

Herbert Schmidt

List of Publications by Year in descending order

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128
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81839

39
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79644

73
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134
all docs

134
docs citations

134
times ranked

4356
citing authors

#	ARTICLE	IF	CITATIONS
1	Pathogenicity Islands in Bacterial Pathogenesis. <i>Clinical Microbiology Reviews</i> , 2004, 17, 14-56.	5.7	603
2	EspP, a novel extracellular serine protease of enterohaemorrhagic <i>Escherichia coli</i> O157:H7 cleaves human coagulation factor V. <i>Molecular Microbiology</i> , 1997, 24, 767-778.	1.2	386
3	A New Shiga Toxin 2 Variant (Stx2f) from <i>Escherichia coli</i> Isolated from Pigeons. <i>Applied and Environmental Microbiology</i> , 2000, 66, 1205-1208.	1.4	284
4	Shiga toxin-encoding bacteriophages " genomes in motion. <i>International Journal of Medical Microbiology</i> , 2004, 294, 115-121.	1.5	233
5	Shiga-toxin-converting bacteriophages. <i>Research in Microbiology</i> , 2001, 152, 687-695.	1.0	200
6	Non-O157:H7 Pathogenic Shiga Toxin-Producing <i>Escherichia coli</i> : Phenotypic and Genetic Profiling of Virulence Traits and Evidence for Clonality. <i>Journal of Infectious Diseases</i> , 1999, 179, 115-123.	1.9	188
7	Epidemiology and diagnosis of Shiga toxin-producing <i>Escherichia coli</i> infections. <i>Diagnostic Microbiology and Infectious Disease</i> , 1999, 34, 229-243.	0.8	177
8	The large plasmids of Shiga-toxin-producing <i>Escherichia coli</i> (STEC) are highly variable genetic elements. <i>Microbiology (United Kingdom)</i> , 1999, 145, 1005-1014.	0.7	146
9	Structural Analysis of Phage-Borne stx Genes and Their Flanking Sequences in Shiga Toxin-Producing <i>Escherichia coli</i> and <i>Shigella dysenteriae</i> Type 1 Strains. <i>Infection and Immunity</i> , 2000, 68, 4856-4864.	1.0	143
10	Transduction of Enteric <i>Escherichia coli</i> Isolates with a Derivative of Shiga Toxin 2-Encoding Bacteriophage ϕ 3538 Isolated from <i>Escherichia coli</i> O157:H7. <i>Applied and Environmental Microbiology</i> , 1999, 65, 3855-3861.	1.4	142
11	A gene cluster closely related to type II secretion pathway operons of Gram-negative bacteria is located on the large plasmid of enterohemorrhagic <i>Escherichia coli</i> O157 strains. <i>FEMS Microbiology Letters</i> , 2006, 148, 265-272.	0.7	133
12	Enteroaggregative, Shiga Toxin-Producing <i>Escherichia coli</i> O111:H2 Associated with an Outbreak of Hemolytic-Uremic Syndrome. <i>Journal of Clinical Microbiology</i> , 1998, 36, 840-842.	1.8	133
13	Molecular Characteristics and Epidemiological Significance of Shiga Toxin-Producing <i>Escherichia coli</i> O26 Strains. <i>Journal of Clinical Microbiology</i> , 2000, 38, 2134-2140.	1.8	125
14	AmpG, a signal transducer in chromosomal β -lactamase induction. <i>Molecular Microbiology</i> , 1993, 9, 703-715.	1.2	118
15	Antibacterials that are used as growth promoters in animal husbandry can affect the release of Shiga-toxin-2-converting bacteriophages and Shiga toxin 2 from <i>Escherichia coli</i> strains. <i>Microbiology (United Kingdom)</i> , 2000, 146, 1085-1090.	0.7	108
16	Analysis of the EHEC hly operon and its location in the physical map of the large plasmid of enterohaemorrhagic <i>Escherichia coli</i> O157:H7. <i>Microbiology (United Kingdom)</i> , 1996, 142, 907-914.	0.7	102
17	Characterization of a Shiga Toxin 2e-Converting Bacteriophage from an <i>Escherichia coli</i> Strain of Human Origin. <i>Infection and Immunity</i> , 2000, 68, 4850-4855.	1.0	100
18	The large-sized plasmids of enterohemorrhagic <i>Escherichia coli</i> O157 strains encode hemolysins which are presumably members of the <i>E. coli</i> α -hemolysin family. <i>FEMS Microbiology Letters</i> , 1994, 117, 189-196.	0.7	96

