Susanne HÃ¤ussler

List of Publications by Year
in descending order

131 papers	7,402 citations	50276 46 h-index	66911 78 g-index
137 all docs	137 docs citations	137 times ranked	8388 citing authors

#	Article	IF	Citations
1	Quorum-Sensing Antagonistic Activities of Azithromycin in Pseudomonas aeruginosa PAO1: a Global Approach. Antimicrobial Agents and Chemotherapy, 2006, 50, 1680-1688.	3.2	279
2	Î²-Lactam Resistance Response Triggered by Inactivation of a Nonessential Penicillin-Binding Protein. PLoS Pathogens, 2009, 5, e1000353.	4.7	258
3	The Pseudomonas aeruginosa quinolone signal (PQS) has an iron-chelating activity. Environmental Microbiology, 2006, 8, 1318-1329.	3.8	239
4	The Pseudomonas aeruginosa Transcriptome in Planktonic Cultures and Static Biofilms Using RNA Sequencing. PLoS ONE, 2012, 7, e31092.	2.5	212
5	The Pseudomonas Quinolone Signal (PQS) Balances Life and Death in Pseudomonas aeruginosa Populations. PLoS Pathogens, 2008, 4, e1000166.	4.7	205
6	Pseudomonas aeruginosa Ceftolozane-Tazobactam Resistance Development Requires Multiple Mutations Leading to Overexpression and Structural Modification of AmpC. Antimicrobial Agents and Chemotherapy, 2014, 58, 3091-3099.	3.2	197
7	Highly adherent small-colony variants of Pseudomonas aeruginosa in cystic fibrosis lung infection. Journal of Medical Microbiology, 2003, 52, 295-301.	1.8	194
8	Global regulation of gene expression by OxyR in an important human opportunistic pathogen. Nucleic Acids Research, 2012, 40, 4320-4333.	14.5	189
9	Quantitative image analysis of microbial communities with BiofilmQ. Nature Microbiology, 2021, 6, 151-156.	13.3	181
10	IL-17â€“induced CXCL12 recruits B cells and induces follicle formation in BALT in the absence of differentiated FDCs. Journal of Experimental Medicine, 2014, 211, 643-651.	8.5	159
11	RhlR Expression in Pseudomonas aeruginosa Is Modulated by the Pseudomonas Quinolone Signal via PhoB-Dependent and -Independent Pathways. Journal of Bacteriology, 2006, 188, 8601-8606.	2.2	141
12	Elucidation of Sigma Factor-Associated Networks in Pseudomonas aeruginosa Reveals a Modular Architecture with Limited and Function-Specific Crosstalk. PLoS Pathogens, 2015, 11, e1004744.	4.7	134
13	Critical Assessment of Metagenome Interpretation: the second round of challenges. Nature Methods, 2022, 19, 429-440.	19.0	133
14	Quantitative Contributions of Target Alteration and Decreased Drug Accumulation to Pseudomonas aeruginosa Fluoroquinolone Resistance. Antimicrobial Agents and Chemotherapy, 2013, 57, 1361-1368.	3.2	130
15	7- O -Malonyl Macrolactin A, a New Macrolactin Antibiotic from Bacillus subtilis Active against Methicillin-Resistant Staphylococcus aureus , Vancomycin-Resistant Enterococci, and a Small-Colony Variant of Burkholderia cepacia. Antimicrobial Agents and Chemotherapy, 2006, 50, 1701-1709.	3.2	129
16	Biofilm formation by the small colony variant phenotype of Pseudomonas aeruginosa. Environmental Microbiology, 2004, 6, 546-551.	3.8	124
17	A 96-well-plateâ€“based optical method for the quantitative and qualitative evaluation of Pseudomonas aeruginosa biofilm formation and its application to susceptibility testing. Nature Protocols, 2010, 5, 1460-1469.	12.0	119
18	Arginine-rhamnosylation as new strategy to activate translation elongation factor P. Nature Chemical Biology, 2015, 11, 266-270.	8.0	116

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19	Biosynthetic Pathway of Pseudomonas aeruginosa 4-Hydroxy-2-Alkylquinolines. Journal of Bacteriology, 2005, 187, 3630-3635.	2.2	115
20	Validation of PqsD as an Anti-biofilm Target in Pseudomonas aeruginosa by Development of Small-Molecule Inhibitors. Journal of the American Chemical Society, 2012, 134, 16143-16146.	13.7	113
