

Christiane Wolz

List of Publications by Year in descending order

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121
papers

8,276
citations

31976

53
h-index

54911

84
g-index

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all docs

125
docs citations

125
times ranked

7523
citing authors

#	ARTICLE	IF	CITATIONS
1	Two-Component Systems of <i>S. aureus</i> : Signaling and Sensing Mechanisms. <i>Genes</i> , 2022, 13, 34.	2.4	29
2	Chemolysin of <i>Staphylococcus aureus</i> impairs thrombus formation. <i>Journal of Thrombosis and Haemostasis</i> , 2022, 20, 1464-1475.	3.8	5
3	The <i>Staphylococcus epidermidis</i> Transcriptional Profile During Carriage. <i>Frontiers in Microbiology</i> , 2022, 13, 896311.	3.5	5
4	Intracellular persistence of <i>Staphylococcus aureus</i> in endothelial cells is promoted by the absence of phenol-soluble modulins. <i>Virulence</i> , 2021, 12, 1186-1198.	4.4	17
5	The Role of hlb-Converting Bacteriophages in <i>Staphylococcus aureus</i> Host Adaptation. <i>Microbial Physiology</i> , 2021, 31, 109-122.	2.4	26
6	Modeling of stringent-response reflects nutrient stress induced growth impairment and essential amino acids in different <i>Staphylococcus aureus</i> mutants. <i>Scientific Reports</i> , 2021, 11, 9651.	3.3	1
7	Interaction between <i>Staphylococcus</i> Agr virulence and neutrophils regulates pathogen expansion in the skin. <i>Cell Host and Microbe</i> , 2021, 29, 930-940.e4.	11.0	18
8	Proteome Dynamics during Antibiotic Persistence and Resuscitation. <i>MSystems</i> , 2021, 6, e0054921.	3.8	4
9	Adaptation of <i>Staphylococcus aureus</i> to the Human Skin Environment Identified Using an ex vivo Tissue Model. <i>Frontiers in Microbiology</i> , 2021, 12, 728989.	3.5	11
10	Exotoxins from <i>Staphylococcus aureus</i> activate 5-lipoxygenase and induce leukotriene biosynthesis. <i>Cellular and Molecular Life Sciences</i> , 2020, 77, 3841-3858.	5.4	16
11	<i>Staphylococcus aureus</i> Skin Colonization Is Enhanced by the Interaction of Neutrophil Extracellular Traps with Keratinocytes. <i>Journal of Investigative Dermatology</i> , 2020, 140, 1054-1065.e4.	0.7	32
12	Small Alarmone Synthetases RelP and RelQ of <i>Staphylococcus aureus</i> Are Involved in Biofilm Formation and Maintenance Under Cell Wall Stress Conditions. <i>Frontiers in Microbiology</i> , 2020, 11, 575882.	3.5	10
13	Structural Basis for Regulation of the Opposing (p)ppGpp Synthetase and Hydrolase within the Stringent Response Orchestrator Rel. <i>Cell Reports</i> , 2020, 32, 108157.	6.4	39
14	The alarmone (p)ppGpp confers tolerance to oxidative stress during the stationary phase by maintenance of redox and iron homeostasis in <i>Staphylococcus aureus</i> . <i>Free Radical Biology and Medicine</i> , 2020, 161, 351-364.	2.9	27
15	Intracellular <i>Staphylococcus aureus</i> persists upon antibiotic exposure. <i>Nature Communications</i> , 2020, 11, 2200.	12.8	197
16	The 5' NAD Cap of RNAlII Modulates Toxin Production in <i>Staphylococcus aureus</i> Isolates. <i>Journal of Bacteriology</i> , 2020, 202, .	2.2	25
17	Inducible expression of (pp)pGpp synthetases in <i>Staphylococcus aureus</i> is associated with activation of stress response genes. <i>PLoS Genetics</i> , 2020, 16, e1009282.	3.5	23
18	Revisiting the regulation of the capsular polysaccharide biosynthesis gene cluster in <i>Staphylococcus aureus</i> . <i>Molecular Microbiology</i> , 2019, 112, 1083-1099.	2.5	17

