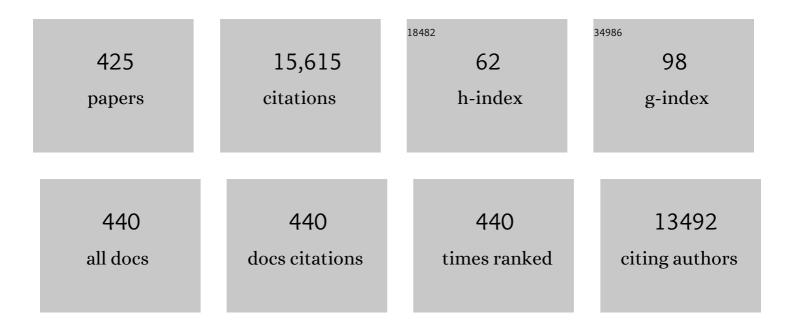
List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3316357/publications.pdf Version: 2024-02-01



#	ARTICLE	IF	CITATIONS
1	proposal of Cronobacter sakazakii gen. nov., comb. nov., Cronobacter malonaticus sp. nov., Cronobacter turicensis sp. nov., Cronobacter muytjensii sp. nov., Cronobacter dublinensis sp. nov., Cronobacter genomospecies 1, and of three subspecies, Cronobacter dublinensis subsp. dublinensis subsp. nov., Cronobacter dublinensis subsp. lausannensis subsp. nov. and Cronobacter dublinensis	1.7	506
2	subsp. lactaridi subsp. nov International Journal of Systematic and Evolutionary Microbiology, 2008, Campylobacter. Veterinary Research, 2005, 36, 351-382.	3.0	389
3	LowWater Activity Foods: Increased Concern as Vehicles of Foodborne Pathogens. Journal of Food Protection, 2013, 76, 150-172.	1.7	355
4	The taxonomy of Enterobacter sakazakii: proposal of a new genus Cronobacter gen. nov. and descriptions of Cronobacter sakazakii comb. nov. Cronobacter sakazakii subsp. sakazakii, comb. nov., Cronobacter sakazakii subsp. malonaticus subsp. nov., Cronobacter turicensis sp. nov., Cronobacter muytjensii sp. nov., Cronobacter dublinensis sp. nov. and Cronobacter genomospecies 1. BMC Evolutionary Biology, 2007, 7, 64.	3.2	275
5	vi>Cronobacter( <i>Enterobacter sakazakii</i> ): An Opportunistic Foodborne Pathogen. Foodborne Pathogens and Disease, 2010, 7, 339-350.	1.8	250
6	A Review on the Applications of Next Generation Sequencing Technologies as Applied to Food-Related Microbiome Studies. Frontiers in Microbiology, 2017, 8, 1829.	3.5	245
7	Mechanisms of survival, responses and sources of Salmonella in low-moisture environments. Frontiers in Microbiology, 2013, 4, 331.	3.5	242
8	The ins and outs of RND efflux pumps in Escherichia coli. Frontiers in Microbiology, 2015, 6, 587.	3.5	219
9	Enterobacter sakazakii: An Emerging Pathogen in Powdered Infant Formula. Clinical Infectious Diseases, 2006, 42, 996-1002.	5.8	206
10	Methicillin-resistant Staphylococcus aureus (MRSA) isolated from animals and veterinary personnel in Ireland. Veterinary Microbiology, 2005, 109, 285-296.	1.9	197
11	The epidemiology of antibiotic resistance in Campylobacter. Microbes and Infection, 2006, 8, 1955-1966.	1.9	192
12	Mechanisms of fluoroquinolone and macrolide resistance in Campylobacter spp Microbes and Infection, 2006, 8, 1967-1971.	1.9	176
13	Toxin A-negative, toxin B-positive Clostridium difficile. International Journal of Infectious Diseases, 2007, 11, 5-10.	3.3	155
14	Efficacy of Biocides Used in the Modern Food Industry To Control Salmonella enterica, and Links between Biocide Tolerance and Resistance to Clinically Relevant Antimicrobial Compounds. Applied and Environmental Microbiology, 2012, 78, 3087-3097.	3.1	151
15	Occurrence of Campylobacter in retail foods in Ireland. International Journal of Food Microbiology, 2004, 95, 111-118.	4.7	150
16	Development and evaluation of rpoB based PCR systems to differentiate the six proposed species within the genus Cronobacter. International Journal of Food Microbiology, 2009, 136, 165-168.	4.7	149
17	Antimicrobial resistance and its association with tolerance to heavy metals in agriculture production. Food Microbiology, 2017, 64, 23-32.	4.2	138
18	Re-examination of the taxonomic status of Enterobacter helveticus, Enterobacter pulveris and Enterobacter turicensis as members of the genus Cronobacter and their reclassification in the genera Franconibacter gen. nov. and Siccibacter gen. nov. as Franconibacter helveticus comb. nov., Franconibacter pulveris comb. nov. and Siccibacter turicensis comb. nov., respectively. International Journal of Systematic and Evolutionary Microbiology, 2014, 64, 3402-3410.	1.7	136

#	Article	IF	CITATIONS
19	Cronobacter species (formerly known as Enterobacter sakazakii) in powdered infant formula: a review of our current understanding of the biology of this bacterium. Journal of Applied Microbiology, 2012, 113, 1-15.	3.1	128
20	Potential role of non-antibiotics (helper compounds) in the treatment of multidrug-resistant Gram-negative infections: mechanisms for their direct and indirect activities. International Journal of Antimicrobial Agents, 2008, 31, 198-208.	2.5	124
21	Salmonellaââ,¬â€œHost Interactions ââ,¬â€œ Modulation of the Host Innate Immune System. Frontiers in Immunology, 2014, 5, 481.	4.8	124
22	Characterisation and transferability of antibiotic resistance genes from lactic acid bacteria isolated from Irish pork and beef abattoirs. Research in Microbiology, 2010, 161, 127-135.	2.1	115
23	Microbial detection and identification methods: Bench top assays to omics approaches. Comprehensive Reviews in Food Science and Food Safety, 2020, 19, 3106-3129.	11.7	115
24	Emergence and Control of Fluoroquinolone-Resistant, Toxin A–Negative, Toxin B–Positive <i>Clostridium difficile</i> . Infection Control and Hospital Epidemiology, 2007, 28, 932-940.	1.8	114
25	Emergence of a Globally Dominant IncHI1 Plasmid Type Associated with Multiple Drug Resistant Typhoid. PLoS Neglected Tropical Diseases, 2011, 5, e1245.	3.0	114
26	Application of pulsed-field gel electrophoresis to characterise and trace the prevalence of Enterobacter sakazakii in an infant formula processing facility. International Journal of Food Microbiology, 2007, 116, 73-81.	4.7	112
27	Antimicrobial Resistance in Foodborne Pathogens - A Cause for Concern?. Current Drug Targets, 2008, 9, 808-815.	2.1	111
28	Whole-genome Sequencing to Track Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Transmission in Nosocomial Outbreaks. Clinical Infectious Diseases, 2021, 72, e727-e735.	5.8	107
29	A Simple Method for Assessment of MDR Bacteria for Over-Expressed Efflux Pumps. Open Microbiology Journal, 2013, 7, 72-82.	0.7	97
30	Serotypes and Virulence Profiles of Non-O157 Shiga Toxin-Producing Escherichia coli Isolates from Bovine Farms. Applied and Environmental Microbiology, 2011, 77, 8662-8668.	3.1	96
31	Antimicrobial Resistance in Wildlife: Implications for Public Health. Zoonoses and Public Health, 2015, 62, 534-542.	2.2	96
32	Multiple Regulatory Pathways Associated with High-Level Ciprofloxacin and Multidrug Resistance in <i>Salmonella enterica</i> Serovar Enteritidis: Involvement of <i>ramA</i> and Other Global Regulators. Antimicrobial Agents and Chemotherapy, 2009, 53, 1080-1087.	3.2	95
33	Inhibition of verocytotoxigenic Escherichia coli in model broth and rumen systems by carvacrol and thymol. International Journal of Food Microbiology, 2010, 139, 70-78.	4.7	95
34	Elucidation of the RamA Regulon in Klebsiella pneumoniae Reveals a Role in LPS Regulation. PLoS Pathogens, 2015, 11, e1004627.	4.7	95
35	Neutral Genomic Microevolution of a Recently Emerged Pathogen, Salmonella enterica Serovar Agona. PLoS Genetics, 2013, 9, e1003471.	3.5	94
36	Identification of " <i>Cronobacter</i> ―spp. ( <i>Enterobacter sakazakii</i> ). Journal of Clinical Microbiology, 2007, 45, 3814-3816.	3.9	93

#	Article	IF	CITATIONS
37	Molecular Analysis of the <i>Enterobacter sakazakii</i> O-Antigen Gene Locus. Applied and Environmental Microbiology, 2008, 74, 3783-3794.	3.1	92
38	Dissemination of <i>Cronobacter</i> spp. ( <i>Enterobacter sakazakii</i> ) in a Powdered Milk Protein Manufacturing Facility. Applied and Environmental Microbiology, 2008, 74, 5913-5917.	3.1	91
39	Development of a real-time multiplex PCR assay for the detection of multiple Salmonella serotypes in chicken samples. BMC Microbiology, 2008, 8, 156.	3.3	88
40	Prevalence and Characterization of Staphylococcus aureus Cultured From Raw Milk Taken From Dairy Cows With Mastitis in Beijing, China. Frontiers in Microbiology, 2018, 9, 1123.	3.5	88
41	High-level resistance to moxifloxacin and gatifloxacin associated with a novel mutation in gyrB in toxin-A-negative, toxin-B-positive Clostridium difficile. Journal of Antimicrobial Chemotherapy, 2006, 58, 1264-1267.	3.0	87
42	Characterization of a collection of Enterobacter sakazakii isolates from environmental and food sources. International Journal of Food Microbiology, 2006, 110, 127-134.	4.7	84
43	An investigation of the molecular mechanisms contributing to high-level erythromycin resistance in Campylobacter. International Journal of Antimicrobial Agents, 2006, 27, 40-45.	2.5	83
44	Characterization of Multidrug-Resistant Escherichia coli Isolates from Animals Presenting at a University Veterinary Hospital. Applied and Environmental Microbiology, 2011, 77, 7104-7112.	3.1	83
45	Whole-Genome Sequencing-Based Characterization of 100 Listeria monocytogenes Isolates Collected from Food Processing Environments over a Four-Year Period. MSphere, 2019, 4, .	2.9	82
46	<i>Yersinia Enterocolitica:</i> A Brief Review of the Issues Relating to the Zoonotic Pathogen, Public Health Challenges, and the Pork Production Chain. Foodborne Pathogens and Disease, 2012, 9, 179-189.	1.8	81
47	Development and application of Multiple-Locus Variable Number of tandem repeat Analysis (MLVA) to subtype a collection of Listeria monocytogenes. International Journal of Food Microbiology, 2007, 115, 187-194.	4.7	80
48	ProP Is Required for the Survival of Desiccated Salmonella enterica Serovar Typhimurium Cells on a Stainless Steel Surface. Applied and Environmental Microbiology, 2013, 79, 4376-4384.	3.1	80
49	Pan-genome analysis of the emerging foodborne pathogen Cronobacter spp. suggests a species-level bidirectional divergence driven by niche adaptation. BMC Genomics, 2013, 14, 366.	2.8	78
50	Influences on antimicrobial prescribing behaviour of veterinary practitioners in cattle practice in Ireland. Veterinary Record, 2013, 172, 14-14.	0.3	78
51	Surveillance and characterisation by Pulsed-Field Gel Electrophoresis of Cronobacter spp. in farming and domestic environments, food production animals and retail foods. International Journal of Food Microbiology, 2009, 136, 198-203.	4.7	75
52	Emergence and Diversity of Salmonella enterica Serovar Indiana Isolates with Concurrent Resistance to Ciprofloxacin and Cefotaxime from Patients and Food-Producing Animals in China. Antimicrobial Agents and Chemotherapy, 2016, 60, 3365-3371.	3.2	75
53	Vertical transmission of highly similar blaCTX-M-1-harboring Incl1 plasmids in Escherichia coli with different MLST types in the poultry production pyramid. Frontiers in Microbiology, 2014, 5, 519.	3.5	74
54	SIRT1-mediated FoxOs pathways protect against apoptosis by promoting autophagy in osteoblast-like MC3T3-E1 cells exposed to sodium fluoride. Oncotarget, 2016, 7, 65218-65230.	1.8	74