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19	Evaluation of lactic acid bacteria for sourdough fermentation of amaranth. <i>International Journal of Food Microbiology</i> , 2009, 136, 75-82.	2.1	94
20	Differentiation in virulence patterns of <i>Escherichia coli</i> possessing <i>eae</i> genes. <i>Medical Microbiology and Immunology</i> , 1994, 183, 23-31.	2.6	89
21	The Nucleotide Sequence of Shiga Toxin (Stx) 2e-Encoding Phage ϕ P27 Is Not Related to Other Stx Phage Genomes, but the Modular Genetic Structure Is Conserved. <i>Infection and Immunity</i> , 2002, 70, 1896-1908.	1.0	88
22	Global Expression of Prophage Genes in <i>Escherichia coli</i> O157:H7 Strain EDL933 in Response to Norfloxacin. <i>Antimicrobial Agents and Chemotherapy</i> , 2005, 49, 931-944.	1.4	79
23	Diarrhea in young children associated with <i>Escherichia coli</i> non-O157 organisms that produce Shiga-like toxin. <i>Journal of Pediatrics</i> , 1996, 128, 341-346.	0.9	71
24	Genetic Analysis of Enteropathogenic and Enterohemorrhagic <i>Escherichia coli</i> Serogroup O103 Strains by Molecular Typing of Virulence and Housekeeping Genes and Pulsed-Field Gel Electrophoresis. <i>Journal of Clinical Microbiology</i> , 2005, 43, 1552-1563.	1.8	70
25	Transduction of Porcine Enteropathogenic <i>Escherichia coli</i> with a Derivative of a Shiga Toxin 2-Encoding Bacteriophage in a Porcine Ligated Ileal Loop System. <i>Applied and Environmental Microbiology</i> , 2003, 69, 7242-7247.	1.4	68
26	Isolation and Characterization of Sorbitol-Fermenting Shiga Toxin (Verocytotoxin)-Producing <i>Escherichia coli</i> O157:H γ strains in the Czech Republic. <i>Journal of Clinical Microbiology</i> , 1998, 36, 2135-2137.	1.8	66
27	Cattle Can Be a Reservoir of Sorbitol-Fermenting Shiga Toxin-Producing <i>Escherichia coli</i> O157:H α Strains and a Source of Human Diseases. <i>Journal of Clinical Microbiology</i> , 2000, 38, 3470-3473.	1.8	66
28	Pore-Forming Properties of the Plasmid-Encoded Hemolysin of Enterohemorrhagic <i>Escherichia coli</i> O157: H7. <i>FEBS Journal</i> , 1996, 241, 594-601.	0.2	65
29	Hemolysin from Shiga toxin-negative <i>Escherichia coli</i> O26 strains injures microvascular endothelium. <i>Microbes and Infection</i> , 2007, 9, 282-290.	1.0	64
30	Anaerobic Conditions Promote Expression of Sfp Fimbriae and Adherence of Sorbitol-Fermenting Enterohemorrhagic <i>Escherichia coli</i> O157:NM to Human Intestinal Epithelial Cells. <i>Applied and Environmental Microbiology</i> , 2008, 74, 1087-1093.	1.4	63
31	Modified Bacteriophage S16 Long Tail Fiber Proteins for Rapid and Specific Immobilization and Detection of <i>Salmonella</i> Cells. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	1.4	59
32	Complete sequence of the large virulence plasmid pSFO157 of the sorbitol-fermenting enterohemorrhagic <i>Escherichia coli</i> O157:H α strain 3072/96. <i>International Journal of Medical Microbiology</i> , 2006, 296, 467-474.	1.5	56
33	Phage-Mediated Shiga Toxin 2 Gene Transfer in Food and Water. <i>Applied and Environmental Microbiology</i> , 2009, 75, 1764-1768.	1.4	55
34	HEp-2 Cell Adherence, Actin Aggregation, and Intimin Types of Attaching and Effacing <i>Escherichia coli</i> Strains Isolated from Healthy Infants in Germany and Australia. <i>Infection and Immunity</i> , 2003, 71, 3995-4002.	1.0	52
35	The Shiga Toxin 1-Converting Bacteriophage BP-4795 Encodes an NleA-Like Type III Effector Protein. <i>Journal of Bacteriology</i> , 2005, 187, 8494-8498.	1.0	47
36	Highly Virulent Non-O157 Enterohemorrhagic <i>Escherichia coli</i> (EHEC) Serotypes Reflect Similar Phylogenetic Lineages, Providing New Insights into the Evolution of EHEC. <i>Applied and Environmental Microbiology</i> , 2015, 81, 7041-7047.	1.4	46

