUM-Plasmid in an Enterococcus faecium ST192 Clone. PLoS ONE, 2014, 9, e103274.	1.1	33
42	Identification of Enterobacteriaceae isolates with OXA-48 and coproduction of OXA-181 and NDM-1 in Norway. Journal of Antimicrobial Chemotherapy, 2013, 68, 1682-1685.	1.3	28
43	Risk factors for acquisition of CTX-M-15 extended-spectrum beta-lactamase-producing Klebsiella pneumoniae during an outbreak in a neonatal intensive care unit in Norway. Scandinavian Journal of Infectious Diseases, 2013, 45, 54-58.	1.5	28
44	ICESluvan, a 94-Kilobase Mosaic Integrative Conjugative Element Conferring Interspecies Transfer of VanB-Type Glycopeptide Resistance, a Novel Bacitracin Resistance Locus, and a Toxin-Antitoxin Stabilization System. Journal of Bacteriology, 2013, 195, 5381-5390.	1.0	17
45	Dissemination of a Carbapenem-Resistant Acinetobacter baumannii Strain Belonging to International Clone II/Sequence Type 2 and Harboring a Novel AbaR4-Like Resistance Island in Latvia. Antimicrobial Agents and Chemotherapy, 2013, 57, 1069-1072.	1.4	26
46	Long-term faecal carriage in infants and intra-household transmission of CTX-M-15-producing Klebsiella pneumoniae following a nosocomial outbreak. Journal of Antimicrobial Chemotherapy, 2013, 68, 1043-1048.	1.3	75
47	Large IncHI2-plasmids encode extended-spectrum \hat{l}^2 -lactamases (ESBLs) in Enterobacter spp. bloodstream isolates, and support ESBL-transfer to Escherichia coli. Clinical Microbiology and Infection, 2013, 19, E516-E518.	2.8	19
48	Prevalence and population structure of Staphylococcus aureus nasal carriage in healthcare workers in a general population. The TromsÃ, Staph and Skin Study. Epidemiology and Infection, 2013, 141, 143-152.	1.0	43
49	A Long-Term Low-Frequency Hospital Outbreak of KPC-Producing Klebsiella pneumoniae Involving Intergenus Plasmid Diffusion and a Persisting Environmental Reservoir. PLoS ONE, 2013, 8, e59015.	1.1	102
50	Characterisation of the Plasmidome within Enterococcus faecalis Isolated from Marginal Periodontitis Patients in Norway. PLoS ONE, 2013, 8, e62248.	1.1	28
51	Risk Factors for Community-Acquired Urinary Tract Infections Caused by ESBL-Producing Enterobacteriaceae –A Case–Control Study in a Low Prevalence Country. PLoS ONE, 2013, 8, e69581.	1.1	170
52	A nationwide study of mechanisms conferring reduced susceptibility to extended-spectrum cephalosporins in clinical Escherichia coli and Klebsiella spp. isolates. Scandinavian Journal of Infectious Diseases, 2012, 44, 927-933.	1.5	6
53	Sequence types and plasmid carriage of uropathogenic Escherichia coli devoid of phenotypically detectable resistance. Journal of Antimicrobial Chemotherapy, 2012, 67, 69-73.	1.3	45
54	Insight into antimicrobial susceptibility and population structure of contemporary human Enterococcus faecalis isolates from Europe. Journal of Antimicrobial Chemotherapy, 2012, 67, 551-558.	1.3	102

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55	Fecal colonization of VIM-1–producing Klebsiella pneumoniae and in vivo transfer of multidrug-resistant IncN plasmid in a renal transplant patient. Diagnostic Microbiology and Infectious Disease, 2012, 72, 363-366.	0.8	6
56	Insights into the global molecular epidemiology of carbapenem non-susceptible clones of Acinetobacter baumannii. Drug Resistance Updates, 2012, 15, 237-247.	6.5	261
57	Increased high-level gentamicin resistance in invasive <i>Enterococcus faecium</i> is associated with <i>ac($6\hat{a}\in^2$)le-aph($2\hat{a}\in^3$)la</i> -encoding transferable megaplasmids hosted by major hospital-adapted lineages. FEMS Immunology and Medical Microbiology, 2012, 66, 166-176.	2.7	44