#	Article	IF	CITATIONS
55	Antimicrobial Resistance and Genetic Diversity of Shigella sonnei Isolates from Western Ireland, an Area of Low Incidence of Infection. Journal of Clinical Microbiology, 2003, 41, 1919-1924.	3.9	70
56	Molecular Characterization of Multidrug-Resistant Escherichia coli Isolates from Irish Cattle Farms. Applied and Environmental Microbiology, 2011, 77, 7121-7127.	3.1	68
57	Nucleotide sequences of 16 transmissible plasmids identified in nine multidrug-resistant Escherichia coli isolates expressing an ESBL phenotype isolated from food-producing animals and healthy humans. Journal of Antimicrobial Chemotherapy, 2014, 69, 2658-2668.	3.0	68
58	Wastewater is a reservoir for clinically relevant carbapenemase- and 16s rRNA methylase-producing Enterobacteriaceae. International Journal of Antimicrobial Agents, 2017, 50, 436-440.	2.5	68
59	Isolation and characterisation of toxin A-negative, toxin B-positive Clostridium difficile in Dublin, Ireland. Clinical Microbiology and Infection, 2007, 13, 298-304.	6.0	67
60	Characterization of Cronobacter recovered from dried milk and related products. BMC Microbiology, 2009, 9, 24.	3.3	66
61	Mortality in patients with Clostridium difficile infection correlates with host pro-inflammatory and humoral immune responses. Journal of Medical Microbiology, 2013, 62, 1453-1460.	1.8	66
62	The prevalence of <i>Cryptosporidium</i> species and subtypes in human faecal samples in Ireland. Epidemiology and Infection, 2009, 137, 270-277.	2.1	65
63	Molecular characterization of blaESBL–harboring conjugative plasmids identified in multi-drug resistant Escherichia coli isolated from food-producing animals and healthy humans. Frontiers in Microbiology, 2013, 4, 188.	3.5	65
64	<i>gyrA</i> Mutations in Fluoroquinolone-resistant <i>Clostridium difficile</i> PCR-027. Emerging Infectious Diseases, 2007, 13, 504-505.	4.3	64
65	Fitness Costs and Stability of a High-Level Ciprofloxacin Resistance Phenotype in <i>Salmonella enterica</i> Serotype Enteritidis: Reduced Infectivity Associated with Decreased Expression of <i>Salmonella</i> Pathogenicity Island 1 Genes. Antimicrobial Agents and Chemotherapy, 2010, 54, 367-374.	3.2	64
66	Enterotoxigenicity and Antimicrobial Resistance of Staphylococcus aureus Isolated from Retail Food in China. Frontiers in Microbiology, 2017, 8, 2256.	3.5	63
67	Prevalence ofCryptosporidiumspecies in intensively farmed pigs in Ireland. Parasitology, 2007, 134, 1575-1582.	1.5	62
68	A 3-year multi-food study of the presence and persistence of Listeria monocytogenes in 54 small food businesses in Ireland. International Journal of Food Microbiology, 2017, 249, 18-26.	4.7	62
69	Complete genome sequence and phenotype microarray analysis of Cronobacter sakazakii SP291: a persistent isolate cultured from a powdered infant formula production facility. Frontiers in Microbiology, 2013, 4, 256.	3.5	61
70	Tracking the Salmonella Status of Pigs and Pork from Lairage through the Slaughter Process in the Republic of Ireland. Journal of Food Protection, 2010, 73, 2148-2160.	1.7	60
71	Mechanisms of Fluoroquinolone Resistance in Escherichia coli Isolates from Food-Producing Animals. Applied and Environmental Microbiology, 2011, 77, 7113-7120.	3.1	60
72	Transfer of Antibiotic Resistance Marker Genes between Lactic Acid Bacteria in Model Rumen and Plant Environments. Applied and Environmental Microbiology, 2009, 75, 3146-3152.	3.1	59

#	Article	IF	CITATIONS
73	Detection of plasmid-mediated tigecycline-resistant gene tet(X4) in Escherichia coli from pork, Sichuan and Shandong Provinces, China, February 2019. Eurosurveillance, 2019, 24, .	7.0	59
74	Antimicrobial resistance islands: resistance gene clusters in Salmonella chromosome and plasmids. Microbes and Infection, 2006, 8, 1923-1930.	1.9	57
75	Prevalence of thermophilic <i>Campylobacter</i> species in household cats and dogs in Ireland. Veterinary Record, 2009, 164, 44-47.	0.3	57
76	Efficacy of UV Light Treatment for the Microbiological Decontamination of Chicken, Associated Packaging, and Contact Surfaces. Journal of Food Protection, 2011, 74, 565-572.	1.7	57
77	Serotypes and virulotypes of non-O157 shiga-toxin producing Escherichia coli (STEC) on bovine hides and carcasses. Food Microbiology, 2012, 32, 223-229.	4.2	57
78	Prevalence, numbers and characteristics of Salmonella spp. on Irish retail pork. International Journal of Food Microbiology, 2009, 131, 233-239.	4.7	56
79	Detection and characterisation of group A rotavirus in asymptomatic piglets in southern Ireland. Archives of Virology, 2010, 155, 1247-1259.	2.1	56
80	Characterization of antimicrobial resistance in Salmonella enterica food and animal isolates from Colombia: identification of a qnrB19-mediated quinolone resistance marker in two novel serovars. FEMS Microbiology Letters, 2010, 313, 10-19.	1.8	55
81	Does Microbicide Use in Consumer Products Promote Antimicrobial Resistance? A Critical Review and Recommendations for a Cohesive Approach to Risk Assessment. Microbial Drug Resistance, 2013, 19, 344-354.	2.0	54
82	Molecular Characterization of Rotavirus in Ireland: Detection of Novel Strains Circulating in the Population. Journal of Clinical Microbiology, 2000, 38, 3370-3374.	3.9	54
83	Genomic Comparisons of Salmonella enterica Serovar Dublin, Agona, and Typhimurium Strains Recently Isolated from Milk Filters and Bovine Samples from Ireland, Using a Salmonella Microarray. Applied and Environmental Microbiology, 2005, 71, 1616-1625.	3.1	53
84	pH Modulation of Efflux Pump Activity of Multi-Drug Resistant Escherichia coli: Protection During Its Passage and Eventual Colonization of the Colon. PLoS ONE, 2009, 4, e6656.	2.5	53
85	Susceptibility of Campylobacter to high intensity near ultraviolet/visible 395±5nm light and its effectiveness for the decontamination of raw chicken and contact surfaces. International Journal of Food Microbiology, 2012, 159, 267-273.	4.7	53
86	Assessment of Antimicrobial Resistance Transfer Between Lactic Acid Bacteria and Potential Foodborne Pathogens Using <i>In Vitro </i> Methods and Mating in a Food Matrix. Foodborne Pathogens and Disease, 2009, 6, 925-933.	1.8	52
87	Efficacy of High-Intensity Pulsed Light for the Microbiological Decontamination of Chicken, Associated Packaging, and Contact Surfaces. Foodborne Pathogens and Disease, 2011, 8, 109-117.	1.8	52
88	Enterobacter sakazakii an emerging bacterial pathogen with implications for infant health. Minerva Pediatrica, 2007, 59, 137-48.	2.7	52
89	Characterization and Chromosomal Mapping of Antimicrobial Resistance Genes in Salmonella enterica Serotype Typhimurium. Applied and Environmental Microbiology, 2000, 66, 4842-4848.	3.1	51
90	Development and Application of a Novel Peptide Nucleic Acid Probe for the Specific Detection of <i>Cronobacter</i> Genomospecies ( <i>Enterobacter sakazakii</i> ) in Powdered Infant Formula. Applied and Environmental Microbiology, 2009, 75, 2925-2930.	3.1	51