21	Predicting antimicrobial resistance in Pseudomonas aeruginosa with machine learningâ€enabled molecular diagnostics. EMBO Molecular Medicine, 2020, 12, e10264.	6.9	111
22	Discovery of Antagonists of PqsR, a Key Player in 2-Alkyl-4-quinolone-Dependent Quorum Sensing in Pseudomonas aeruginosa. Chemistry and Biology, 2012, 19, 381-390.	6.0	110
23	Genomewide Identification of Genetic Determinants of Antimicrobial Drug Resistance in Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2009, 53, 2522-2531.	3.2	108
24	Pseudomonas aeruginosa cupA-encoded fimbriae expression is regulated by a GGDEF and EAL domain-dependent modulation of the intracellular level of cyclic diguanylate. Environmental Microbiology, 2007, 9, 2475-2485.	3.8	107
25	The LasB Elastase of Pseudomonas aeruginosa Acts in Concert with Alkaline Protease AprA To Prevent Flagellin-Mediated Immune Recognition. Infection and Immunity, 2016, 84, 162-171.	2.2	106
26	The YfiBNR Signal Transduction Mechanism Reveals Novel Targets for the Evolution of Persistent Pseudomonas aeruginosa in Cystic Fibrosis Airways. PLoS Pathogens, 2012, 8, e1002760.	4.7	105
27	Pseudomonas aeruginosa population structure revisited under environmental focus: impact of water quality and phage pressure. Environmental Microbiology, 2012, 14, 1952-1967.	3.8	104
28	Production of medium chain length polyhydroxyalkanoate in metabolic flux optimized Pseudomonas putida. Microbial Cell Factories, 2014, 13, 88.	4.0	98
29	Genetic determinants of Pseudomonas aeruginosa biofilm establishment. Microbiology (United) Tj ETQq1 1 0.784314 rgBT /Overlock	1.8	97
30	Expression Analysis of a Highly Adherent and Cytotoxic Small Colony Variant of Pseudomonas aeruginosa Isolated from a Lung of a Patient with Cystic Fibrosis. Journal of Bacteriology, 2004, 186, 3837-3847.	2.2	93
31	Phenotypic and Genome-Wide Analysis of an Antibiotic-Resistant Small Colony Variant (SCV) of Pseudomonas aeruginosa. PLoS ONE, 2011, 6, e29276.	2.5	81
32	The Pseudomonas aeruginosa Transcriptional Landscape Is Shaped by Environmental Heterogeneity and Genetic Variation. MBio, 2015, 6, e00749.	4.1	73
33	Genetically diverse Pseudomonas aeruginosa populations display similar transcriptomic profiles in a cystic fibrosis explanted lung. Nature Communications, 2019, 10, 3397.	12.8	68
34	Transcriptome Profiling of Antimicrobial Resistance in Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2016, 60, 4722-4733.	3.2	67
35	Inter- and Intraclonal Diversity of the Pseudomonas aeruginosa Proteome Manifests within the Secretome. Journal of Bacteriology, 2003, 185, 5807-5814.	2.2	65
36	Structure of an Acidic Exopolysaccharide of Burkholderia pseudomallei. FEBS Journal, 1997, 250, 608-616.	0.2	63

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37	In Vivo mRNA Profiling of Uropathogenic Escherichia coli from Diverse Phylogroups Reveals Common and Group-Specific Gene Expression Profiles. MBio, 2014, 5, e01075-14.	4.1	63
38	aroA -Deficient Salmonella enterica Serovar Typhimurium Is More Than a Metabolically Attenuated Mutant. MBio, 2016, 7, .	4.1	62
39	Biofilms 2012: New Discoveries and Significant Wrinkles in a Dynamic Field. Journal of Bacteriology, 2013, 195, 2947-2958.	2.2	59
40	The Pseudomonas aeruginosa Chemotaxis Methyltransferase CheR1 Impacts on Bacterial Surface Sampling. PLoS ONE, 2011, 6, e18184.	2.5	59
41	Structure Elucidation and Preliminary Assessment of Hydrolase Activity of PqsE, thePseudomonasQuinolone Signal (PQS) Response Protein. Biochemistry, 2009, 48, 10298-10307.	2.5	57