58	First outbreak of extendedâ€spectrum βâ€lactamaseâ€producing <i><scp>K</scp>lebsiella pneumoniae</i> in a <scp>N</scp> orwegian neonatal intensive care unit; associated with contaminated breast milk and resolved by strict cohorting. Apmis, 2012, 120, 612-621.	0.9	44
59	Evaluation of Rosco Neoâ€Sensitabs for phenotypic detection and subgrouping of ESBLâ€, AmpC†and carbapenemaseâ€producing Enterobacteriaceae. Apmis, 2012, 120, 724-732.	0.9	23
60	Enterococcus faecalis from patients with chronic periodontitis: virulence and antimicrobial resistance traits and determinants. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 267-272.	1.3	39
61	Multi-Locus Variable Number of Tandem Repeat Analysis for Rapid and Accurate Typing of Virulent Multidrug Resistant Escherichia coli Clones. PLoS ONE, 2012, 7, e41232.	1.1	32
62	Emergence of OXA-carbapenemase- and 16S rRNA methylase-producing international clones of Acinetobacter baumannii in Norway. Journal of Medical Microbiology, 2011, 60, 515-521.	0.7	56
63	A Diversity of OXA-Carbapenemases and Class 1 Integrons Among Carbapenem-Resistant <i>Acinetobacter baumannii</i> Clinical Isolates from Sweden Belonging to Different International Clonal Lineages. Microbial Drug Resistance, 2011, 17, 545-549.	0.9	31
64	The CTX-M Conundrum: Dissemination of Plasmids and <i>Escherichia coli </i> Clones. Microbial Drug Resistance, 2011, 17, 83-97.	0.9	198
65	Structural and Computational Investigations of VIM-7: Insights into the Substrate Specificity of VIM Metallo-β-Lactamases. Journal of Molecular Biology, 2011, 411, 174-189.	2.0	35
66	A sensitive and specific phenotypic assay for detection of metallo- \hat{l}^2 -lactamases and KPC in Klebsiella pneumoniae with the use of meropenem disks supplemented with aminophenylboronic acid, dipicolinic acid and cloxacillin. Clinical Microbiology and Infection, 2011, 17, 552-556.	2.8	178
67	Comparison of disk diffusion, Etest and VITEK2 for detection of carbapenemaseâ€producing Klebsiella pneumoniae with the EUCAST and CLSI breakpoint systems. Clinical Microbiology and Infection, 2011, 17, 668-674.	2.8	54
68	Molecular characterization of VIM-producing Klebsiella pneumoniae from Scandinavia reveals genetic relatedness with international clonal complexes encoding transferable multidrug resistance. Clinical Microbiology and Infection, 2011, 17, 1811-1816.	2.8	70
69	Clustering of polyclonal VanBâ€type vancomycinâ€resistant <i>Enterococcus faecium</i> in a lowâ€endemic area was associated with CC17â€genogroup strains harbouring transferable <i>vanB2</i> and pRUMâ€like <i>repA</i> containing plasmids with <i>axeâ€txe</i> plasmid addiction systems. Apmis, 2011. 119. 247-258.	0.9	21
70	Host range of enterococcal vanA plasmids among Gram-positive intestinal bacteria. Journal of Antimicrobial Chemotherapy, 2011, 66, 273-282.	1.3	55
71	Identification of NDM-1-producing Enterobacteriaceae in Norway. Journal of Antimicrobial Chemotherapy, 2011, 66, 670-672.	1.3	65
72	Retrospective evidence for a biological cost of vancomycin resistance determinants in the absence of glycopeptide selective pressures. Journal of Antimicrobial Chemotherapy, 2011, 66, 608-610.	1.3	51

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7 3	Species identification and molecular characterization of Acinetobacter spp. blood culture isolates from Norway. Journal of Antimicrobial Chemotherapy, 2011, 66, 738-744.	1.3	110