#	Article	IF	CITATIONS
91	Prevalence of thermophilic Campylobacter species in cats and dogs in two animal shelters in Ireland. Veterinary Record, 2006, 158, 51-54.	0.3	50
92	Deteminants of cross ontamination during home food preparation. British Food Journal, 2011, 113, 280-297.	2.9	50
93	Elucidating the Regulon of Multidrug Resistance Regulator RarA in Klebsiella pneumoniae. Antimicrobial Agents and Chemotherapy, 2013, 57, 1603-1609.	3.2	50
94	Antimicrobial resistant bacteria in wild mammals and birds: a coincidence or cause for concern?. Irish Veterinary Journal, 2014, 67, 8.	2.1	50
95	Molecular characterisation and analysis of bovine rotavirus strains circulating in Ireland 2002–2004. Veterinary Microbiology, 2006, 117, 242-247.	1.9	47
96	An Original Deal for New Molecule: Reversal of Efflux Pump Activity, A Rational Strategy to Combat Gram-Negative Resistant Bacteria. Current Medicinal Chemistry, 2011, 18, 2969-2980.	2.4	47
97	Molecular Analysis of <i>Pseudomonas aeruginosa</i> : Epidemiological Investigation of Mastitis Outbreaks in Irish Dairy Herds. Applied and Environmental Microbiology, 1999, 65, 2723-2729.	3.1	47
98	Phylogenetic Profiles of In-House Microflora in Drains at a Food Production Facility: Comparison and Biocontrol Implications of Listeria-Positive and -Negative Bacterial Populations. Applied and Environmental Microbiology, 2014, 80, 3369-3374.	3.1	46
99	Multi-Drug Resistance in Salmonella enterica: Efflux Mechanisms and Their Relationships with the Development of Chromosomal Resistance Gene Clusters. Current Drug Targets, 2006, 7, 849-860.	2.1	45
100	Survival characteristics of environmental and clinically derived strains of Cronobacter sakazakii in in infant milk formula (IMF) and ingredients. Journal of Applied Microbiology, 2011, 110, 697-703.	3.1	44
101	Characterisation of multidrug-resistant Shiga toxin-producing Escherichia coli cultured from pigs in China: co-occurrence of extended-spectrum l²-lactamase- and mcr-1-encoding genes on plasmids. International Journal of Antimicrobial Agents, 2016, 48, 445-448.	2.5	44
102	Molecular Characterization of Irish Salmonella enterica Serotype Typhimurium: Detection of Class I Integrons and Assessment of Genetic Relationships by DNA Amplification Fingerprinting. Applied and Environmental Microbiology, 2000, 66, 614-619.	3.1	43
103	Development and validation of a rapid real-time PCR based method for the specific detection of Salmonella on fresh meat. Meat Science, 2009, 83, 555-562.	5.5	43
104	Antimicrobial Resistance of Faecal <i>Escherichia coli</i> Isolates from Pig Farms with Different Durations of Inâ€feed Antimicrobial Use. Zoonoses and Public Health, 2016, 63, 241-250.	2.2	43
105	Genomic diversity of Salmonella enterica -The UoWUCC 10K genomes project. Wellcome Open Research, 2020, 5, 223.	1.8	43
106	Trends in antimicrobial susceptibility among isolates of <i>Campylobacter</i> species in Ireland and the emergence of resistance to ciprofloxacin. Veterinary Record, 2002, 151, 317-320.	0.3	42
107	Detection of Enterobacter sakazakii in Dried Infant Milk Formula by Cationic-Magnetic-Bead Capture. Applied and Environmental Microbiology, 2006, 72, 6325-6330.	3.1	42
108	Transfer of ampicillin resistance from Salmonella Typhimurium DT104 to Escherichia coli K12 in food. Letters in Applied Microbiology, 2008, 46, 210-215.	2.2	42

#	Article	IF	CITATIONS
109	Changing profile of the bovine rotavirus G6 population in the south of Ireland from 2002 to 2009. Veterinary Microbiology, 2010, 146, 238-244.	1.9	42
110	The potential for biocide tolerance in Escherichia coli and its impact on the response to food processing stresses. Food Control, 2012, 26, 98-106.	5.5	42
111	Characterization of Five Escherichia coli Isolates Co-expressing ESBL and MCR-1 Resistance Mechanisms From Different Origins in China. Frontiers in Microbiology, 2019, 10, 1994.	3.5	42
112	Efflux Pumps of Gramâ€Negative Bacteria: Genetic Responses to Stress and the Modulation of their Activity by pH, Inhibitors, and Phenothiazines. Advances in Enzymology and Related Areas of Molecular Biology, 2011, 77, 61-108.	1.3	41
113	Incorporation of commercially-derived antimicrobials into gelatin-based films and assessment of their antimicrobial activity and impact on physical film properties. Food Control, 2016, 64, 202-211.	5.5	41
114	Identification of efflux pump-mediated multidrug-resistant bacteria by the ethidium bromide-agar cartwheel method. In Vivo, 2011, 25, 171-8.	1.3	41
115	<i>Enterobacter sakazakii</i> survives spray drying. International Journal of Dairy Technology, 2008, 61, 102-108.	2.8	40
116	Cronobacter: An Emergent Pathogen Causing Meningitis to Neonates through their Feeds. Science Progress, 2014, 97, 154-172.	1.9	40
117	RNA Sequencing-Based Transcriptional Overview of Xerotolerance in Cronobacter sakazakii SP291. Applied and Environmental Microbiology, 2019, 85, .	3.1	40
118	Molecular Epidemiology of Campylobacter Isolates from Poultry Production Units in Southern Ireland. PLoS ONE, 2011, 6, e28490.	2.5	40
119	Emergence of G3 and G9 rotavirus and increased incidence of mixed infections in the southern region of Ireland 2001-2004. Journal of Medical Virology, 2005, 77, 571-578.	5.0	39
120	Identification of critical points during domestic food preparation: an observational study. British Food Journal, 2011, 113, 766-783.	2.9	39
121	An Evaluation of the Potential of High-Intensity Ultrasound for Improving the Microbial Safety of Poultry. Food and Bioprocess Technology, 2012, 5, 992-998.	4.7	39
122	Comparative Genotypic and Phenotypic Analysis of Cronobacter Species Cultured from Four Powdered Infant Formula Production Facilities: Indication of Pathoadaptation along the Food Chain. Applied and Environmental Microbiology, 2015, 81, 4388-4402.	3.1	39
123	Comparison of multidrug resistance gene regions between two geographically unrelated Salmonella serotypes. Journal of Antimicrobial Chemotherapy, 2005, 55, 558-561.	3.0	38
124	Enterobacter sakazakii: biological properties and significance in dried infant milk formula (IMF) powder+. International Journal of Dairy Technology, 2006, 59, 102-111.	2.8	38
125	PCR ribotype prevalence and molecular basis of macrolide-lincosamide-streptogramin B (MLSB) and fluoroquinolone resistance in Irish clinical Clostridium difficile isolates. Journal of Antimicrobial Chemotherapy, 2011, 66, 1976-1982.	3.0	38
126	Acid environments affect biofilm formation and gene expression in isolates of Salmonella enterica Typhimurium DT104. International Journal of Food Microbiology, 2015, 206, 7-16.	4.7	38

#	Article	IF	CITATIONS
127	Complete genetic analysis of a Salmonella enterica serovar Indiana isolate accompanying four plasmids carrying mcr-1, ESBL and other resistance genes in China. Veterinary Microbiology, 2017, 210, 142-146.	1.9	38
128	First Report of Klebsiella oxytoca Strain Simultaneously Producing NDM-1, IMP-4, and KPC-2 Carbapenemases. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	38
129	Identification and Characterization of Five New Molecular Serogroups of Cronobacter spp Foodborne Pathogens and Disease, 2013, 10, 343-352.	1.8	37
130	Surface attachment of active antimicrobial coatings onto conventional plastic-based laminates and performance assessment of these materials on the storage life of vacuum packaged beef sub-primals. Food Microbiology, 2017, 62, 196-201.	4.2	37
131	<i>Klebsiella pneumoniae</i> : Prevalence, Reservoirs, Antimicrobial Resistance, Pathogenicity, and Infection: A Hitherto Unrecognized Zoonotic Bacterium. Foodborne Pathogens and Disease, 2021, 18, 63-84.	1.8	36
132	Detection of numerous verotoxigenic E. coli serotypes, with multiple antibiotic resistance from cattle faeces and soil. Veterinary Microbiology, 2009, 134, 288-293.	1.9	35
133	Comparative proteomic analysis of Salmonella tolerance to the biocide active agent triclosan. Journal of Proteomics, 2012, 75, 4505-4519.	2.4	35
134	Detection of Escherichia coli O157 by Peptide Nucleic Acid Fluorescence <i>In Situ</i> Hybridization (PNA-FISH) and Comparison to a Standard Culture Method. Applied and Environmental Microbiology, 2013, 79, 6293-6300.	3.1	35
135	Exploring the Genome and Phenotype of Multi-Drug Resistant Klebsiella pneumoniae of Clinical Origin. Frontiers in Microbiology, 2017, 8, 1913.	3.5	35
136	Genomic insights into persistence of Listeria species in the food processing environment. Journal of Applied Microbiology, 2021, 131, 2082-2094.	3.1	35
137	Rotavirus in Ireland: national estimates of disease burden, 1997 to 1998. Pediatric Infectious Disease Journal, 2001, 20, 693-698.	2.0	35
138	Phenothiazines, bacterial efflux pumps and targeting the macrophage for enhanced killing of intracellular XDRTB. In Vivo, 2010, 24, 409-24.	1.3	35
139	Comparative Genomics of the Listeria monocytogenes ST204 Subgroup. Frontiers in Microbiology, 2016, 7, 2057.	3.5	34
140	Investigation of in-feed organic acids as a low cost strategy to combat Salmonella in grower pigs. Preventive Veterinary Medicine, 2017, 139, 50-57.	1.9	34
141	Prevalence of <i>Cronobacter</i> species ( <i>Enterobacter sakazakii</i> ) in follow-on infant formulae and infant drinks. Letters in Applied Microbiology, 2009, 48, 536-541.	2.2	33
142	A Stereoselective Switch: Enantiodivergent Approach to the Synthesis of Isoflavanones. Chemistry - A European Journal, 2014, 20, 15354-15359.	3.3	33
143	Prevalence and numbers of <i>Salmonella</i> spp. and Enterobacteriaceae on pork cuts in abattoirs in the Republic of Ireland. Journal of Applied Microbiology, 2008, 105, 1209-1219.	3.1	32
144	Ethidium bromide efflux by Salmonella: modulation by metabolic energy, pH, ions and phenothiazines. International Journal of Antimicrobial Agents, 2011, 38, 140-145.	2.5	32