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19	Function and regulation of <i>Staphylococcus aureus</i> wall teichoic acids and capsular polysaccharides. <i>International Journal of Medical Microbiology</i> , 2019, 309, 151333.	3.6	31
20	Oxidative stress drives the selection of quorum sensing mutants in the <i>Staphylococcus aureus</i> population. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 19145-19154.	7.1	28
21	Temperate Phages of <i>Staphylococcus aureus</i> . <i>Microbiology Spectrum</i> , 2019, 7, .	3.0	43
22	Temperate Phages of <i>Staphylococcus aureus</i> . , 2019, , 521-535.		2
23	Structural and mechanistic divergence of the small (p)ppGpp synthetases RelP and RelQ. <i>Scientific Reports</i> , 2018, 8, 2195.	3.3	51
24	Methicillin-resistant <i>Staphylococcus aureus</i> alters cell wall glycosylation to evade immunity. <i>Nature</i> , 2018, 563, 705-709.	27.8	137
25	Long Noncoding RNA SSR42 Controls <i>Staphylococcus aureus</i> Alpha-Toxin Transcription in Response to Environmental Stimuli. <i>Journal of Bacteriology</i> , 2018, 200, .	2.2	15
26	Regulation of the opposing (p)ppGpp synthetase and hydrolase activities in a bifunctional RelA/SpoT homologue from <i>Staphylococcus aureus</i> . <i>PLoS Genetics</i> , 2018, 14, e1007514.	3.5	67
27	Inactivation of TCA cycle enhances <i>Staphylococcus aureus</i> persister cell formation in stationary phase. <i>Scientific Reports</i> , 2018, 8, 10849.	3.3	68
28	European external quality assessments for identification, molecular typing and characterization of <i>Staphylococcus aureus</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2018, 73, 2662-2666.	3.0	6
29	Wall teichoic acids mediate increased virulence in <i>Staphylococcus aureus</i> . <i>Nature Microbiology</i> , 2017, 2, 16257.	13.3	81
30	Human NACHT, LRR, and PYD domain-containing protein 3 (NLRP3) inflammasome activity is regulated by and potentially targetable through Bruton tyrosine kinase. <i>Journal of Allergy and Clinical Immunology</i> , 2017, 140, 1054-1067.e10.	2.9	105
31	Absence of ppGpp Leads to Increased Mobilization of Intermediately Accumulated Poly(3-Hydroxybutyrate) in <i>Ralstonia eutropha</i> H16. <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	33
32	Downstream element determines RNase Y cleavage of the saePQRS operon in <i>Staphylococcus aureus</i> . <i>Nucleic Acids Research</i> , 2017, 45, 5980-5994.	14.5	21
33	Keratinocytes as sensors and central players in the immune defense against <i>Staphylococcus aureus</i> in the skin. <i>Journal of Dermatological Science</i> , 2017, 87, 215-220.	1.9	65
34	Commercial Biocides Induce Transfer of Prophage ϕ 13 from Human Strains of <i>Staphylococcus aureus</i> to Livestock CC398. <i>Frontiers in Microbiology</i> , 2017, 8, 2418.	3.5	23
35	An essential role for the baseplate protein Gp45 in phage adsorption to <i>Staphylococcus aureus</i> . <i>Scientific Reports</i> , 2016, 6, 26455.	3.3	61
36	RpiRc Is a Pleiotropic Effector of Virulence Determinant Synthesis and Attenuates Pathogenicity in <i>Staphylococcus aureus</i> . <i>Infection and Immunity</i> , 2016, 84, 2031-2041.	2.2	26