42	The PqsR and RhlR Transcriptional Regulators Determine the Level of Pseudomonas Quinolone Signal Synthesis in Pseudomonas aeruginosa by Producing Two Different pqsABCDE mRNA Isoforms. Journal of Bacteriology, 2014, 196, 4163-4171.	2.2	57
43	The extensive set of accessoryPseudomonas aeruginosagenomic components. FEMS Microbiology Letters, 2014, 356, 235-241.	1.8	55
44	Identification of the Alternative Sigma Factor SigX Regulon and Its Implications for Pseudomonas aeruginosa Pathogenicity. Journal of Bacteriology, 2014, 196, 345-356.	2.2	55
45	Cross talk between the response regulators PhoB and TctD allows for the integration of diverse environmental signals inPseudomonas aeruginosa. Nucleic Acids Research, 2015, 43, 6413-6425.	14.5	54
46	Regulation of Flagellum Biosynthesis in Response to Cell Envelope Stress in Salmonella enterica Serovar Typhimurium. MBio, 2018, 9, .	4.1	53
47	A chemical proteomics approach to identify c-di-GMP binding proteins in Pseudomonas aeruginosa. Journal of Microbiological Methods, 2012, 88, 229-236.	1.6	52
48	Multiple roles of Pseudomonas aeruginosa TBCF10839 PilY1 in motility, transport and infection. Molecular Microbiology, 2009, 71, 730-747.	2.5	50
49	Biofilms 2009: New Perspectives at the Heart of Surface-Associated Microbial Communities. Journal of Bacteriology, 2010, 192, 2941-2949.	2.2	50
50	The virulence factor LecB varies in clinical isolates: consequences for ligand binding and drug discovery. Chemical Science, 2016, 7, 4990-5001.	7.4	50
51	Constitutive production of c-di-GMP is associated with mutations in a variant of Pseudomonas aeruginosa with altered membrane composition. Science Signaling, 2015, 8, ra36.	3.6	49
52	Mining zebrafish microbiota reveals key community-level resistance against fish pathogen infection. ISME Journal, 2021, 15, 702-719.	9.8	49
53	Evolutionary conservation of essential and highly expressed genes in Pseudomonas aeruginosa. BMC Genomics, 2010, 11, 234.	2.8	48
54	Contribution of Veillonella parvula to Pseudomonas aeruginosa-Mediated Pathogenicity in a Murine Tumor Model System. Infection and Immunity, 2015, 83, 417-429.	2.2	47

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55	Inoculation density and nutrient level determine the formation of mushroom-shaped structures in Pseudomonas aeruginosa biofilms. Scientific Reports, 2016, 6, 32097.	3.3	47
56	cGAS-Mediated Innate Immunity Spreads Intercellularly through HIV-1 Env-Induced Membrane Fusion Sites. Cell Host and Microbe, 2016, 20, 443-457.	11.0	46
57	Highly resistant Burkholderia pseudomallei small colony variants isolated in vitro and in experimental melioidosis. Medical Microbiology and Immunology, 1999, 188, 91-97.	4.8	45
58	Cross-regulation by CrcZ RNA controls anoxic biofilm formation in Pseudomonas aeruginosa. Scientific Reports, 2016, 6, 39621.	3.3	45
59	Use of Single-Frequency Impedance Spectroscopy to Characterize the Growth Dynamics of Biofilm Formation in Pseudomonas aeruginosa. Scientific Reports, 2017, 7, 5223.	3.3	44
60	RNASeq Based Transcriptional Profiling of Pseudomonas aeruginosa PA14 after Short- and Long-Term Anoxic Cultivation in Synthetic Cystic Fibrosis Sputum Medium. PLoS ONE, 2016, 11, e0147811.	2.5	42
61	An oral multispecies biofilm model for high content screening applications. PLoS ONE, 2017, 12, e0173973.	2.5	42
62	Ex vivo transcriptional profiling reveals a common set of genes important for the adaptation of <scp>P</scp>seudomonas aeruginosa to chronically infected host sites. Environmental Microbiology, 2013, 15, 570-587.	3.8	41
63	Recent Advances and Current Trends in Nucleotide Second Messenger Signaling in Bacteria. Journal of Molecular Biology, 2019, 431, 908-927.	4.2	41