#	Article	IF	CITATIONS
145	Incidence and survival of non-O157 verocytotoxigenic Escherichia coli in soil. Journal of Applied Microbiology, 2011, 111, 484-490.	3.1	32
146	Antimicrobial Resistance in <i>Listeria</i> Species. Microbiology Spectrum, 2018, 6, .	3.0	32
147	A scoping review on the prevalence of Shigaâ€ŧoxigenic <i>Escherichia coli </i> in wild animal species. Zoonoses and Public Health, 2018, 65, 911-920.	2.2	32
148	Antimicrobial Resistance in NontyphoidalSalmonellafrom Food Sources in Colombia: Evidence for an Unusual Plasmid-Localized Class 1 Integron in Serotypes Typhimurium and Anatum. Microbial Drug Resistance, 2006, 12, 269-277.	2.0	31
149	Antibiotic-resistant Campylobacter: could efflux pump inhibitors control infection?. Journal of Antimicrobial Chemotherapy, 2007, 59, 1230-1236.	3.0	31
150	Efficacy of pulsed electric fields for the inactivation of indicator microorganisms and foodborne pathogens in liquids and raw chicken. Food Control, 2012, 25, 131-135.	5.5	31
151	A Preliminary Study of <i>Salmonella</i> , Verocytotoxigenic <i>Escherichia coli</i> / <i>Escherichia coli</i> /on Four Mixed Farms. Zoonoses and Public Health, 2012, 59, 217-228.	2.2	31
152	A multiplex real-time PCR assay for the identification and differentiation of Salmonella enterica serovar Typhimurium and monophasic serovar 4,[5],12:i:–. International Journal of Food Microbiology, 2013, 166, 48-53.	4.7	31
153	Cronobacter spp.—Opportunistic Foodborne Pathogens: an Update on Evolution, Osmotic Adaptation and Pathogenesis. Current Clinical Microbiology Reports, 2018, 5, 97-105.	3.4	31
154	Novel IS26-mediated hybrid plasmid harbouring tet(X4) in Escherichia coli. Journal of Global Antimicrobial Resistance, 2020, 21, 162-168.	2.2	31
155	Antimicrobial resistance profiling andfla-typing of Irish thermophillicCampylobacterspp. of human and poultry origin. Letters in Applied Microbiology, 2006, 43, 560-565.	2.2	30
156	An Investigation into the Efficacy of Washing Trucks Following the Transportation of Pigs—A <i>Salmonella</i> Perspective. Foodborne Pathogens and Disease, 2008, 5, 261-271.	1.8	30
157	Molecular Characterization and Environmental Mapping of <i>Campylobacter</i> Isolates in a Subset of Intensive Poultry Flocks in Ireland. Foodborne Pathogens and Disease, 2011, 8, 99-108.	1.8	30
158	Microarray-based comparative genomic indexing of the Cronobacter genus (Enterobacter sakazakii). International Journal of Food Microbiology, 2009, 136, 159-164.	4.7	29
159	A Comparison of Different Culture Methods for the Recovery of <i>Campylobacter</i> Species from Pets. Zoonoses and Public Health, 2009, 56, 490-495.	2.2	29
160	Development and Validation of a PulseNet Standardized Protocol for Subtyping Isolates of <i>Cronobacter</i> Species. Foodborne Pathogens and Disease, 2012, 9, 861-867.	1.8	29
161	A proposed harmonized LPS molecular-subtyping scheme for Cronobacter species. Food Microbiology, 2015, 50, 38-43.	4.2	29
162	Genomic characterization of a large plasmid containing a bla NDM-1 gene carried on Salmonella enterica serovar Indiana C629 isolate from China. BMC Infectious Diseases, 2017, 17, 479.	2.9	29

#	Article	IF	CITATIONS
163	Draft genomes of Cronobacter sakazakii strains isolated from dried spices bring unique insights into the diversity of plant-associated strains. Standards in Genomic Sciences, 2018, 13, 35.	1.5	29
164	The Secretion of Toxins and Other Exoproteins of Cronobacter: Role in Virulence, Adaption, and Persistence. Microorganisms, 2020, 8, 229.	3.6	29
165	An instrument-free method for the demonstration of efflux pump activity of bacteria. In Vivo, 2006, 20, 657-64.	1.3	29
166	Development of Multiple-Locus Variable-Number Tandem-Repeat Analysis for the Molecular Subtyping of <i>Enterobacter sakazakii</i> . Applied and Environmental Microbiology, 2008, 74, 1223-1231.	3.1	28
167	Clindamycin-Resistant Clone ofClostridium difficilePCR Ribotype 027, Europe. Emerging Infectious Diseases, 2008, 14, 1485-1487.	4.3	28
168	Application of multiple locus variable number of tandem repeat analysis (MLVA), phage typing and antimicrobial susceptibility testing to subtype Salmonella enterica serovar Typhimurium isolated from pig farms, pork slaughterhouses and meat producing plants in Ireland. Food Microbiology, 2011, 28, 1087-1094.	4.2	28
169	Identification and genetic characterization of a novel picornavirus from chickens. Journal of General Virology, 2014, 95, 1094-1103.	2.9	28
170	Longitudinal Study of Two Irish Dairy Herds: Low Numbers of Shiga Toxin-Producing Escherichia coli O157 and O26 Super-Shedders Identified. Frontiers in Microbiology, 2016, 7, 1850.	3.5	28
171	Analysis and Characterization of Proteins Associated with Outer Membrane Vesicles Secreted by Cronobacter spp Frontiers in Microbiology, 2017, 8, 134.	3.5	28
172	<i>Campylobacter</i> spp. in Irish Feedlot Cattle: A Longitudinal Study Involving Preâ€harvest and Harvest Phases of the Food Chain. Zoonoses and Public Health, 2004, 51, 28-33.	1.4	27
173	Antimicrobial resistance in Irish isolates of verocytotoxigenic Escherichia coli (E. coli)—VTEC. International Journal of Food Microbiology, 2006, 109, 173-178.	4.7	27
174	Comparison of Listeria monocytogenes strain types in Irish smoked salmon and other foods. International Journal of Hygiene and Environmental Health, 2006, 209, 527-534.	4.3	27
175	Changing profile of rotavirus in Ireland: Predominance of P[8] and emergence of P[6] and P[9] in mixed infections. Journal of Medical Virology, 2008, 80, 524-530.	5.0	27
176	An AcrAB-mediated multidrug-resistant phenotype is maintained following restoration of wild-type activities by efflux pump genes and their regulators. International Journal of Antimicrobial Agents, 2009, 34, 602-604.	2.5	27
177	Emerging Infectious Disease Implications of Invasive Mammalian Species: The Greater White-Toothed Shrew (Crocidura russula) Is Associated With a Novel Serovar of Pathogenic Leptospira in Ireland. PLoS Neglected Tropical Diseases, 2016, 10, e0005174.	3.0	27
178	Biocide susceptibility testing of bacteria: Development of a broth microdilution method. Veterinary Microbiology, 2020, 248, 108791.	1.9	27
179	Integronlike structures in Campylobacter spp. of human and animal origin. Emerging Infectious Diseases, 2000, 6, 50-5.	4.3	27
180	Molecular epidemiology of cystic fibrosis-linked Burkholderia cepacia complex isolates from three national referral centres in Ireland. Journal of Applied Microbiology, 2002, 92, 992-1004.	3.1	26

#	Article	IF	CITATIONS
181	Molecular characterization of class 1 integrons from Irish thermophilic Campylobacter spp Journal of Antimicrobial Chemotherapy, 2004, 53, 952-957.	3.0	26
182	In-line milk filter analysis: Escherichia coli O157 surveillance of milk production holdings. International Journal of Hygiene and Environmental Health, 2005, 208, 407-413.	4.3	26
183	Surveillance of Dairy Production Holdings Supplying Raw Milk to the Farmhouse Cheese Sector for Escherichia coli O157, O26 and O111. Zoonoses and Public Health, 2007, 54, 358-365.	2.2	26
184	Comparison of disc diffusion and epsilometer (E-test) testing techniques to determine antimicrobial susceptibiliy of Campylobacter isolates of food and human clinical origin. Journal of Microbiological Methods, 2009, 79, 238-241.	1.6	26
185	Effect of crust freezing applied alone and in combination with ultraviolet light on the survival of Campylobacter on raw chicken. Food Microbiology, 2012, 32, 147-151.	4.2	26
186	Evidence of Metabolic Switching and Implications for Food Safety from the Phenome(s) of Salmonella enterica Serovar Typhimurium DT104 Cultured at Selected Points across the Pork Production Food Chain. Applied and Environmental Microbiology, 2013, 79, 5437-5449.	3.1	26
187	Development of a Custom-Designed, Pan Genomic DNA Microarray to Characterize Strain-Level Diversity among Cronobacter spp Frontiers in Pediatrics, 2015, 3, 36.	1.9	26
188	Serovar diversity and antimicrobial resistance of non-typhoidal Salmonella enterica recovered from retail chicken carcasses for sale inÂdifferent regions of China. Food Control, 2017, 81, 46-54.	5.5	26
189	Sentinel Listeriosis Surveillance in Selected Hospitals, China, 2013–2017. Emerging Infectious Diseases, 2019, 25, 2274-2277.	4.3	26
190	Epidemiological Study on Prevalence, Serovar Diversity, Multidrug Resistance, and CTX-M-Type Extended-Spectrum β-Lactamases of <i>Salmonella</i> spp. from Patients with Diarrhea, Food of Animal Origin, and Pets in Several Provinces of China. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	26
191	<i>Escherichia coli</i> serogroup O2 and O28ac O-antigen gene cluster sequences and detection of pathogenic <i>E. coli</i> O2 and O28ac by PCR. Canadian Journal of Microbiology, 2010, 56, 308-316.	1.7	25
192	Development and evaluation of DNA and RNA real-time assays for food analysis using the hilA gene of Salmonella enterica subspecies enterica. Food Microbiology, 2011, 28, 447-456.	4.2	25
193	Verocytotoxigenic Escherichia coli O157 in beef and sheep abattoirs in Ireland and characterisation of isolates by Pulsed-Field Gel Electrophoresis and Multi-Locus Variable Number of Tandem Repeat Analysis. International Journal of Food Microbiology, 2011, 144, 519-527.	4.7	25
194	Processing environment monitoring in low moisture food production facilities: Are we looking for the right microorganisms?. International Journal of Food Microbiology, 2021, 356, 109351.	4.7	25
195	Exposure of Chlorpromazine to 266 nm Laser Beam Generates New Species with Antibacterial Properties: Contributions to Development of a New Process for Drug Discovery. PLoS ONE, 2013, 8, e55767.	2.5	25
196	Relative contribution of target gene mutation and efflux to varying quinolone resistance in Irish Campylobacter isolates. FEMS Microbiology Letters, 2005, 253, 39-46.	1.8	24
197	Serotypes and virulence profiles of atypical enteropathogenic Escherichia coli (EPEC) isolated from bovine farms and abattoirs. Journal of Applied Microbiology, 2013, 114, 595-603.	3.1	24
198	Genome Sequence of Cronobacter sakazakiiSP291, a Persistent Thermotolerant Isolate Derived from a Factory Producing Powdered Infant Formula. Genome Announcements, 2013, 1, e0008213.	0.8	24