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37	Toll-like receptor 2 activation depends on lipopeptide shedding by bacterial surfactants. <i>Nature Communications</i> , 2016, 7, 12304.	12.8	86
38	Influence of Sae-regulated and Agr-regulated factors on the escape of <i>Staphylococcus aureus</i> from human macrophages. <i>Cellular Microbiology</i> , 2016, 18, 1172-1183.	2.1	67
39	rRNA regulation during growth and under stringent conditions in <i>Staphylococcus aureus</i> . <i>Environmental Microbiology</i> , 2015, 17, 4394-4405.	3.8	30
40	Phenotypic heterogeneity and temporal expression of the capsular polysaccharide in <i>Staphylococcus aureus</i> . <i>Molecular Microbiology</i> , 2015, 98, 1073-1088.	2.5	27
41	Fine-tuning recA expression in <i>Staphylococcus aureus</i> for antimicrobial photoinactivation: importance of photo-induced DNA damage in the photoinactivation mechanism. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 9161-9176.	3.6	46
42	Emergence of trimethoprim resistance gene dfrG in <i>Staphylococcus aureus</i> causing human infection and colonization in sub-Saharan Africa and its import to Europe. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 2361-2368.	3.0	87
43	Cytoplasmic replication of <i>Staphylococcus aureus</i> upon phagosomal escape triggered by phenol-soluble modulins. <i>Cellular Microbiology</i> , 2014, 16, 451-465.	2.1	160
44	The Catabolite Control Protein E (CcpE) Affects Virulence Determinant Production and Pathogenesis of <i>Staphylococcus aureus</i> . <i>Journal of Biological Chemistry</i> , 2014, 289, 29701-29711.	3.4	27
45	Intersection of the stringent response and the CodY regulon in low GC Gram-positive bacteria. <i>International Journal of Medical Microbiology</i> , 2014, 304, 150-155.	3.6	103
46	Phages of <i>Staphylococcus aureus</i> and their impact on host evolution. <i>Infection, Genetics and Evolution</i> , 2014, 21, 593-601.	2.3	184
47	Two Small (p)ppGpp Synthases in <i>Staphylococcus aureus</i> Mediate Tolerance against Cell Envelope Stress Conditions. <i>Journal of Bacteriology</i> , 2014, 196, 894-902.	2.2	159
48	Altering gene expression by aminocoumarins: the role of DNA supercoiling in <i>Staphylococcus aureus</i> . <i>BMC Genomics</i> , 2014, 15, 291.	2.8	22
49	Heterogeneity of Host TLR2 Stimulation by <i>Staphylococcus aureus</i> Isolates. <i>PLoS ONE</i> , 2014, 9, e96416.	2.5	25
50	Production of capsular polysaccharide does not influence <i>Staphylococcus aureus</i> vancomycin susceptibility. <i>BMC Microbiology</i> , 2013, 13, 65.	3.3	31
51	A semi-quantitative model of Quorum-Sensing in <i>Staphylococcus aureus</i> , approved by microarray meta-analyses and tested by mutation studies. <i>Molecular BioSystems</i> , 2013, 9, 2665.	2.9	16
52	Opposing effects of aminocoumarins and fluoroquinolones on the SOS response and adaptability in <i>Staphylococcus aureus</i> . <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 529-538.	3.0	48
53	Methionine Biosynthesis in <i>Staphylococcus aureus</i> Is Tightly Controlled by a Hierarchical Network Involving an Initiator tRNA-Specific T-box Riboswitch. <i>PLoS Pathogens</i> , 2013, 9, e1003606.	4.7	23
54	Diversification of Clonal Complex 5 Methicillin-Resistant <i>Staphylococcus aureus</i> Strains (Rhine-Hesse) Tj ETQq0 0 0,rgBT /Overlock 10 Tf	3.9	23