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37	Virulence Genes and Molecular Typing of Different Groups of <i>Escherichia coli</i> O157 Strains in Cattle. <i>Applied and Environmental Microbiology</i> , 2009, 75, 6282-6291.	1.4	45
38	Recent advances in cured raw ham manufacture. <i>Critical Reviews in Food Science and Nutrition</i> , 2018, 58, 610-630.	5.4	45
39	Reduced incidence of postoperative infection after intravenous administration of an immunoglobulin A- and immunoglobulin M-enriched preparation in anergic patients undergoing cardiac surgery. <i>Critical Care Medicine</i> , 1999, 27, 1281-1287.	0.4	45
40	Shiga Toxin-Producing <i>Escherichia coli</i> Infections in Germany. <i>Journal of Food Protection</i> , 1997, 60, 1454-1457.	0.8	44
41	Bacteriophages of Shiga Toxin-Producing <i>Escherichia coli</i> and Their Contribution to Pathogenicity. <i>Pathogens</i> , 2021, 10, 404.	1.2	44
42	<i>Escherichia coli</i> O157:H7 Strain EDL933 Harbors Multiple Functional Prophage-Associated Genes Necessary for the Utilization of 5- <i>N</i> -Acetyl-9- <i>O</i> -Acetyl Neuraminic Acid as a Growth Substrate. <i>Applied and Environmental Microbiology</i> , 2016, 82, 5940-5950.	1.4	40
43	Characterization of two major groups of diarrheagenic <i>Escherichia coli</i> O26 strains which are globally spread in human patients and domestic animals of different species. <i>FEMS Microbiology Letters</i> , 2005, 249, 335-342.	0.7	39
44	Phylogenetic and Molecular Analysis of Food-Borne Shiga Toxin-Producing <i>Escherichia coli</i> . <i>Applied and Environmental Microbiology</i> , 2013, 79, 2731-2740.	1.4	37
45	Molecular Analysis of H Antigens Reveals that Human Diarrheagenic <i>Escherichia coli</i> O26 Strains That Carry the <i>eae</i> Gene Belong to the H11 Clonal Complex. <i>Journal of Clinical Microbiology</i> , 2000, 38, 2989-2993.	1.8	33
46	Genetic structure and chromosomal integration site of the cryptic prophage CP-1639 encoding Shiga toxin 1. <i>Microbiology (United Kingdom)</i> , 2005, 151, 941-950.	0.7	31
47	Development of a rapid detection system for opportunistic pathogenic <i>Cronobacter</i> spp. in powdered milk products. <i>Food Microbiology</i> , 2014, 42, 19-25.	2.1	30
48	Distribution, Functional Expression, and Genetic Organization of Cif, a Phage-Encoded Type III-Secreted Effector from Enteropathogenic and Enterohemorrhagic <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2008, 190, 275-285.	1.0	29
49	Nitrate reductase activity of <i>Staphylococcus carnosus</i> affecting the color formation in cured raw ham. <i>Food Research International</i> , 2016, 85, 113-120.	2.9	29
50	A secretome view of colonisation factors in Shiga toxin-encoding <i>Escherichia coli</i> (STEC): from enterohaemorrhagic <i>E. coli</i> (EHEC) to related enteropathotypes. <i>FEMS Microbiology Letters</i> , 2016, 363, fnw179.	0.7	29
51	Molecular analysis of subtilase cytotoxin genes of food-borne Shiga toxin-producing <i>Escherichia coli</i> reveals a new allelic subAB variant. <i>BMC Microbiology</i> , 2013, 13, 230.	1.3	28
52	How bacterial pathogens of the gastrointestinal tract use the mucosal glyco-code to harness mucus and microbiota: New ways to study an ancient bag of tricks. <i>International Journal of Medical Microbiology</i> , 2020, 310, 151392.	1.5	28
53	Serotypes and intimin types of intestinal and faecal strains of <i>eae+</i> <i>Escherichia coli</i> from weaned pigs. <i>Veterinary Microbiology</i> , 2006, 114, 82-93.	0.8	26
54	Growth Media Simulating Ileal and Colonic Environments Affect the Intracellular Proteome and Carbon Fluxes of Enterohemorrhagic <i>Escherichia coli</i> O157:H7 Strain EDL933. <i>Applied and Environmental Microbiology</i> , 2013, 79, 3703-3715.	1.4	26