74	Sporadic occurrence of CMY-2-producing multidrug-resistant Escherichia coli of ST-complexes 38 and 448, and ST131 in Norway. Clinical Microbiology and Infection, 2010, 16, 171-178.	2.8	66
7 5	Enterococcus research: recent developments and clinical challenges. Clinical Microbiology and Infection, 2010, 16, 525-526.	2.8	12
76	Mobile genetic elements and their contribution to the emergence of antimicrobial resistant Enterococcus faecalis and Enterococcus faecium. Clinical Microbiology and Infection, 2010, 16, 541-554.	2.8	285
77	PCR-based plasmid typing in <i>Enterococcus faecium</i> strains reveals widely distributed pRE25-, pRUM-, pIP501- and pHTl²-related replicons associated with glycopeptide resistance and stabilizing toxin–antitoxin systems. FEMS Immunology and Medical Microbiology, 2010, 58, 254-268.	2.7	101
78	Tn1546 is part of a larger plasmid-encoded genetic unit horizontally disseminated among clonal Enterococcus faecium lineages. Journal of Antimicrobial Chemotherapy, 2010, 65, 1894-1906.	1.3	56
79	Molecular Epidemiology of Metallo- \hat{l}^2 -Lactamase-Producing <i>Pseudomonas aeruginosa</i> Isolates from Norway and Sweden Shows Import of International Clones and Local Clonal Expansion. Antimicrobial Agents and Chemotherapy, 2010, 54, 346-352.	1.4	136
80	Urinary tract isolates of Enterobacteriaceae from hospitalized patients in the Arkhangelsk region, Russia: Antimicrobial susceptibility and characterization of extended-spectrum beta-lactamases strains. Scandinavian Journal of Infectious Diseases, 2010, 42, 797-800.	1.5	1
81	Plasmid-mediated quinolone resistance determinants qnr and aac(6′)-lb-cr in Escherichia coli and Klebsiella spp. from Norway and Sweden. Diagnostic Microbiology and Infectious Disease, 2010, 66, 425-431.	0.8	66
82	ccrAB Ent serine recombinase genes are widely distributed in the Enterococcus faecium and Enterococcus casseliflavus species groups and are expressed in E. faecium. Microbiology (United) Tj ETQq0 0 0 rg	B ō/ Øverlo	c k 10 Tf 50
83	<i>Enterococcus</i> research - recent developments and clinical challenges. Clinical Microbiology and Infection, 2010, , .	2.8	O
84	The First Metallo- \hat{l}^2 -Lactamase Identified in Norway Is Associated with a TniC-Like Transposon in a Pseudomonas aeruginosa Isolate of Sequence Type 233 Imported from Ghana. Antimicrobial Agents and Chemotherapy, 2009, 53, 331-332.	1.4	26
85	Occurrence, Population Structure, and Antimicrobial Resistance of Enterococci in Marginal and Apical Periodontitis. Journal of Clinical Microbiology, 2009, 47, 2218-2225.	1.8	44
86	Molecular characterization of CTXâ€Mâ€15â€producing clinical isolates of <i>Escherichia coli</i> reveals the spread of multidrugâ€resistant ST131 (O25:H4) and ST964 (O102:H6) strains in Norway. Apmis, 2009, 117, 526-536.	0.9	80
87	Emergence of clonally related Klebsiella pneumoniae isolates of sequence type 258 producing plasmid-mediated KPC carbapenemase in Norway and Sweden. Journal of Antimicrobial Chemotherapy, 2009, 63, 654-658.	1.3	156
88	Factors affecting the reversal of antimicrobial-drug resistance. Lancet Infectious Diseases, The, 2009, 9, 357-364.	4.6	112
89	Clinical isolates of <i>Staphylococcus aureus </i> from the Arkhangelsk region, Russia: antimicrobial susceptibility, molecular epidemiology, and distribution of Pantonâ€Valentine leucocidin genes. Apmis, 2008, 116, 877-887.	0.9	34
90	Complete sequence of Enterococcus faecium pVEF3 and the detection of an ï‰-ε-ζ toxin–antitoxin module and an ABC transporter. Plasmid, 2008, 60, 75-85.	0.4	60