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55	SDS Interferes with SaeS Signaling of <i>Staphylococcus aureus</i> Independently of SaePQ. <i>PLoS ONE</i> , 2013, 8, e71644.	2.5	9
56	Two Distinct Coagulase-Dependent Barriers Protect <i>Staphylococcus aureus</i> from Neutrophils in a Three Dimensional in vitro Infection Model. <i>PLoS Pathogens</i> , 2012, 8, e1002434.	4.7	77
57	The Stringent Response of <i>Staphylococcus aureus</i> and Its Impact on Survival after Phagocytosis through the Induction of Intracellular PSMs Expression. <i>PLoS Pathogens</i> , 2012, 8, e1003016.	4.7	209
58	Import and Spread of Panton-Valentine Leukocidin in <i>Staphylococcus aureus</i> Through Nasal Carriage and Skin Infections in Travelers Returning From the Tropics and Subtropics. <i>Clinical Infectious Diseases</i> , 2012, 54, 483-492.	5.8	78
59	Global Analysis of the <i>Staphylococcus aureus</i> Response to Mupirocin. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 787-804.	3.2	88
60	Reply to Mimica. <i>Clinical Infectious Diseases</i> , 2012, 54, 1518-1519.	5.8	0
61	<i>Staphylococcus aureus</i> determinants for nasal colonization. <i>Trends in Microbiology</i> , 2012, 20, 243-250.	7.7	127
62	RNase Y of <i>Staphylococcus aureus</i> and its role in the activation of virulence genes. <i>Molecular Microbiology</i> , 2012, 85, 817-832.	2.5	72
63	Expression of staphylococcal superantigens during nasal colonization is not sufficient to induce a systemic neutralizing antibody response in humans. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2012, 31, 251-256.	2.9	24
64	The CodY pleiotropic repressor controls virulence in gram-positive pathogens. <i>FEMS Immunology and Medical Microbiology</i> , 2011, 62, 123-139.	2.7	94
65	<i>Staphylococcus aureus</i> physiological growth limitations: Insights from flux calculations built on proteomics and external metabolite data. <i>Proteomics</i> , 2011, 11, 1915-1935.	2.2	27
66	β -Lactams Interfering with PBP1 Induce Panton-Valentine Leukocidin Expression by Triggering <i>sarA</i> and <i>rot</i> Global Regulators of <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 3261-3271.	3.2	61
67	Vectors for improved Tet repressor-dependent gradual gene induction or silencing in <i>Staphylococcus aureus</i> . <i>Microbiology (United Kingdom)</i> , 2011, 157, 3314-3323.	1.8	87
68	High-Level Fluorescence Labeling of Gram-Positive Pathogens. <i>PLoS ONE</i> , 2011, 6, e19822.	2.5	43
69	Insertion of host DNA into PVL-encoding phages of the <i>Staphylococcus aureus</i> lineage ST80 by intra-chromosomal recombination. <i>Virology</i> , 2010, 406, 322-327.	2.4	14
70	Changing the phospholipid composition of <i>Staphylococcus aureus</i> causes distinct changes in membrane proteome and membrane sensory regulators. <i>Proteomics</i> , 2010, 10, 1685-1693.	2.2	27
71	Staphylococcal superantigen-like genes, <i>ssl5</i> and <i>ssl8</i> , are positively regulated by Sae and negatively by Agr in the Newman strain. <i>FEMS Microbiology Letters</i> , 2010, 308, no-no.	1.8	29
72	Differential Target Gene Activation by the <i>Staphylococcus aureus</i> Two-Component System <i>saeRS</i> . <i>Journal of Bacteriology</i> , 2010, 192, 613-623.	2.2	150

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73	Temporal Expression of Adhesion Factors and Activity of Global Regulators during Establishment of <i>Staphylococcus aureus</i> Nasal Colonization. <i>Journal of Infectious Diseases</i> , 2010, 201, 1414-1421.	4.0	114
74	Role of the (p)ppGpp Synthase RSH, a RelA/SpoT Homolog, in Stringent Response and Virulence of <i>Staphylococcus aureus</i> . <i>Infection and Immunity</i> , 2010, 78, 1873-1883.	2.2	123
75	The synthesis and function of the alarmone (p)ppGpp in firmicutes. <i>International Journal of Medical Microbiology</i> , 2010, 300, 142-147.	3.6	57
76	Adaptation of <i>Staphylococcus aureus</i> to the cystic fibrosis lung. <i>International Journal of Medical Microbiology</i> , 2010, 300, 520-525.	3.6	108
77	Regulatory Adaptation of <i>Staphylococcus aureus</i> during Nasal Colonization of Humans. <i>PLoS ONE</i> , 2010, 5, e10040.	2.5	101
78	Diversity of Prophages in Dominant <i>Staphylococcus aureus</i> Clonal Lineages. <i>Journal of Bacteriology</i> , 2009, 191, 3462-3468.	2.2	257
79	Transcriptional regulation of the novobiocin biosynthetic gene cluster. <i>Microbiology (United Kingdom)</i> 151:1077-1087	1.8	15
80	Functional Characterization of the σ^B -Dependent <i>yabJ</i> - <i>spoVG</i> Operon in <i>Staphylococcus aureus</i> : Role in Methicillin and Glycopeptide Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 1832-1839.	3.2	70
81	A Point Mutation in the Sensor Histidine Kinase SaeS of <i>Staphylococcus aureus</i> Strain Newman Alters the Response to Biocide Exposure. <i>Journal of Bacteriology</i> , 2009, 191, 7306-7314.	2.2	40
82	Transcription Analysis of the Extracellular Adherence Protein from <i>Staphylococcus aureus</i> in Authentic Human Infection and In Vitro. <i>Journal of Infectious Diseases</i> , 2009, 199, 1471-1478.	4.0	40
83	CodY in <i>Staphylococcus aureus</i> : a Regulatory Link between Metabolism and Virulence Gene Expression. <i>Journal of Bacteriology</i> , 2009, 191, 2953-2963.	2.2	195
84	Transcription of the phage-encoded Pantonâ€“Valentine leukocidin of <i>Staphylococcus aureus</i> is dependent on the phage life-cycle and on the host background. <i>Microbiology (United Kingdom)</i> , 2009, 155, 3491-3499.	1.8	51
85	Dermcidin-Derived Peptides Show a Different Mode of Action than the Cathelicidin LL-37 against <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2499-2509.	3.2	61
86	Acquisition of antibiotic-resistant <i>Enterococcus faecium</i> strains during long-term hospitalization and fast adaptation of enterococcal flora to antibiotic treatment: A case report. <i>International Journal of Hygiene and Environmental Health</i> , 2009, 212, 105-108.	4.3	1
87	Resistance to dermcidin-derived peptides is independent of bacterial protease activity. <i>International Journal of Antimicrobial Agents</i> , 2009, 34, 86-90.	2.5	10
88	Bioluminescence imaging to study the promoter activity of <i>hla</i> of <i>Staphylococcus aureus</i> in vitro and in vivo. <i>International Journal of Medical Microbiology</i> , 2008, 298, 599-605.	3.6	11
89	<i>Staphylococcus aureus</i> CcpA Affects Biofilm Formation. <i>Infection and Immunity</i> , 2008, 76, 2044-2050.	2.2	153
90	The Virulence Regulator Sae of <i>Staphylococcus aureus</i> : Promoter Activities and Response to Phagocytosis-Related Signals. <i>Journal of Bacteriology</i> , 2008, 190, 3419-3428.	2.2	166