#	Article	IF	CITATIONS
199	Characterization of the genetic environment of blaESBL genes, integrons and toxin-antitoxin systems identified on large transferrable plasmids in multi-drug resistant Escherichia coli. Frontiers in Microbiology, 2015, 5, 716.	3.5	24
200	Prevalence and Molecular Characteristics of Extended-Spectrum β-Lactamase Genes in Escherichia coli Isolated from Diarrheic Patients in China. Frontiers in Microbiology, 2017, 8, 144.	3.5	24
201	Complete Genomic Analysis of a Salmonella enterica Serovar Typhimurium Isolate Cultured From Ready-to-Eat Pork in China Carrying One Large Plasmid Containing mcr-1. Frontiers in Microbiology, 2018, 9, 616.	3.5	24
202	Characterization of acmeABCOperon in a Quinolone- ResistantCampylobacter coliIsolate of Irish Origin. Microbial Drug Resistance, 2005, 11, 303-308.	2.0	23
203	Evaluation of an Automated Repetitive Sequence–Based PCR System for Subtyping Enterobacter sakazakii. Journal of Food Protection, 2008, 71, 1372-1378.	1.7	23
204	Molecular characterization of group A rotaviruses detected in children with gastroenteritis in Ireland in 2006–2009. Epidemiology and Infection, 2012, 140, 247-259.	2.1	23
205	Proteomic and phenotypic analysis of triclosan tolerant verocytotoxigenic Escherichia coli O157:H19. Journal of Proteomics, 2013, 80, 78-90.	2.4	23
206	Genomic characterization of an extensively-drug resistance Salmonella enterica serotype Indiana strain harboring blaNDM-1 gene isolated from a chicken carcass in China. Microbiological Research, 2017, 204, 48-54.	5.3	23
207	Antimicrobial resistance profiling and DNA Amplification Fingerprinting (DAF) of thermophilic Campylobacter spp. in human, poultry and porcine samples from the Cork region of Ireland. Journal of Applied Microbiology, 2000, 89, 727-734.	3.1	22
208	Comparison of phenotypic and genotypic characteristics of Salmonella bredeney associated with a poultry-related outbreak of gastroenteritis in Northern Ireland. Journal of Infection, 2003, 47, 33-39.	3.3	22
209	THERMAL RESISTANCE OF ANTIBIOTIC-RESISTANT AND ANTIBIOTIC-SENSITIVE SALMONELLA SPP. ON CHICKEN MEAT. Journal of Food Safety, 2005, 25, 288-302.	2.3	22
210	Characterization of a 2.6 kbp variable region within a class 1 integron found in an Acinetobacter baumannii strain isolated from a horse. Journal of Antimicrobial Chemotherapy, 2005, 55, 367-370.	3.0	22
211	Risk factors for the introduction and within-herd transmission of Mycobacterium avium subspecies paratuberculosis (MAP) infection on 59 Irish dairy herds. Irish Veterinary Journal, 2008, 61, 464-7.	2.1	22
212	A Real-Time Multiplex SYBR Green I Polymerase Chain Reaction Assay for Rapid Screening of <i>Salmonella</i> Serotypes Prevalent in the European Union. Foodborne Pathogens and Disease, 2011, 8, 769-780.	1.8	22
213	Strategies for the Identification and Tracking of Cronobacter Species: An Opportunistic Pathogen of Concern to Neonatal Health. Frontiers in Pediatrics, 2015, 3, 38.	1.9	22
214	Control of Listeria species food safety at a poultry food production facility. Food Microbiology, 2015, 51, 81-86.	4.2	22
215	Molecular characterization ofblaESBL-producingEscherichia colicultured from pig farms in Ireland. Journal of Antimicrobial Chemotherapy, 2016, 71, 3062-3065.	3.0	22
216	Genomic characterization of malonate positive Cronobacter sakazakii serotype O:2, sequence type 64 strains, isolated from clinical, food, and environment samples. Gut Pathogens, 2018, 10, 11.	3.4	22

#	Article	IF	CITATIONS
217	Multi-drug resistant Escherichia coli in diarrhoeagenic foals: Pulsotyping, phylotyping, serotyping, antibiotic resistance and virulence profiling. Veterinary Microbiology, 2018, 223, 144-152.	1.9	22
218	<i>Salmonella</i> harbouring the <i>mcr-1</i> gene isolated from food in China between 2012 and 2016. Journal of Antimicrobial Chemotherapy, 2019, 74, 826-828.	3.0	22
219	Silver Nanoparticles Induce a Triclosan-Like Antibacterial Action Mechanism in Multi-Drug Resistant Klebsiella pneumoniae. Frontiers in Microbiology, 2021, 12, 638640.	3.5	22
220	Natural co-occurrence of multi-mycotoxins in unprocessed wheat grains from China. Food Control, 2021, 130, 108321.	5.5	22
221	Genetic response of Salmonella enterica serotype Enteritidis to thioridazine rendering the organism resistant to the agent. International Journal of Antimicrobial Agents, 2012, 39, 16-21.	2.5	21
222	Characteristics of Aerococcus viridans isolated from bovine subclinical mastitis and its effect on milk SCC, yield, and composition. Tropical Animal Health and Production, 2017, 49, 843-849.	1.4	21
223	Identification of a novel hybrid plasmid coproducing MCR-1 and MCR-3 variant from an Escherichia coli strain. Journal of Antimicrobial Chemotherapy, 2019, 74, 1517-1520.	3.0	21
224	Genome-wide survey of efflux pump-coding genes associated with Cronobacter survival, osmotic adaptation, and persistence. Current Opinion in Food Science, 2019, 30, 32-42.	8.0	21
225	Inhibition of verocytotoxigenic Escherichia coli by antimicrobial peptides caseicin A and B and the factors affecting their antimicrobial activities. International Journal of Food Microbiology, 2012, 153, 260-268.	4.7	20
226	Microbiological study of biofilm formation in isolates of Salmonella enterica Typhimurium DT104 and DT104b cultured from the modern pork chain. International Journal of Food Microbiology, 2013, 161, 36-43.	4.7	20
227	Phenotypic Characterization of Salmonella Isolated from Food Production Environments Associated with Low–Water Activity Foods. Journal of Food Protection, 2013, 76, 1488-1499.	1.7	20
228	Molecular Analysis of OXA-48-Carrying Conjugative IncL/M-Like Plasmids in Clinical Isolates of <i>Klebsiella pneumoniae</i> in Ireland. Microbial Drug Resistance, 2014, 20, 270-274.	2.0	20
229	Comparative analysis of Salmonella susceptibility and tolerance to the biocide chlorhexidine identifies a complex cellular defense network. Frontiers in Microbiology, 2014, 5, 373.	3.5	20
230	Molecular and Phenotypic Characterization of Aerococcus viridans Associated with Subclinical Bovine Mastitis. PLoS ONE, 2015, 10, e0125001.	2.5	20
231	Exposure to Sub-inhibitory Concentrations of the Chemosensitizer 1-(1-Naphthylmethyl)-Piperazine Creates Membrane Destabilization in Multi-Drug Resistant Klebsiella pneumoniae. Frontiers in Microbiology, 2019, 10, 92.	3.5	20
232	Transcriptomic Analysis of Triclosan-Susceptible and -Tolerant <i>Escherichia coli</i> O157:H19 in Response to Triclosan Exposure. Microbial Drug Resistance, 2014, 20, 91-103.	2.0	19
233	Exposure of Salmonella enterica Serovar Typhimurium to Three Humectants Used in the Food Industry Induces Different Osmoadaptation Systems. Applied and Environmental Microbiology, 2015, 81, 6800-6811.	3.1	19
234	Complete Genome Sequence of Clostridium estertheticum DSM 8809, a Microbe Identified in Spoiled Vacuum Packed Beef. Frontiers in Microbiology, 2016, 7, 1764.	3.5	19

#	Article	IF	CITATIONS
235	Genomic Evolution of SARS-CoV-2 Virus in Immunocompromised Patient, Ireland. Emerging Infectious Diseases, 2021, 27, 2499-2501.	4.3	19
236	Direct Modification of Bioactive Phenothiazines by Exposure to Laser Radiation. Recent Patents on Anti-infective Drug Discovery, 2011, 6, 147-157.	0.8	19
237	Salmonella enterica Serotype Bredeney: Antimicrobial Susceptibility and Molecular Diversity of Isolates from Ireland and Northern Ireland. Applied and Environmental Microbiology, 2002, 68, 181-186.	3.1	18
238	Evidence for a Chromosomally Located Third Integron in Salmonella enterica Serovar Typhimurium DT104b. Antimicrobial Agents and Chemotherapy, 2004, 48, 1350-1352.	3.2	18
239	Molecular characterization of Irish E. coli O157:H7 isolates of human, bovine, ovine and porcine origin. Journal of Applied Microbiology, 2009, 107, 1340-1349.	3.1	18
240	Genetic diversity among <i>Campylobacter jejuni</i> isolates from pets in Ireland. Veterinary Record, 2010, 166, 102-106.	0.3	18
241	Heat Adaptation and Survival of <i>Cronobacter</i> spp. (Formerly <i>Enterobacter sakazakii</i> ). Foodborne Pathogens and Disease, 2011, 8, 975-981.	1.8	18
242	A novel disrupted <i>mcr-1</i> gene and a lysogenized phage P1-like sequence detected from a large conjugative plasmid, cultured from a human atypical enteropathogenic <i>Escherichia coli</i> (aEPEC) recovered in China. Journal of Antimicrobial Chemotherapy, 2017, 72, dkw564.	3.0	18
243	Characterizing the Multidrug Resistance of non-O157 Shiga Toxin-ProducingEscherichia colilsolates from Cattle Farms and Abattoirs. Microbial Drug Resistance, 2017, 23, 781-790.	2.0	18
244	Susceptibility (re)-testing of a large collection of Listeria monocytogenes from foods in China from 2012 to 2015 and WGS characterization of resistant isolates. Journal of Antimicrobial Chemotherapy, 2019, 74, 1786-1794.	3.0	18
245	Enriching antimicrobial peptides from milk hydrolysates using pectin/alginate food-gels. Food Chemistry, 2021, 352, 129220.	8.2	18
246	Investigation of a methicillin-resistant Staphylococcus aureus (MRSA) outbreak in an Irish hospital: triplex PCR and DNA amplification fingerprinting. Journal of Hospital Infection, 1997, 36, 37-47.	2.9	17
247	Antibiotic resistance of retail food and human Campylobacter isolates on the island of Ireland from 2001–2002. Epidemiology and Infection, 2006, 134, 1282-1291.	2.1	17
248	Detection and characterisation of bovine rotavirus in Ireland from 2006–2008. Irish Veterinary Journal, 2014, 67, 13.	2.1	17
249	Changing patterns of rotavirus strains circulating in Ireland: Re-emergence of G2P[4] and identification of novel genotypes in Ireland. Journal of Medical Virology, 2015, 87, 764-773.	5.0	17
250	Potential for transfer of Escherichia coli O157:H7, Listeria monocytogenes and Salmonella Senftenberg from contaminated food waste derived compost and anaerobic digestate liquid to lettuce plants. Food Microbiology, 2016, 59, 7-13.	4.2	17
251	Analysis of the Molecular Diversity Among Cronobacter Species Isolated From Filth Flies Using Targeted PCR, Pan Genomic DNA Microarray, and Whole Genome Sequencing Analyses. Frontiers in Microbiology, 2020, 11, 561204.	3.5	17
252	Application of Whole Genome Sequencing to Aid in Deciphering the Persistence Potential of Listeria monocytogenes in Food Production Environments. Microorganisms, 2021, 9, 1856.	3.6	17