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55	Genetics, Toxicity, and Distribution of Enterohemorrhagic <i>Escherichia coli</i> Hemolysin. <i>Toxins</i> , 2019, 11, 502.	1.5	26
56	Molecular Characterization and Distribution of Genes Encoding Members of the Type III Effector NleA Family among Pathogenic <i>Escherichia coli</i> Strains. <i>Journal of Clinical Microbiology</i> , 2007, 45, 2498-2507.	1.8	24
57	Impact of different washing procedures on quality of fresh-cut iceberg lettuce (<i>Lactuca sativa</i> var.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 229-241.	1.6	23
58	Surface adhesins and exopolymers of selected foodborne pathogens. <i>Microbiology (United Kingdom)</i> , 2014, 160, 2561-2582.	0.7	23
59	Lysogenic conversion of atypical enteropathogenic <i>Escherichia coli</i> (aEPEC) from human, murine, and bovine origin with bacteriophage ϕ 3538 [†] stx::cat proves their enterohemorrhagic <i>E. coli</i> (EHEC) progeny. <i>International Journal of Medical Microbiology</i> , 2018, 308, 890-898.	1.5	23
60	Highly conserved B-subunit genes of Shiga-like toxin II variants found in <i>Escherichia coli</i> O157 strains. <i>FEMS Microbiology Letters</i> , 1994, 118, 335-340.	0.7	22
61	Development of a Multiplex Real-Time Polymerase Chain Reaction for Simultaneous Detection of Enterohemorrhagic <i>Escherichia coli</i> and Enteropathogenic <i>Escherichia coli</i> Strains. <i>Foodborne Pathogens and Disease</i> , 2010, 7, 801-808.	0.8	22
62	Evolutionary analysis and distribution of type III effector genes in pathogenic <i>Escherichia coli</i> from human, animal and food sources. <i>Environmental Microbiology</i> , 2011, 13, 439-452.	1.8	22
63	Impact of Fatty Acid Chain Length of Rosmarinate Esters on Their Antimicrobial Activity against <i>Staphylococcus camosus</i> LTH1502 and <i>Escherichia coli</i> K-12 LTH4263. <i>Journal of Food Protection</i> , 2013, 76, 1539-1548.	0.8	22
64	A Novel Glutamyl (Aspartyl)-Specific Aminopeptidase A from <i>Lactobacillus delbrueckii</i> with Promising Properties for Application. <i>PLoS ONE</i> , 2016, 11, e0152139.	1.1	22
65	Characterization of in vitro antifungal activities of small and American cranberry (<i>Vaccinium</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 sugar reduced fruit spreads. <i>International Journal of Food Microbiology</i> , 2015, 204, 111-117.	2.1	20
66	Quality Improvement of Fresh-Cut Endive (<i>Cichorium endivia</i> L.) and Recycling of Washing Water by Low-Dose UV-C Irradiation. <i>Food and Bioprocess Technology</i> , 2016, 9, 1979-1990.	2.6	20
67	Effect of Water Jet Cutting and Moderate Heat Treatment on Quality of Fresh-Cut Red Oak Leaf Lettuce (<i>Lactuca sativa</i> L. var. <i>crispa</i>). <i>Food and Bioprocess Technology</i> , 2014, 7, 3478-3492.	2.6	19
68	Identification and characterisation of <i>Escherichia coli</i> strains of O157 and non-O157 serogroups containing three distinct Shiga toxin genes. <i>Journal of Medical Microbiology</i> , 2000, 49, 383-386.	0.7	19
69	Growth advantage of <i>Escherichia coli</i> O104:H4 strains on 5- N -acetyl-9- O -acetyl neuraminic acid as a carbon source is dependent on heterogeneous phage-Borne nanS-p esterases. <i>International Journal of Medical Microbiology</i> , 2018, 308, 459-468.	1.5	18
70	The signal transducer encoded by <i>ampG</i> is essential for induction of chromosomal AmpC β -lactamase in <i>Escherichia coli</i> by β -lactam antibiotics and unspecific inducers. <i>Microbiology (United Kingdom)</i> , 1995, 141, 1085-1092.	0.7	17
71	Protective effects of <i>Lactobacilli</i> , <i>Bifidobacteria</i> and <i>Staphylococci</i> on the infection of cultured HT29 cells with different enterohemorrhagic <i>Escherichia coli</i> serotypes are strain-specific. <i>International Journal of Food Microbiology</i> , 2010, 144, 133-140.	2.1	17
72	Cytotoxic and Apoptotic Effects of Recombinant Subtilase Cytotoxin Variants of Shiga Toxin-Producing <i>Escherichia coli</i> . <i>Infection and Immunity</i> , 2015, 83, 2338-2349.	1.0	17