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91	Evaluation of phenotypic tests for the detection of metallo-Â-lactamase-producing Pseudomonas aeruginosa in a low prevalence country. Journal of Antimicrobial Chemotherapy, 2008, 61, 827-830.	1.3	31
92	Redefining extended-spectrum \hat{A} -lactamases: balancing science and clinical need. Journal of Antimicrobial Chemotherapy, 2008, 63, 1-4.	1.3	92
93	The AmpC phenotype in Norwegian clinical isolates of Escherichia coli is associated with an acquired ISEcp1-like ampC element or hyperproduction of the endogenous AmpC. Journal of Antimicrobial Chemotherapy, 2008, 62, 694-702.	1.3	47
94	Alterations of Porin, Pumps, and Penicillin-Binding Proteins in Carbapenem Resistant Clinical Isolates of <i>Pseudomonas aeruginosa</i> . Microbial Drug Resistance, 2008, 14, 23-30.	0.9	33
95	Transfer of plasmid and chromosomal glycopeptide resistance determinants occurs more readily in the digestive tract of mice than in vitro and exconjugants can persist stably in vivo in the absence of glycopeptide selection. Journal of Antimicrobial Chemotherapy, 2007, 59, 478-486.	1.3	59
96	Effects of Phenotype and Genotype on Methods for Detection of Extended-Spectrum-Î ² -Lactamase-Producing Clinical Isolates of Escherichia coli and Klebsiella pneumoniae in Norway. Journal of Clinical Microbiology, 2007, 45, 199-205.	1.8	121
97	Comparative DNA Analysis of Two vanA Plasmids from Enterococcus faecium Strains Isolated from Poultry and a Poultry Farmer in Norway. Antimicrobial Agents and Chemotherapy, 2007, 51, 736-739.	1.4	58
98	VanA-Type Enterococci from Humans, Animals, and Food: Species Distribution, Population Structure, Tn 1546 Typing and Location, and Virulence Determinants. Applied and Environmental Microbiology, 2007, 73, 3307-3319.	1.4	77
99	Nosocomial outbreak of CTX-M-15-producing E. coli in Norway. Apmis, 2007, 115, 120-126.	0.9	41
100	Macrolide-Resistant Streptococcus pyogenes in Norway: Population Structure and Resistance Determinants. Antimicrobial Agents and Chemotherapy, 2006, 50, 1896-1899.	1.4	31
101	Clonal Spread of Macrolide- and Tetracycline-Resistant [erm (A) tet (O)] emm 77 Streptococcus pyogenes Isolates in Italy and Norway. Antimicrobial Agents and Chemotherapy, 2006, 50, 4229-4230.	1.4	10
102	Molecular detection of antibiotic resistance: when and where?. Journal of Antimicrobial Chemotherapy, 2005, 56, 259-261.	1.3	59
103	Persistence of Animal and Human Glycopeptide-Resistant Enterococci on Two Norwegian Poultry Farms Formerly Exposed to Avoparcin Is Associated with a Widespread Plasmid-Mediated vanA Element within a Polyclonal Enterococcus faecium Population. Applied and Environmental Microbiology, 2005, 71, 159-168.	1.4	71
104	Molecular Epidemiology of Macrolide-Resistant Isolates of Streptococcus pneumoniae Collected from Blood and Respiratory Specimens in Norway. Journal of Clinical Microbiology, 2005, 43, 2125-2132.	1.8	20
105	mef(A), mef(E) and a new mef allele in macrolide-resistant Streptococcus spp. isolates from Norway. Journal of Antimicrobial Chemotherapy, 2005, 56, 841-846.	1.3	25
106	Characterization of a carbapenemase-producing clinical isolate of Bacteroides fragilis in Scandinavia: Genetic analysis of a unique insertion sequence. Scandinavian Journal of Infectious Diseases, 2005, 37, 676-679.	1.5	9
107	Phenotypic and genotypic aminoglycoside resistance in blood culture isolates of coagulase-negative staphylococci from a single neonatal intensive care unit, 1989–2000. Journal of Antimicrobial Chemotherapy, 2004, 54, 889-896.	1.3	50