#	Article	IF	CITATIONS
253	Characterization of Staphylococcus aureus Isolated from Dairy Animals in Ireland. Zoonoses and Public Health, 2010, 57, 249-257.	2.2	16
254	Patterns of antimicrobial resistance in pathogenic Escherichia coli isolates from cases of calf enteritis during the spring-calving season. Veterinary Microbiology, 2014, 170, 73-80.	1.9	16
255	Insights into antimicrobial resistance among long distance migratory East Canadian High Arctic light-bellied Brent geese (Branta bernicla hrota). Irish Veterinary Journal, 2015, 69, 13.	2.1	16
256	Increased Virulence of Bloodstream Over Peripheral Isolates of P. aeruginosa Identified Through Post-transcriptional Regulation of Virulence Factors. Frontiers in Cellular and Infection Microbiology, 2018, 8, 357.	3.9	16
257	The prevalence of Clostridioides difficile on farms, in abattoirs and in retail foods in Ireland. Food Microbiology, 2021, 98, 103781.	4.2	16
258	Multiple-locus variable number of tandem repeat analysis (MLVA) of Irish verocytotoxigenic Escherichia coliO157 from feedlot cattle: uncovering strain dissemination routes. BMC Veterinary Research, 2008, 4, 2.	1.9	15
259	Antimicrobial Resistance Profiles and Mechanisms of Resistance in <i>Campylobacter jejuni</i> Isolates from Pets. Foodborne Pathogens and Disease, 2009, 6, 705-710.	1.8	15
260	A study of the molecular basis of quinolone and macrolide resistance in a selection of Campylobacter isolates from intensive poultry flocks. Food Control, 2013, 30, 222-226.	5.5	15
261	A Comparative Study of the Susceptibility of Listeria Species to Sanitizer Treatments When Grown under Planktonic and Biofilm Conditions. Journal of Food Protection, 2018, 81, 1481-1490.	1.7	15
262	Effects of metal and metalloid pollutants on the microbiota composition of feces obtained from twelve commercial pig farms across China. Science of the Total Environment, 2019, 647, 577-586.	8.0	15
263	Yersinia hibernica sp. nov., isolated from pig-production environments. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 2023-2027.	1.7	15
264	Application of Restriction Fragment Length Polymorphism Analysis of VP7-Encoding Genes: Fine Comparison of Irish and Global Rotavirus Isolates. Journal of Clinical Microbiology, 2002, 40, 524-531.	3.9	14
265	A novel quantitative reverse-transcription PCR (qRT-PCR) for the enumeration of total bacteria, using meat micro-flora as a model. Journal of Microbiological Methods, 2009, 77, 1-7.	1.6	14
266	Assessment of horizontal gene transfer in Lactic acid bacteria – A comparison of mating techniques with a view to optimising conjugation conditions. Journal of Microbiological Methods, 2009, 77, 23-28.	1.6	14
267	Mechanisms of Antibiotic Resistance in Salmonella: Efflux Pumps, Genetics, Quorum Sensing and Biofilm Formation. Letters in Drug Design and Discovery, 2011, 8, 114-123.	0.7	14
268	Flow cytometry to assess biochemical pathways in heat-stressed Cronobacter spp. (formerly) Tj ETQq0 0 0 rgBT ,	Oyerlock	10 <sub>14</sub> 50 142
269	Application of PCR for rapid detection and serotyping of Salmonella spp. from porcine carcass swabs following enrichment in semi-solid agar. Food Research International, 2012, 45, 993-999.	6.2	14
270	Complete genomic sequence analyses of the first group A giraffe rotavirus reveals close evolutionary relationship with rotaviruses infecting other members of the Artiodactyla. Veterinary Microbiology, 2014, 170, 151-156.	1.9	14

#	Article	IF	CITATIONS
271	<i>Salmonella</i> in breeding pigs: Shedding pattern, transmission of infection and the role of environmental contamination in Irish commercial farrowâ€toâ€finish herds. Zoonoses and Public Health, 2018, 65, e196-e206.	2.2	14
272	Modulation of antimicrobial resistance in clinical isolates of Enterobacter aerogenes: A strategy combining antibiotics and chemosensitisers. Journal of Global Antimicrobial Resistance, 2019, 16, 187-198.	2.2	14
273	Investigation of the Causes of Shigatoxigenic Escherichia coli PCR Positive and Culture Negative Samples. Microorganisms, 2020, 8, 587.	3.6	14
274	Emergence of a <i>Salmonella enterica</i> serovar Typhimurium ST34 isolate, CFSA629, carrying a novel <i>mcr-1.19</i> variant cultured from egg in China. Journal of Antimicrobial Chemotherapy, 2021, 76, 1776-1785.	3.0	14
275	Colonisation dynamics of Listeria monocytogenes strains isolated from food production environments. Scientific Reports, 2021, 11, 12195.	3.3	14
276	Insights into the mechanisms of Cronobacter sakazakii virulence. Microbial Pathogenesis, 2022, 169, 105643.	2.9	14
277	Nucleotide and deduced amino acid sequences of Rhizobium meliloti 102F34 lacZ gene: comparison with prokaryotic β-galactosidases and human β-glucuronidase. Gene, 1994, 141, 91-96.	2.2	13
278	Prevalence of <i>Campylobacter</i> spp. in a subset of intensive poultry flocks in Ireland. Letters in Applied Microbiology, 2009, 49, 305-310.	2.2	13
279	First report: Yersinia enterocolitica recovered from canine tonsils. Veterinary Microbiology, 2010, 146, 336-339.	1.9	13
280	Roles of Diet and the Acid Tolerance Response in Survival of Common Salmonella Serotypes in Feces of Finishing Pigs. Applied and Environmental Microbiology, 2012, 78, 110-119.	3.1	13
281	THE BIODIVERSITY OF THERMODURIC BACTERIA ISOLATED FROM WHEY. Journal of Food Safety, 2012, 32, 255-261.	2.3	13
282	Norovirus contamination in retail oysters from Beijing and Qingdao, China. Food Control, 2018, 86, 415-419.	5.5	13
283	Analysis of the Oxidative Stress Regulon Identifies <i>soxS</i> as a Genetic Target for Resistance Reversal in Multidrug-Resistant Klebsiella pneumoniae. MBio, 2021, 12, e0086721.	4.1	13
284	Evaluation of a New One-Step Enrichment in Conjunction with a Chromogenic Medium for the Detection of Cronobacter spp. (Enterobacter sakazakii)in Powdered Infant Formula. Journal of Food Protection, 2009, 72, 1472-1475.	1.7	13
285	Genomic Epidemiology of ST34 Monophasic Salmonella enterica Serovar Typhimurium from Clinical Patients from 2008 to 2017 in Henan, China. Engineering, 2022, 15, 34-44.	6.7	13
286	Use of Aprotinin Therapy in a Patient with Factor V Leiden. Anesthesia and Analgesia, 1997, 84, 694-698.	2.2	12
287	Prevalence and Characterization ofEscherichia coliO26 and O111 in Retail Minced Beef in Ireland. Foodborne Pathogens and Disease, 2005, 2, 357-360.	1.8	12
288	Molecular characterisation of a bovine-like rotavirus detected from a giraffe. BMC Veterinary Research, 2008, 4, 46.	1.9	12

#	Article	IF	CITATIONS
289	Complete nucleotide sequence of pVQS1 containing a quinolone resistance determinant from Salmonella enterica serovar Virchow associated with foreign travel. Journal of Antimicrobial Chemotherapy, 2012, 67, 1861-1864.	3.0	12
290	Detection and characterisation of novel bocavirus (genus <i>Bocaparvovirus</i> ) and gastroenteritis viruses from asymptomatic pigs in Ireland. Infection Ecology and Epidemiology, 2015, 5, 27270.	0.8	12
291	Flow Cytometric and 16S Sequencing Methodologies for Monitoring the Physiological Status of the Microbiome in Powdered Infant Formula Production. Frontiers in Microbiology, 2016, 7, 968.	3.5	12
292	An Investigation of the Innate Immune Response in Bovine Mammary Epithelial Cells Challenged by Prototheca zopfii. Mycopathologia, 2016, 181, 823-832.	3.1	12
293	Genomic insights into the pathogenicity and environmental adaptability of <i>Enterococcus hirae</i> R17 isolated from pork offered for retail sale. MicrobiologyOpen, 2017, 6, e00514.	3.0	12
294	Salmonella tel-el-kebir and terrapins. Journal of Infection, 1999, 38, 182-184.	3.3	11
295	Validation of DNA and RNA real-time assays for food analysis using the hilA gene of Salmonella enterica serovars. Journal of Microbiological Methods, 2011, 84, 19-26.	1.6	11
296	Complete nucleotide sequence of a conjugative IncF plasmid from an <i>Escherichia coli</i> isolate of equine origin containing <i>bla</i> <sub>CMY-2</sub> within a novel genetic context. FEMS Microbiology Letters, 2014, 352, 123-127.	1.8	11
297	The effect of Quaternary Ammonium Compounds on the attachment of wild and adapted Pseudomonas putida strains to different contact materials used in the food sector. Food Control, 2014, 42, 277-283.	5.5	11
298	Cloning and characterization of a novel β-galactosidase-coding gene from rhizobium meliloti. Gene, 1988, 71, 57-64.	2.2	10
299	Antimicrobial Resistance in Isolates of Salmonella spp. from Pigs and the Characterization of an S. Infantis Gene Cassette. Foodborne Pathogens and Disease, 2005, 2, 274-281.	1.8	10
300	Investigation of a global collection of nontyphoidalSalmonellaof various serotypes cultured between 1953 and 2004 for the presence of class 1 integrons. FEMS Microbiology Letters, 2007, 266, 170-176.	1.8	10
301	Genotypic characterisation and cluster analysis of Campylobacter jejuniisolates from domestic pets, human clinical cases and retail food. Irish Veterinary Journal, 2011, 64, 6.	2.1	10
302	Effect of surface properties of different food contact materials on the efficiency of quaternary ammonium compounds residue recovery and persistence. International Journal of Food Science and Technology, 2013, 48, 1791-1797.	2.7	10
303	A longitudinal study of Staphylococcus aureus colonization in pigs in Ireland. Veterinary Microbiology, 2014, 174, 504-513.	1.9	10
304	A Novel Tn3-Like Composite Transposon HarboringblaVIM-1inKlebsiella pneumoniaespp.pneumoniaelsolated from River Water. Microbial Drug Resistance, 2015, 21, 43-49.	2.0	10
305	The role of selenium in insulin-like growth factor I receptor (IGF-IR) expression and regulation of apoptosis in mouse osteoblasts. Chemosphere, 2016, 144, 2158-2164.	8.2	10
306	Interaction of matrix metalloproteinase-9 and Zpx in <i>Cronobacter turicensis</i> LMG 23827 <sup>T</sup> mediated infections in the zebrafish model. Cellular Microbiology, 2018, 20, e12888.	2.1	10