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73	Safety assessment of selected <i>Staphylococcus carnosus</i> strains with regard to their application as meat starter culture. <i>Food Control</i> , 2016, 66, 93-99.	2.8	17
74	Bacteriophage 933W encodes a functional esterase downstream of the Shiga toxin 2a operon. <i>International Journal of Medical Microbiology</i> , 2014, 304, 269-274.	1.5	16
75	Quality of fresh-cut radicchio cv. Rosso di Chioggia (<i>Cichorium intybus</i> L. var. <i>foliosum</i> Hegi) as affected by water jet cutting and different washing procedures. <i>European Food Research and Technology</i> , 2015, 240, 159-172.	1.6	16
76	Antimicrobial Mechanism and Activity of Dodecyl Rosmarinate against <i>Staphylococcus carnosus</i> LTH1502 as Influenced by Addition of Salt and Change in pH. <i>Journal of Food Protection</i> , 2014, 77, 444-452.	0.8	15
77	Prevalence of subtilase cytotoxin-encoding subAB variants among Shiga toxin-producing <i>Escherichia coli</i> strains isolated from wild ruminants and sheep differs from that of cattle and pigs and is predominated by the new allelic variant subAB2-2. <i>International Journal of Medical Microbiology</i> , 2015, 305, 124-128.	1.5	15
78	Selection of <i>Staphylococcus carnosus</i> strains based on in vitro analysis of technologically relevant physiological activities. <i>Annals of Microbiology</i> , 2016, 66, 479-487.	1.1	15
79	Toxins of Locus of Enterocyte Effacement-Negative Shiga Toxin-Producing <i>Escherichia coli</i> . <i>Toxins</i> , 2018, 10, 241.	1.5	15
80	Adherence factors of enterohemorrhagic <i>Escherichia coli</i> O157:H7 strain Sakai influence its uptake into the roots of <i>Valerianella locusta</i> grown in soil. <i>Food Microbiology</i> , 2018, 76, 245-256.	2.1	15
81	Effects of <i>Quillaja saponaria</i> extract and N ^ε -lauroyl-L-arginine ethyl ester on reducing selected foodborne pathogens in vitro and maintaining quality of fresh-cut endive (<i>Cichorium endivia</i> L.) at pilot plant scale. <i>Food Control</i> , 2017, 73, 393-400.	2.8	14
82	Kinetics of migration of colloidal particles in meat muscles in the absence and presence of a proteolytic enzyme to simulate non-motile bacteria penetration. <i>Food Research International</i> , 2015, 75, 79-88.	2.9	13
83	Regulation of <i>nleA</i> in Shiga Toxin-Producing <i>Escherichia coli</i> O84:H4 Strain 4795/97. <i>Journal of Bacteriology</i> , 2011, 193, 832-841.	1.0	12
84	Effects of gallotannin treatment on attachment, growth, and survival of <i>Escherichia coli</i> O157:H7 and <i>Listeria monocytogenes</i> on spinach and lettuce. <i>European Food Research and Technology</i> , 2012, 234, 1081-1090.	1.6	12
85	Type III effector genes and other virulence factors of Shiga toxin-encoding <i>Escherichia coli</i> isolated from wastewater. <i>Environmental Microbiology Reports</i> , 2012, 4, 147-155.	1.0	12
86	Antimicrobial effect of lauroyl arginate ethyl on <i>Escherichia coli</i> O157:H7 and <i>Listeria monocytogenes</i> on red oak leaf lettuce. <i>European Food Research and Technology</i> , 2017, 243, 879-887.	1.6	12
87	Endemic occurrence of infections by multidrug-resistant <i>Escherichia coli</i> of four unique serotypes in the elderly population of Israel. <i>FEMS Microbiology Letters</i> , 2004, 239, 249-254.	0.7	11
88	Repression of the Locus of the Enterocyte Effacement-Encoded Regulator of Gene Transcription of <i>Escherichia coli</i> O157:H7 by <i>Lactobacillus reuteri</i> Culture Supernatants Is LuxS and Strain Dependent. <i>Applied and Environmental Microbiology</i> , 2008, 74, 3310-3314.	1.4	11
89	Plant variety and soil type influence <i>Escherichia coli</i> O104:H4 strain C227/11 ^{cu} adherence to and internalization into the roots of lettuce plants. <i>Food Microbiology</i> , 2020, 86, 103316.	2.1	11
90	Survival of spoilage bacteria subjected to sequential eugenol and temperature treatments. <i>International Journal of Food Microbiology</i> , 2016, 218, 6-16.	2.1	10