64	Effects of Green Tea Compound Epigallocatechin-3-Gallate against Stenotrophomonas maltophilia Infection and Biofilm. PLoS ONE, 2014, 9, e92876.	2.5	40
65	Human airway mucus alters susceptibility of Pseudomonas aeruginosa biofilms to tobramycin, but not colistin. Journal of Antimicrobial Chemotherapy, 2018, 73, 2762-2769.	3.0	39
66	BACTOMEâ€"a reference database to explore the sequence- and gene expression-variation landscape ofPseudomonas aeruginosaclinical isolates. Nucleic Acids Research, 2019, 47, D716-D720.	14.5	38
67	Multicellular signalling and growth of Pseudomonas aeruginosa. International Journal of Medical Microbiology, 2010, 300, 544-548.	3.6	37
68	Parallel evolutionary paths to produce more than one Pseudomonas aeruginosa biofilm phenotype. Npj Biofilms and Microbiomes, 2020, 6, 2.	6.4	36
69	Extensively Drug-Resistant Pseudomonas aeruginosa Isolates Containing bla _{VIM-2} and Elements of Salmonella Genomic Island 2: a New Genetic Resistance Determinant in Northeast Ohio. Antimicrobial Agents and Chemotherapy, 2014, 58, 5929-5935.	3.2	34
70	Biofilm formation by Pseudomonas aeruginosa in solid murine tumors â€“ a novel model system. Microbes and Infection, 2012, 14, 951-958.	1.9	32
71	Identification of a Pseudomonas aeruginosa PAO1 DNA Methyltransferase, Its Targets, and Physiological Roles. MBio, 2017, 8, .	4.1	32
72	Deep transcriptome profiling of clinical <scp>K</scp>lebsiella pneumoniae isolates reveals strain and sequence typeâ€specific adaptation. Environmental Microbiology, 2015, 17, 4690-4710.	3.8	31

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73	Azetidineâ€Containing Alkaloids Produced by a Quorumâ€Sensing Regulated Nonribosomal Peptide Synthetase Pathway in Pseudomonas aeruginosa. Angewandte Chemie - International Edition, 2019, 58, 3178-3182.	13.8	31
74	Mutation in Elongation Factor G Confers Resistance to the Antibiotic Argyrin in the Opportunistic Pathogen Pseudomonas aeruginosa. ChemBioChem, 2012, 13, 2339-2345.	2.6	30
75	In VivoEfficacy of Antimicrobials against Biofilm-Producing Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2015, 59, 4974-4981.	3.2	30
76	An orphan sensor kinase controls quinolone signal production via MexT in Pseudomonas aeruginosa. Molecular Microbiology, 2012, 83, 536-547.	2.5	29
77	Insights into hostâ€“pathogen interactions from stateâ€ofâ€theâ€art animal models of respiratory Pseudomonas aeruginosa infections. FEBS Letters, 2016, 590, 3941-3959.	2.8	29
78	Detrimental Effect of Type I IFNs During Acute Lung Infection With Pseudomonas aeruginosa Is Mediated Through the Stimulation of Neutrophil NETosis. Frontiers in Immunology, 2019, 10, 2190.	4.8	29
79	Neutrophil elastase-mediated increase in airway temperature during inflammation. Journal of Cystic Fibrosis, 2014, 13, 623-631.	0.7	26
80	The Core Proteome of Biofilm-Grown Clinical Pseudomonas aeruginosa Isolates. Cells, 2019, 8, 1129.	4.1	26
81	Establishment of an induced memory response in Pseudomonas aeruginosa during infection of a eukaryotic host. ISME Journal, 2019, 13, 2018-2030.	9.8	26
82	The Pho regulon influences biofilm formation and type three secretion in Pseudomonas aeruginosa. Environmental Microbiology Reports, 2009, 1, 488-494.	2.4	23
83	Recycling of Peptidyl-tRNAs by Peptidyl-tRNA Hydrolase Counteracts Azithromycin-Mediated Effects on Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2013, 57, 1617-1624.	3.2	23
84	Breaking the Vicious Cycle of Antibiotic Killing and Regrowth of Biofilm-Residing Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	23
85	Nonâ€invasive, ratiometric determination of intracellular pH in Pseudomonas species using a novel genetically encoded indicator. Microbial Biotechnology, 2019, 12, 799-813.	4.2	23