#	Article	IF	CITATIONS
307	A quantitative real time PCR assay to detect and enumerate Escherichia coli O157 and O26 serogroups in sheep recto-anal swabs. Journal of Microbiological Methods, 2019, 165, 105703.	1.6	10
308	Molecular characterisation of multi-drug resistant Escherichia coli of bovine origin. Veterinary Microbiology, 2020, 242, 108566.	1.9	10
309	Detection of Hepatitis E Virus in the Pig Livers and Retail Pork Samples Collected in Selected Cities in China. Foodborne Pathogens and Disease, 2021, 18, 97-103.	1.8	10
310	Comparative Genomic Analysis of the Foodborne Pathogen Burkholderia gladioli pv. cocovenenans Harboring a Bongkrekic Acid Biosynthesis Gene Cluster. Frontiers in Microbiology, 2021, 12, 628538.	3.5	10
311	Yersinia canariae sp. nov., isolated from a human yersiniosis case. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 2382-2387.	1.7	10
312	Investigation of the persistence and transmission of MRSA CC 5 in pigs following intra-nasal inoculation. Veterinary Microbiology, 2013, 162, 771-778.	1.9	9
313	Phenotypic changes contributing to <i>Enterobacter gergoviae</i> biocide resistance. Letters in Applied Microbiology, 2015, 61, 121-129.	2.2	9
314	Silent antibiotic resistance genes: A threat to antimicrobial therapy. International Journal of Infectious Diseases, 2019, 79, 20.	3.3	9
315	Ram locus is a key regulator to trigger multidrug resistance in Enterobacter aerogenes. Journal of Medical Microbiology, 2018, 67, 148-159.	1.8	9
316	First isolation and report of clusters of Clostridium difficile PCR 027 cases in Ireland. , 2007, 12, E070426.3.		9
317	Prevalence and Whole-Genome Sequence-Based Analysis of Shiga Toxin-Producing Escherichia coli Isolates from the Recto-Anal Junction of Slaughter-Age Irish Sheep. Applied and Environmental Microbiology, 2021, 87, e0138421.	3.1	9
318	Inactivation and Recovery of High Quality RNA From Positive SARS-CoV-2 Rapid Antigen Tests Suitable for Whole Virus Genome Sequencing. Frontiers in Public Health, 2022, 10, 863862.	2.7	9
319	Cronobacter Species. , 2019, , 389-414.		8
320	Nosocomial cross-infection of hypervirulent Listeria monocytogenes sequence type 87 in China. Annals of Translational Medicine, 2020, 8, 603-603.	1.7	8
321	An Overview of Shiga-Toxin Producing <i>Escherichia coli</i> Carriage and Prevalence in the Ovine Meat Production Chain. Foodborne Pathogens and Disease, 2021, 18, 147-168.	1.8	8
322	Alterations in the Transcriptional Landscape Allow Differential Desiccation Tolerance in Clinical Cronobacter sakazakii. Applied and Environmental Microbiology, 2021, 87, e0083021.	3.1	8
323	Comparison Between Full-Length 16S rRNA Metabarcoding and Whole Metagenome Sequencing Suggests the Use of Either Is Suitable for Large-Scale Microbiome Studies. Foodborne Pathogens and Disease, 2022, 19, 495-504.	1.8	8
324	Effect of divalent metal cations on <u>Rhizobium meliloti</u> B-galactosidase. Biochemical Society Transactions, 1991, 19, 19S-19S.	3.4	7

#	Article	IF	CITATIONS
325	Purification and some characteristics of a recombinant dimeric rhizobium meliloti β-galactosidase expressed in escherichia coli. Enzyme and Microbial Technology, 2001, 28, 682-688.	3.2	7
326	Development and assessment of a rapid method to detect Escherichia coli O26, O111 and O157 in retail minced beef. International Journal of Hygiene and Environmental Health, 2007, 210, 155-161.	4.3	7
327	Potential of a commercially-available water acidification product for reducingCampylobacterin broilers prior to slaughter. British Poultry Science, 2013, 54, 130320024130007.	1.7	7
328	Complete Genome Sequence of Leptospira alstonii Serovar Room22 Strain GWTS #1. Genome Announcements, 2016, 4, .	0.8	7
329	Draft Genome Sequences of 15 Isolates of Listeria monocytogenes Serotype 1/2a, Subgroup ST204. Genome Announcements, 2016, 4, .	0.8	7
330	Characterisation of Early Positive mcr-1 Resistance Gene and Plasmidome in Escherichia coli Pathogenic Strains Associated with Variable Phylogroups under Colistin Selection. Antibiotics, 2021, 10, 1041.	3.7	7
331	Isolation and Identification of Enterobacter sakazakii. , 0, , 27-59.		7
332	Genomic fingerprinting Acinetobacter baumannii: amplification of multiple inter-repetitive extragenic palindromic sequences. Journal of Hospital Infection, 1995, 31, 33-40.	2.9	6
333	Towards "Molecular Esperanto―or the Tower of Babel? (The Need for Harmonization of Techniques) Tj ETQq1	1 0.7843 3.9	14 rgBT /O
334	Comparison of Salmonella species recovered from Irish liquid milk production holdings with temporal clinical veterinary isolates. International Journal of Hygiene and Environmental Health, 2008, 211, 283-291.	4.3	6
335	The Safety of Raw Liquid Milk. , 0, , 139-167.		6
336	Clostridium difficileInfection in the Republic of Ireland: Results of a 1-Month National Surveillance and Ribotyping Project, March 2009. Infection Control and Hospital Epidemiology, 2010, 31, 1085-1087.	1.8	6
337	An investigation of the subtype diversity of clinical isolates of Irish Clostridium difficile ribotypes 027 and 078 by repetitive-extragenic palindromic PCR. Journal of Medical Microbiology, 2011, 60, 1080-1087.	1.8	6
338	Evaluation of Carvacrol for the Control of <i>Escherichia coli</i> O157 on Cattle Hide and Carcass Cuts. Foodborne Pathogens and Disease, 2012, 9, 1049-1052.	1.8	6
339	Use of a Pan–Genomic DNA Microarray in Determination of the Phylogenetic Relatedness among Cronobacter spp. and Its Use as a Data Mining Tool to Understand Cronobacter Biology. Microarrays (Basel, Switzerland), 2017, 6, 6.	1.4	6
340	Controlling Blown Pack Spoilage Using Anti-Microbial Packaging. Foods, 2017, 6, 67.	4.3	6
341	Reversing Antimicrobial Resistance in Multidrug-Resistant <i>Klebsiella pneumoniae</i> of Clinical Origin Using 1-(1-Naphthylmethyl)-Piperazine. Microbial Drug Resistance, 2018, 24, 1497-1506.	2.0	6
342	Atypical Salmonella enterica Serovars in Murine and Human Macrophage Infection Models. Infection and Immunity, 2020, 88, .	2.2	6

#	Article	IF	CITATIONS
343	Characterization of Cronobacter sakazakii Strains Originating from Plant-Origin Foods Using Comparative Genomic Analyses and Zebrafish Infectivity Studies. Microorganisms, 2022, 10, 1396.	3.6	6
344	Shotgun metagenomic sequencing of bulk tank milk filters reveals the role of Moraxellaceae and Enterobacteriaceae as carriers of antimicrobial resistance genes. Food Research International, 2022, 158, 111579.	6.2	6
345	Integron Analysis and Genetic Mapping of Antimicrobial Resistance Genes in <i>Salmonella enterica</i> serotype Typhimurium. , 2004, 268, 015-032.		5
346	Application of quantitative reverseâ€ŧranscription PCR (qRTâ€PCR) for the determination of the total viable count (TVC) on meat samples. Journal of Applied Microbiology, 2010, 109, 91-98.	3.1	5
347	Genome Sequence of an Enterobacter helveticus Strain, 1159/04 (LMG 23733), Isolated from Fruit Powder. Genome Announcements, 2013, 1, .	0.8	5
348	Targets to increase food production: One Health implications. Infection Ecology and Epidemiology, 2015, 5, 27708.	0.8	5
349	Contamination and characterization of multiple pathogens in powdered formula at retail collected between 2014 and 2015 in China. Food Control, 2018, 87, 40-45.	5.5	5
350	Complete Genome and Plasmid Sequences of Seven Isolates of Salmonella enterica subsp. enterica Harboring the mcr-1 Gene Obtained from Food in China. Microbiology Resource Announcements, 2019, 8, .	0.6	5
351	Comparison of two methods for cell count determination in the course of biocide susceptibility testing. Veterinary Microbiology, 2020, 251, 108831.	1.9	5
352	Investigation of tigecycline resistant Escherichia coli from raw meat reveals potential transmission among food-producing animals. Food Control, 2021, 121, 107633.	5.5	5
353	Pulsed-Field Gel Electrophoresis (PFGE) for Pathogenic Cronobacter Species. Methods in Molecular Biology, 2015, 1301, 55-69.	0.9	5
354	Essential veterinary education on the development of antimicrobial and anti parasitic resistance: consequences for animal health and food safety and the need for vigilance. OIE Revue Scientifique Et Technique, 2009, 28, 575-582.	1.2	5
355	Survival Characteristics of <i>Cronobacter</i> spp. in Model Bovine Gut and in the Environment. Foodborne Pathogens and Disease, 2010, 7, 671-675.	1.8	4
356	Genome Sequence of Enterobacter turicensis Strain 610/05 (LMG 23731), Isolated from Fruit Powder. Genome Announcements, 2013, 1, .	0.8	4
357	Comparison of Listeria monocytogenes Isolates across the Island of Ireland. Journal of Food Protection, 2014, 77, 1402-1406.	1.7	4
358	Epidemiological and genetic characterization of Clostridium butyricum cultured from neonatal cases of necrotizing enterocolitis in China. Infection Control and Hospital Epidemiology, 2020, 41, 900-907.	1.8	4
359	Characterization of Non-O157 Shiga Toxin-Producing Escherichia coli Cultured from Cattle Farms in Xinjiang Uygur Autonomous Region, China, During 2016–2017. Foodborne Pathogens and Disease, 2021, 18, 761-770.	1.8	4
360	Antimicrobial Resistance and Genomic Characterization of Two mcr-1-Harboring Foodborne Salmonella Isolates Recovered in China, 2016. Frontiers in Microbiology, 2021, 12, 636284.	3.5	4