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91	Influence of application sequence and timing of eugenol and lauric arginate (LAE) on survival of spoilage organisms. <i>Food Microbiology</i> , 2017, 64, 210-218.	2.1	10
92	Bacteriophages as modulator for the human gut microbiota: Release from dairy food systems and survival in a dynamic human gastrointestinal model. <i>LWT - Food Science and Technology</i> , 2018, 91, 235-241.	2.5	10
93	The Role of Bacteriophages in the Generation and Spread of Bacterial Pathogens. , 2008, , 79-112.		9
94	Comparison of ultra-high-pressure water jet and conventional rotating blade cutting for the production of fresh-cut iceberg (<i>Lactuca sativa</i> L.) and endive (<i>Cichorium endivia</i> L.). <i>European Food Research and Technology</i> , 2016, 242, 2071-2081.	1.6	9
95	Subtilase contributes to the cytotoxicity of a Shiga toxin-producing <i>Escherichia coli</i> strain encoding three different toxins. <i>International Journal of Food Microbiology</i> , 2016, 217, 156-161.	2.1	8
96	Effect of mechanical curing treatments on particle distribution to simulate non-motile bacteria migration in cured raw ham. <i>Journal of Food Engineering</i> , 2017, 194, 58-66.	2.7	8
97	Amplification methods in diagnostic bacteriology (selected examples). <i>Journal of Microbiological Methods</i> , 1995, 23, 55-73.	0.7	7
98	Mechanisms of enterohemorrhagic <i>Escherichia coli</i> spread along the food-chain and precautionary measures. <i>Journal Fur Verbraucherschutz Und Lebensmittelsicherheit</i> , 2011, 6, 503-510.	0.5	7
99	Genetic Background and Mobility of Variants of the Gene <i>nleA</i> in Attaching and Effacing <i>Escherichia coli</i> . <i>Applied and Environmental Microbiology</i> , 2011, 77, 8705-8713.	1.4	7
100	Draft Genome Sequences of Five Shiga Toxin-Producing <i>Escherichia coli</i> Isolates Harboring the New and Recently Described Subtilase Cytotoxin Allelic Variant subAB 2-3. <i>Genome Announcements</i> , 2017, 5, .	0.8	7
101	Molecular analysis of nosocomial infection by oxacillin-resistant <i>Staphylococcus aureus</i> lacking protein A and clumping factor. <i>Lancet, The</i> , 1992, 340, 621.	6.3	6
102	Variation of the <i>Pseudomonas</i> community structure on oak leaf lettuce during storage detected by culture-dependent and -independent methods. <i>International Journal of Food Microbiology</i> , 2016, 216, 95-103.	2.1	6
103	Determination of virulence and fitness genes associated with the <i>pheU</i> , <i>pheV</i> and <i>selC</i> integration sites of LEE-negative food-borne Shiga toxin-producing <i>Escherichia coli</i> strains. <i>Gut Pathogens</i> , 2018, 10, 43.	1.6	6
104	Differential transcriptome analysis of enterohemorrhagic <i>Escherichia coli</i> strains reveals differences in response to plant-derived compounds. <i>BMC Microbiology</i> , 2019, 19, 212.	1.3	6
105	Effect of exopolysaccharides produced by <i>Lactobacillus sanfranciscensis</i> on the processing properties of wheat doughs. <i>European Food Research and Technology</i> , 2020, 246, 461-469.	1.6	6
106	Antibody Reactivity of a Standardized Human Serum Protein Solution Against a Spectrum of Microbial Pathogens and Toxins: Comparison with Fresh Frozen Plasma. <i>Therapeutic Apheresis and Dialysis</i> , 2002, 6, 145-153.	0.4	5
107	Genetic diversity and population structure of food-borne <i>Staphylococcus carnosus</i> strains. <i>Systematic and Applied Microbiology</i> , 2017, 40, 34-41.	1.2	5
108	Kinetics of volatile marker compounds during ripening of cured loins inoculated with <i>Staphylococcus carnosus</i> . <i>Journal of the Science of Food and Agriculture</i> , 2017, 97, 3050-3057.	1.7	5