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91	Emergence of increasing linezolid-resistance in enterococci in a post-outbreak situation with vancomycin-resistant <i>Enterococcus faecium</i> . <i>Epidemiology and Infection</i> , 2008, 136, 1131-1133.	2.1	33
92	Ĵf ^B and the Ĵf ^B-Dependent <i>arlRS</i> and <i>yabJ-spoVG</i> Loci Affect Capsule Formation in <i>Staphylococcus aureus</i>. <i>Infection and Immunity</i> , 2007, 75, 4562-4571.	2.2	72
93	High phenotypic diversity in infecting but not in colonizing <i>Staphylococcus aureus</i> populations. <i>Environmental Microbiology</i> , 2007, 9, 3134-3142.	3.8	49
94	The staphylococcal respiratory response regulator SrrAB induces <i>ica</i> gene transcription and polysaccharide intercellular adhesin expression, protecting <i>Staphylococcus aureus</i> from neutrophil killing under anaerobic growth conditions. <i>Molecular Microbiology</i> , 2007, 65, 1276-1287.	2.5	94
95	The staphylococcal respiratory response regulator SrrAB induces <i>ica</i> gene transcription and polysaccharide intercellular adhesin expression, protecting <i>Staphylococcus aureus</i> from neutrophil killing under anaerobic growth conditions. <i>Molecular Microbiology</i> , 2007, 66, 278-278.	2.5	2
96	Discrimination between epidemic and non-epidemic glycopeptide-resistant <i>E. faecium</i> in a post-outbreak situation. <i>Journal of Hospital Infection</i> , 2007, 67, 49-55.	2.9	29
97	Suicidal erythrocyte death in sepsis. <i>Journal of Molecular Medicine</i> , 2007, 85, 273-281.	3.9	277
98	Extensive phage dynamics in <i>Staphylococcus aureus</i> contributes to adaptation to the human host during infection. <i>Molecular Microbiology</i> , 2006, 61, 1673-1685.	2.5	136
99	<i>Staphylococcus aureus</i> CcpA Affects Virulence Determinant Production and Antibiotic Resistance. <i>Antimicrobial Agents and Chemotherapy</i> , 2006, 50, 1183-1194.	3.2	179
100	Ciprofloxacin and Trimethoprim Cause Phage Induction and Virulence Modulation in <i>Staphylococcus aureus</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2006, 50, 171-177.	3.2	190
101	Influence of the Two-Component System SaeRS on Global Gene Expression in Two Different <i>Staphylococcus aureus</i> Strains. <i>Journal of Bacteriology</i> , 2006, 188, 7742-7758.	2.2	164
102	Influence of clindamycin on the stability of <i>coa</i> and <i>fnbB</i> transcripts and adherence properties of <i>Staphylococcus aureus</i> Newman. <i>FEMS Microbiology Letters</i> , 2005, 252, 73-78.	1.8	16
103	<i>sae</i> is essential for expression of the staphylococcal adhesins Eap and Emp. <i>Microbiology (United Kingdom)</i> 157:1073-1081 (2005)	1.8	69
104	Biofilm Formation, <i>icaADBC</i> Transcription, and Polysaccharide Intercellular Adhesin Synthesis by <i>Staphylococci</i> in a Device-Related Infection Model. <i>Infection and Immunity</i> , 2005, 73, 1811-1819.	2.2	157
105	Molecular basis of florfenicol-induced increase in adherence of <i>Staphylococcus aureus</i> strain Newman. <i>Journal of Antimicrobial Chemotherapy</i> , 2005, 56, 315-323.	3.0	25
106	Role of <i>Staphylococcus aureus</i> Global Regulators <i>sae</i> and ĴfB in Virulence Gene Expression during Device-Related Infection. <i>Infection and Immunity</i> , 2005, 73, 3415-3421.	2.2	111
107	strain designation by and polymorphism typing and delineation of diversification by sequence analysis. <i>International Journal of Medical Microbiology</i> , 2005, 295, 67-75.	3.6	58
108	Increased Frequency of Genomic Alterations in <i>Staphylococcus aureus</i> during Chronic Infection Is in Part Due to Phage Mobilization. <i>Journal of Infectious Diseases</i> , 2004, 189, 724-734.	4.0	99