86	Importance of flagella in acute and chronic Pseudomonas aeruginosa infections. Environmental Microbiology, 2019, 21, 883-897.	3.8	23
87	Evaluation of a microarray-hybridization based method applicable for discovery of single nucleotide polymorphisms (SNPs) in the Pseudomonas aeruginosa genome. BMC Genomics, 2009, 10, 29.	2.8	22
88	Functional modules of sigma factor regulons guarantee adaptability and evolvability. Scientific Reports, 2016, 6, 22212.	3.3	22
89	Diagnostics and Resistance Profiling of Bacterial Pathogens. Current Topics in Microbiology and Immunology, 2016, 398, 89-102.	1.1	22
90	The peptide chain release factor methyltransferase <scp>PrmC</scp> is essential for pathogenicity and environmental adaptation of <scp>P</scp>seudomonas aeruginosa<scp>PA</scp>14. Environmental Microbiology, 2013, 15, 597-609.	3.8	21

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91	Towards individualized diagnostics of biofilm-associated infections: a case study. Npj Biofilms and Microbiomes, 2017, 3, 22.	6.4	21
92	Untargeted LC-MS Metabolomics Differentiates Between Virulent and Avirulent Clinical Strains of Pseudomonas aeruginosa. Biomolecules, 2020, 10, 1041.	4.0	21
93	Transcriptional and Mutational Profiling of an Aminoglycoside-Resistant Pseudomonas aeruginosa Small-Colony Variant. Antimicrobial Agents and Chemotherapy, 2017, 61, .	3.2	19
94	Environmentâ€driven changes of mRNA and protein levels in Pseudomonas aeruginosa. Environmental Microbiology, 2018, 20, 3952-3963.	3.8	19
95	Production of Norspermidine Contributes to Aminoglycoside Resistance in pmrAB Mutants of Pseudomonas aeruginosa. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	18
96	Evolution of biofilm-adapted gene expression profiles in lasR-deficient clinical Pseudomonas aeruginosa isolates. Npj Biofilms and Microbiomes, 2022, 8, 6.	6.4	17
97	Genetic determinants of Pseudomonas aeruginosa fitness during biofilm growth. Biofilm, 2020, 2, 100023.	3.8	16
98	Chromatin Immunoprecipitation for ChIP-chip and ChIP-seq. Methods in Molecular Biology, 2014, 1149, 591-605.	0.9	16
99	Exopolysaccharides of Burkholderia pseudomallei. Acta Tropica, 2000, 74, 211-214.	2.0	15
100	Application of Synthetic Peptide Arrays To Uncover Cyclic Di-GMP Binding Motifs. Journal of Bacteriology, 2016, 198, 138-146.	2.2	15
101	The Anaerobically Induced sRNA Pail Affects Denitrification in Pseudomonas aeruginosa PA14. Frontiers in Microbiology, 2017, 8, 2312.	3.5	14
102	Expression of the MexXY Aminoglycoside Efflux Pump and Presence of an Aminoglycoside-Modifying Enzyme in Clinical Pseudomonas aeruginosa Isolates Are Highly Correlated. Antimicrobial Agents and Chemotherapy, 2020, 65, .	3.2	14
103	Identification of Burkholderia cepacia complex pathogens by rapid-cycle PCR with fluorescent hybridization probes. Journal of Medical Microbiology, 2006, 55, 721-727.	1.8	13
104	Complete Genome Sequence of Highly Adherent Pseudomonas aeruginosa Small-Colony Variant SCV20265. Genome Announcements, 2014, 2, .	0.8	13
105	Targeting bioenergetics is key to counteracting the drug-tolerant state of biofilm-grown bacteria. PLoS Pathogens, 2020, 16, e1009126.	4.7	13
106	A German external quality survey of diagnostic microbiology of respiratory tract infections in patients with cystic fibrosis. Journal of Cystic Fibrosis, 2008, 7, 7-14.	0.7	12
107	Global Genotype-Phenotype Correlations in Pseudomonas aeruginosa. PLoS Pathogens, 2010, 6, e1001074.	4.7	12
108	Spatiotemporal control of FlgZ activity impacts Pseudomonas aeruginosa flagellar motility. Molecular Microbiology, 2019, 111, 1544-1557.	2.5	12