108	Genetic methods for detection of antimicrobial resistance. Apmis, 2004, 112, 815-837.	0.9	119

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109	Nonconjugative Transposition of the vanB -Containing Tn 5382 -Like Element in Enterococcus faecium. Antimicrobial Agents and Chemotherapy, 2003, 47, 786-789.	1.4	29
110	Transferable vanB2 Tn 5382 -Containing Elements in Fecal Streptococcal Strains from Veal Calves. Antimicrobial Agents and Chemotherapy, 2003, 47, 2579-2583.	1.4	34
111	Stability, Persistence, and Evolution of Plasmid-Encoded VanA Glycopeptide Resistance in Enterococci in the Absence of Antibiotic SelectionIn Vitroand in Gnotobiotic Mice. Microbial Drug Resistance, 2002, 8, 161-170.	0.9	69
112	The prevalence of faecal carriage of ampicillin-resistant and high-level gentamicin-resistant enterococci among inpatients at 10 major Norwegian hospitals. Journal of Hospital Infection, 2002, 50, 145-154.	1.4	26
113	Typeability of Tn <i>1546</i> like Elements in Vancomycin-Resistant Enterococci Using Long-Range PCRs and Specific Analysis of Polymorphic Regions. Microbial Drug Resistance, 2000, 6, 49-57.	0.9	41
114	Genetic linkage of the vanB2 gene cluster to Tn5382 in vancomycin-resistant enterococci and characterization of two novel insertion sequences GenBank and GenPept accession numbers are given in Table 3 T3 Microbiology (United Kingdom), 2000, 146, 1469-1479.	0.7	65
115	Heterogeneity in the <i>vanB</i> Gene Cluster of Genomically Diverse Clinical Strains of Vancomycin-Resistant Enterococci. Antimicrobial Agents and Chemotherapy, 1999, 43, 1105-1110.	1.4	118
116	BK and JC Viruses in Patients with Systemic Lupus Erythematosus: Prevalent and Persistent BK Viruria, Sequence Stability of the Viral Regulatory Regions, and Nondetectable Viremia. Journal of Infectious Diseases, 1999, 180, 1-9.	1.9	103
117	Transmission of VanA-Type Vancomycin-Resistant Enterococci andvanAResistance Elements between Chicken and Humans at Avoparcin-Exposed Farms. Microbial Drug Resistance, 1998, 4, 313-318.	0.9	82
118	Intrahospital Spread of Vancomycin-resistant Enterococcus faecium in Sweden. Scandinavian Journal of Infectious Diseases, 1997, 29, 259-263.	1.5	33
119	Nucleic Acid Amplification Techniques in Detection and Diagnosis of Medically Important Viral Infections. , 1997, , 183-199.		2
120	Experimental expression in mice and spontaneous expression in human SLE of polyomavirus T-antigen. A molecular basis for induction of antibodies to DNA and eukaryotic transcription factors Journal of Clinical Investigation, 1997, 99, 2045-2054.	3.9	106
121	On the biological origin of anti-double-stranded (ds)DNA antibodies: systemic lupus erythematosus-related anti-dsDNA antibodies are induced by polyomavirus BK in lupus-prone (NZBxNZW) F1 hybrids, but not in normal mice. European Journal of Immunology, 1994, 24, 66-70.	1.6	53
122	Amplification and sequencing of the control regions of BK and JC virus from human urine by polymerase chain reaction. Virology, 1991, 180, 553-560.	1.1	163
123	BK virus early RNA transcripts in stably transformed cells: enhanced levels induced by dibutyryl cyclic AMP, forskolin and 12-O-tetradecanoylphorbol-13-acetate treatment. Journal of General Virology, 1990, 71, 1461-1471.	1.3	22
124	Peripheral Facial Palsy and Coincidental Cytomegalovirus Infection or Reactivation. Scandinavian Journal of Infectious Diseases, 1983, 15, 233-238.	1.5	27