#	Article	IF	CITATIONS
361	Detection of the heat-stable toxin coding gene (ST-gene) in enterotoxigenic Escherichia coli: development of a colour amplified PCR detection system. British Journal of Biomedical Science, 1995, 52, 317-20.	1.3	4
362	Complete genome sequences and genomic characterization of five plasmids harbored by environmentally persistent Cronobacter sakazakii strains ST83 H322 and ST64 GK1025B obtained from powdered infant formula manufacturing facilities. Gut Pathogens, 2022, 14, .	3.4	4
363	16S rRNA Based Profiling of Bacterial Communities Colonizing Bakery-Production Environments. Foodborne Pathogens and Disease, 2022, 19, 485-494.	1.8	4
364	Introductory note to the Cronobacter special issue. International Journal of Food Microbiology, 2009, 136, 151.	4.7	3
365	The effect of bovine diet on Salmonella survival in synthetic abomasal fluid. Journal of Applied Microbiology, 2010, 109, 2060-2068.	3.1	3
366	Current evidence for human yersiniosis in Ireland. European Journal of Clinical Microbiology and Infectious Diseases, 2012, 31, 2969-2981.	2.9	3
367	Further Characterization of Three Yersinia enterocolitica Strains with a Nalidixic Acid–Resistant Phenotype Isolated from Humans with Diarrhea. Foodborne Pathogens and Disease, 2013, 10, 744-746.	1.8	3
368	Genome Sequences of Two Enterobacter pulveris Strains, 601/05 T (=LMG 24057 T =DSM 19144 T ) and 1160/04 (=LMG 24058 =DSM 19146), Isolated from Fruit Powder. Genome Announcements, 2013, 1, .	0.8	3
369	Veterinary Drugs Residues: Antibacterials. , 2014, , 39-44.		3
370	A comparison of the efficiency of ELISA and selected primer sets to detect Norovirus isolates in southern Ireland over a four-year period (2002-2006): variation in detection rates and evidence for continuing predominance of NoV GII.4 genotype. Archives of Virology, 2014, 159, 1697-1705.	2.1	3
371	Complete Genome Sequence of Listeria monocytogenes Strain DPC6895, a Serotype 1/2b Isolate from Bovine Raw Milk. Genome Announcements, 2015, 3, .	0.8	3
372	Quaternary Ammonium Compounds (QACs) induced inactivation of Pseudomonas spp.: Effect of material surface. Food and Bioproducts Processing, 2016, 98, 71-78.	3.6	3
373	Effect of Exposure to Chlorhexidine Residues at "During Use―Concentrations on Antimicrobial Susceptibility Profile, Efflux, Conjugative Plasmid Transfer, and Metabolism of Escherichia coli. Antimicrobial Agents and Chemotherapy, 2020, 64, .	3.2	3
374	Sporulation and Biofilms as Survival Mechanisms of <i>Bacillus</i> Species in Low-Moisture Food Production Environments. Foodborne Pathogens and Disease, 2022, 19, 448-462.	1.8	3
375	Molecular-Based Identification and Typing of <1>Campylobacter jejuni and <1>C. coli. , 2004, 268, 033-048.		2
376	AN INVESTIGATION OF TRANSPORT, LAIRAGE AND HIDE CLEANLINESS ON CAMPYLOBACTER PREVALENCE IN FEEDLOT CATTLE AND DRESSED CARCASSES. Journal of Food Safety, 2004, 24, 37-52.	2.3	2
377	In VitroStudies on the Effect of pH and Volatile Fatty Acid Concentration, as Influenced by Diet, on the Survival of Inoculated Nonacid- and Acid-AdaptedSalmonellain Bovine Rumen Fluid and Feces. Foodborne Pathogens and Disease, 2011, 8, 609-614.	1.8	2
378	Tracing pathogens in red meat and game production chains and at the abattoir. , 2011, , 393-432.		2

#	Article	IF	CITATIONS
379	Polypharmacy Is an Independent Risk Factor for Oropharyngeal Isolation of Gramâ€Negative Bacilli in Older Persons. Journal of the American Geriatrics Society, 2012, 60, 182-183.	2.6	2
380	Molecular Characterization of Salmonella Serovars Anatum and Ealing Associated with Two Historical Outbreaks, Linked to Contaminated Powdered Infant Formula. Frontiers in Microbiology, 2016, 7, 1664.	3.5	2
381	Antimicrobial Resistance in Listeria Species. , 2018, , 237-259.		2
382	The Therapeutic Potential of the $\hat{a} {\in} \infty$ Yin-Yang $\hat{a} {\in} {f G}$ arden in Our Gut. , 2019, , .		2
383	Draft genome sequences of <i>Salmonella</i> Oslo isolated from seafood and its laboratory generated auxotrophic mutant. Journal of Genomics, 2020, 8, 7-10.	0.9	2
384	Epidemiology of Norovirus among outpatients presenting with acute diarrhea in Dalian, China. Biosafety and Health, 2020, 2, 60-63.	2.7	2
385	Molecular characterization of two novel NDM-1-producing atypical enteroaggregative Escherichia coli isolates from patients. Plasmid, 2021, 115, 102568.	1.4	2
386	Whole Genome Analysis of Three Multi-Drug Resistant Listeria innocua and Genomic Insights Into Their Relatedness With Resistant Listeria monocytogenes. Frontiers in Microbiology, 2021, 12, 694361.	3.5	2
387	Safety of Meat Products. Food Science Text Series, 2012, , 147-158.	0.3	2
388	Surveillance of Shiga-toxin producing Escherichia coli in Irish sheep. Access Microbiology, 2019, 1, .	0.5	2
389	Differences in antimicrobial susceptibility testing complicating management of IMP carbapenemase-producing Enterobacterales infection. Journal of Global Antimicrobial Resistance, 2021, 27, 284-288.	2.2	2
390	Does Silver in Different Forms Affect Bacterial Susceptibility and Resistance? A Mechanistic Perspective. ACS Applied Bio Materials, 2022, 5, 801-817.	4.6	2
391	Molecular Genetics of Factor V Leiden: Genetic Origins and Modern DNA-Based Detection Strategies. Seminars in Cardiothoracic and Vascular Anesthesia, 1997, 1, 333-341.	1.0	1
392	Phenotypic and Genotypic Anti-Microbial Resistance Profiles of Campylobacters from Untreated Feedlot Cattle and Their Environment. Zoonoses and Public Health, 2006, 53, 181-187.	1.4	1
393	SUBSPECIES CHARACTERIZATION OF UREASE-POSITIVE THERMOPHILIC CAMPYLOBACTER (UPTC) ISOLATED FROM SHELLFISH EMPLOYING MODIFIED FLAGELLIN (flaA) RESTRICTION FRAGMENT LENGTH POLYMORPHISM (RFLP) TYPING. Journal of Shellfish Research, 2006, 25, 625-629.	0.9	1
394	Shiga toxin-producing Escherichia coli isolated from human and pig origin induce different gene expression profiles in human Caco-2 epithelial cells. Livestock Science, 2010, 133, 189-191.	1.6	1
395	Pathogens in Milk   Enterobacter spp , 2011, , 72-80.		1

Cronobacter Species (formerly Enterobacter sakazakii). , 2013, , 251-258.

#	Article	IF	CITATIONS
397	Draft Genome Sequence of Escherichia coli 26R 793, a Plasmid-Free Recipient Strain Commonly Used in Conjugation Assays. Genome Announcements, 2016, 4, .	0.8	1
398	Occurrence of CTX-M-123-producing Salmonella Indiana in chicken carcasses: a new challenge for the poultry industry and food safety. Journal of Antimicrobial Chemotherapy, 2019, 74, 3637-3639.	3.0	1
399	The Role of Genomics in Food Quality and Safety Management: Possibilities and Limitations. , 2021, , 127-137.		1
400	Cronobacter species. , 2021, , 265-283.		1
401	Investigation of the Anti-Methicillin-Resistant Staphylococcus aureus Activity of (+)-Tanikolide- and (+)-Malyngolide-Based Analogues Prepared by Asymmetric Synthesis. International Journal of Molecular Sciences, 2021, 22, 6400.	4.1	1
402	Enterobacter Species. , 2022, , 469-481.		1
403	Transcriptomic Responses of Salmonella Species to Desiccation and Low-Moisture Environments: Extending Our Knowledge of How Bacteria Cope with Low-Moisture Stress. , 2014, , 49-66.		1
404	Pulsed-Field Gel Electrophoresis for Listeria monocytogenes. Methods in Molecular Biology, 2015, 1301, 43-53.	0.9	1
405	Comparative genomic insights into Yersinia hibernica – a commonly misidentified Yersinia enterocolitica-like organism. Microbial Genomics, 2020, 6, .	2.0	1
406	High-Throughput Characterization of Listeria monocytogenes Using the OmniLog Phenotypic Microarray. Methods in Molecular Biology, 2021, 2220, 107-113.	0.9	1
407	Molecular and Epidemiological Characterization of Infant Botulism in Beijing, China. Biomedical and Environmental Sciences, 2017, 30, 460-464.	0.2	1
408	A 16S rRNA Sequencing Study Describing the Environmental Microbiota of Two Powdered Infant Formula Built Facilities. Foodborne Pathogens and Disease, 2022, 19, 473-484.	1.8	1
409	Non-radioactive detection of the Staphylococcus aureus protein a gene (spa) using the polymerase chain reaction (PCR). Biochemical Education, 1994, 22, 52-55.	0.1	Ο
410	Applications of the polymerase chain reaction (PCR) in diagnosis. Irish Journal of Medical Science, 1995, 164, 116-121.	1.5	0
411	Royal Academy of Medicine in Ireland Section of Pathology. Irish Journal of Medical Science, 1997, 166, 267-269.	1.5	0
412	Motif-Dependent Polymerase Chain Reaction (PCR): DNA Fingerprinting Enterotoxigenic Escherichia coli. , 1998, 92, 115-126.		0
413	Cronobacter Species in Powdered Infant Formula. , 2014, , 307-317.		0
414	Pathogens in Milk: Enterobacter Species. , 2016, , .		0

24

#	Article	IF	CITATIONS
415	Molecular Characterization of CTX-M Producing Salmonella Isolates with Concurrent Resistance to Ciprofloxacin and Cefotaxime from Slaughtered Chicken Carcasses in Qingdao, China. Journal of Antimicrobial Agents, 2018, 04, .	0.2	Ο
416	Pathogens in Milk: Campylobacter spp , 2020, , 419-419.		0
417	<i>Call for Special Issue Papers:</i> Special Issue on Detecting and Tracking Zoonotic Bacteria Using Non-Culture-Based Methods. Foodborne Pathogens and Disease, 2021, 18, 297-297.	1.8	0
418	The Genomics Revolution: Agri-Food Research in the 21st Century. , 2021, , 2-18.		0
419	BM0701: Antibiotic Transport and Efflux: New Strategies to Combat Bacterial Resistance (ATENS). Letters in Drug Design and Discovery, 2011, 8, 101-101.	0.7	Ο
420	Preliminary Survey Regarding Yersiniosis in Ireland. Advances in Experimental Medicine and Biology, 2012, 954, 59-61.	1.6	0
421	Sampling, Transport, and Sample Preparation in Emergency Situations and Rapid Response. , 0, , 93-101.		0
422	Analysis of phenotypic traits which may impact long term survival of different Escherichia coli pathotypes. Access Microbiology, 2019, 1, .	0.5	0
423	Draft Genome Sequences of Three Novel Acinetobacter Isolates from an Irish Commercial Pig Farm. Microbiology Resource Announcements, 2019, 8, .	0.6	Ο
424	Whole-Genome Sequencing Confirms SARS-CoV-2 Transmission between Healthcare Workers and Patients. SSRN Electronic Journal, 0, , .	0.4	0
425	Detecting and Tracking Zoonotic Bacteria Using Nonculture-Based Methods. Foodborne Pathogens and Disease, 0, , .	1.8	Ο