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109	Dual Effect: High NADH Levels Contribute to Efflux-Mediated Antibiotic Resistance but Drive Lethality Mediated by Reactive Oxygen Species. MBio, 2022, 13, e0243421.	4.1	12
110	Evolution of Pseudomonas aeruginosa toward higher fitness under standard laboratory conditions. ISME Journal, 2021, 15, 1165-1177.	9.8	11
111	Identification and Quantification of (t)RNA Modifications in Pseudomonas aeruginosa by Liquid Chromatographyâ€“Tandem Mass Spectrometry. ChemBioChem, 2019, 20, 1430-1437.	2.6	10
112	Single-Nucleotide Polymorphism-Based Genetic Diversity Analysis of Clinical Pseudomonas aeruginosa Isolates. Genome Biology and Evolution, 2020, 12, 396-406.	2.5	10
113	Antisense transcription in Pseudomonas aeruginosa. Microbiology (United Kingdom), 2018, 164, 889-895.	1.8	10
114	Monoclonal IgA class-switch variants against bacterial surface antigens: molecular forms and transport into murine respiratory secretions. European Journal of Immunology, 1994, 24, 2855-2862.	2.9	9
115	Pseudomonas aeruginosa LysR PA4203 Regulator NmoR Acts as a Repressor of the PA4202 nmoA Gene, Encoding a Nitronate Monooxygenase. Journal of Bacteriology, 2015, 197, 1026-1039.	2.2	9
116	Pseudomonas aeruginosa Is More Tolerant Under Biofilm Than Under Planktonic Growth Conditions: A Multi-Isolate Survey. Frontiers in Cellular and Infection Microbiology, 2022, 12, 851784.	3.9	9
117	Comprehensive MALDI-TOF Biotyping of the Non-Redundant Harvard Pseudomonas aeruginosa PA14 Transposon Insertion Mutant Library. PLoS ONE, 2015, 10, e0117144.	2.5	7
118	The immunogenic potential of bacterial flagella for Salmonellaâ€mediated tumor therapy. International Journal of Cancer, 2020, 147, 448-460.	5.1	7
119	The Two-Component System 09 Regulates Pneumococcal Carbohydrate Metabolism and Capsule Expression. Microorganisms, 2021, 9, 468.	3.6	7
120	The Pseudomonas aeruginosa whole genome sequence: A 20th anniversary celebration. Advances in Microbial Physiology, 2021, 79, 25-88.	2.4	7
121	Unravelling postâ€transcriptional PrmCâ€dependent regulatory mechanisms in Pseudomonas aeruginosa. Environmental Microbiology, 2016, 18, 3583-3592.	3.8	6
122	Host-induced spermidine production in motile Pseudomonas aeruginosa triggers phagocytic uptake. ELife, 2020, 9, .	6.0	6
123	Pseudomonas aeruginosa postâ€translational responses to elevated <scp>câ€diâ€GMP</scp> levels. Molecular Microbiology, 2022, 117, 1213-1226.	2.5	6
124	Organism-specific depletion of highly abundant RNA species from bacterial total RNA. Access Microbiology, 2020, 2, acmi000159.	0.5	5
125	High plasmidome diversity of extended-spectrum beta-lactam-resistant Escherichia coli isolates collected during one year in one community hospital. Genomics, 2022, 114, 110368.	2.9	5
126	Analysis of the organization and expression patterns of the convergent Pseudomonas aeruginosa lasR/rsaL gene pair uncovers mutual influence. Molecular Microbiology, 2021, 115, 643-657.	2.5	4

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127	Targeting Bacterial Gyrase with Cystobactamid, Fluoroquinolone, and Aminocoumarin Antibiotics Induces Distinct Molecular Signatures in Pseudomonas aeruginosa. MSystems, 2021, 6, e0061021.	3.8	3
128	Phenome-ing Microbes. Springer Protocols, 2015, , 83-96.	0.3	1
129	The Peptide Chain Release Factor Methyltransferase PrmC Influences the Pseudomonas aeruginosa PA14 Endo- and Exometabolome. Metabolites, 2020, 10, 417.	2.9	0
130	Quo vadis clinical diagnostic microbiology?. Clinical Microbiology and Infection, 2021, 27, 1562-1564.	6.0	0
131	Pseudomonas aeruginosa flagellin modulates the immune response in ex vivo lung tissue slices. , 2019, , .		0