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109	Regulatory and genomic plasticity of <i>Staphylococcus aureus</i> during persistent colonization and infection. <i>International Journal of Medical Microbiology</i> , 2004, 294, 195-202.	3.6	80
110	Evaluation of Intraspecies Interference Due to agr Polymorphism in <i>Staphylococcus aureus</i> during Infection and Colonization. <i>Journal of Infectious Diseases</i> , 2003, 188, 250-256.	4.0	39
111	Molecular Architecture of the Regulatory Locus sae of <i>Staphylococcus aureus</i> and Its Impact on Expression of Virulence Factors. <i>Journal of Bacteriology</i> , 2003, 185, 6278-6286.	2.2	120
112	Transcription of Clumping Factor A in Attached and Unattached <i>Staphylococcus aureus</i> In Vitro and during Device-Related Infection. <i>Infection and Immunity</i> , 2002, 70, 2758-2762.	2.2	76
113	Quantification of Bacterial Transcripts during Infection Using Competitive Reverse Transcription-PCR (RT-PCR) and LightCycler RT-PCR. <i>Vaccine Journal</i> , 2001, 8, 279-282.	2.6	50
114	Impact of the regulatory loci agr, sarA and sae of <i>Staphylococcus aureus</i> on the induction of alpha-toxin during device-related infection resolved by direct quantitative transcript analysis. <i>Molecular Microbiology</i> , 2001, 40, 1439-1447.	2.5	155
115	Regulation of <i>Staphylococcus aureus</i> Type 5 and Type 8 Capsular Polysaccharides by CO ₂ . <i>Journal of Bacteriology</i> , 2001, 183, 4609-4613.	2.2	35
116	Agr-independent regulation of fibronectin-binding protein(s) by the regulatory locus sar in <i>Staphylococcus aureus</i> . <i>Molecular Microbiology</i> , 2000, 36, 230-243.	2.5	123
117	Direct Quantitative Transcript Analysis of the agr Regulon of <i>Staphylococcus aureus</i> during Human Infection in Comparison to the Expression Profile In Vitro. <i>Infection and Immunity</i> , 2000, 68, 1304-1311.	2.2	207
118	Molecular Epidemiology of Community-Acquired <i>Staphylococcus aureus</i> in Families with and without Cystic Fibrosis Patients. <i>Journal of Infectious Diseases</i> , 2000, 181, 984-989.	4.0	57
119	Adherence of <i>Staphylococcus aureus</i> to Endothelial Cells: Influence of Capsular Polysaccharide, Global Regulator agr, and Bacterial Growth Phase. <i>Infection and Immunity</i> , 2000, 68, 4865-4871.	2.2	117
120	Characterization of a sar Homolog of <i>Staphylococcus epidermidis</i> . <i>Infection and Immunity</i> , 1998, 66, 2871-2878.	2.2	30
121	<i>Pseudomonas aeruginosa</i> cross-colonization and persistence in patients with cystic fibrosis. Use of a DNA probe. <i>Epidemiology and Infection</i> , 1989, 102, 205-214.	2.1	98