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109	Transcription of the Subtilase Cytotoxin Gene <i>stxAB1</i> in Shiga Toxin-Producing <i>Escherichia coli</i> Is Dependent on <i>hfq</i> and <i>hns</i> . <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	5
110	Variants of <i>Escherichia coli</i> Subtilase Cytotoxin Subunits Show Differences in Complex Formation In Vitro. <i>Toxins</i> , 2019, 11, 703.	1.5	5
111	Effects of Protein, Calcium, and pH on Gene Transcription, Cell-Envelope Peptidase Activity of <i>Lactococcus lactis</i> Strains, and the Formation of Bitter Peptides. <i>Foods</i> , 2021, 10, 1588.	1.9	5
112	Survival of Enterohemorrhagic <i>Escherichia coli</i> O104:H4 Strain C227/111 in Agricultural Soils Depends on <i>rpoS</i> and Environmental Factors. <i>Pathogens</i> , 2021, 10, 1443.	1.2	4
113	IgA/IgM and Secretory Immunity. <i>Sepsis</i> , 1999, 3, 219-224.	0.5	3
114	Detection and Characterization of EHEC-Hemolysin. , 2003, 73, 151-164.		3
115	Phage-bacterium Co-evolution and Its Implication for Bacterial Pathogenesis. , 0, , 49-78.		3
116	Analysis of the survival of <i>Listeria monocytogenes</i> in food-grade lubricants. <i>European Food Research and Technology</i> , 2012, 234, 323-331.	1.6	3
117	The enzyme subunit SubA of Shiga toxin-producing <i>E. coli</i> strains demonstrates comparable intracellular transport and cytotoxic activity as the holotoxin SubAB in HeLa and HCT116 cells in vitro. <i>Archives of Toxicology</i> , 2021, 95, 975-983.	1.9	3
118	Comparison of net growth of Shiga toxin-producing <i>Escherichia coli</i> strains of serogroups O26, O103, and O157 in ground meat at different temperatures. <i>European Food Research and Technology</i> , 2014, 238, 163-168.	1.6	2
119	Draft Genome Sequence of <i>Staphylococcus carnosus</i> subsp. <i>utilis</i> LTH 7013, Isolated from South Tyrolean Ham. <i>Genome Announcements</i> , 2015, 3, .	0.8	2
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121	Morphological and Dose-Dependent Study on the Effect of Methyl, Hexyl, and Dodecyl Rosmarinate on <i>Staphylococcus carnosus</i> LTH1502: Use of the Weibull Model. <i>Journal of Food Protection</i> , 2018, 81, 598-605.	0.8	2
122	Application of MALDI-TOF mass spectrometry and specific PCR for tracking of <i>E. coli</i> O157:H ⁺ strain 431/97 in Batavia lettuce. <i>Chemical and Biological Technologies in Agriculture</i> , 2019, 6, .	1.9	2
123	Effect of Different Wash Water Additives and Deep-Frozen Storage on the Quality of Curly Parsley (<i>Petroselinum crispum</i> var. <i>crispum</i>). <i>Food and Bioprocess Technology</i> , 2019, 12, 158-165.	2.6	2
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126	Disturbance of Peptidoglycan Synthesis by Glycine and D-Methionine Creates a Signal for the ampG-Mediated Induction of AmpC- β -Lactamase in <i>Escherichia coli</i> . , 1993, , 341-346.		1

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127	Genomic or Pathogenicity Islands in <i>Streptococcus pneumoniae</i> . , 0, , 217-236.		0
128	Novel Aspects of the SubA Subunit of the Subtilase Cytotoxin. <i>Toxins</i> , 2022, 14, 156.	1.5	0