

Thorsten Mascher

List of Publications by Year in descending order

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82
papers

5,585
citations

94433

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85541

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90
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90
docs citations

90
times ranked

4468
citing authors

#	ARTICLE	IF	CITATIONS
1	Stimulus Perception in Bacterial Signal-Transducing Histidine Kinases. <i>Microbiology and Molecular Biology Reviews</i> , 2006, 70, 910-938.	6.6	592
2	The third pillar of bacterial signal transduction: classification of the extracytoplasmic function (ECF) σ factor protein family. <i>Molecular Microbiology</i> , 2009, 74, 557-581.	2.5	374
3	Cell envelope stress response in Gram-positive bacteria. <i>FEMS Microbiology Reviews</i> , 2008, 32, 107-146.	8.6	323
4	Cell wall stress responses in <i>Bacillus subtilis</i> : the regulatory network of the bacitracin stimulon. <i>Molecular Microbiology</i> , 2003, 50, 1591-1604.	2.5	290
5	Antibiotic-Inducible Promoter Regulated by the Cell Envelope Stress-Sensing Two-Component System LiaRS of <i>Bacillus subtilis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2004, 48, 2888-2896.	3.2	277
6	The Bacillus BioBrick Box: generation and evaluation of essential genetic building blocks for standardized work with <i>Bacillus subtilis</i> . <i>Journal of Biological Engineering</i> , 2013, 7, 29.	4.7	195
7	Regulation of LiaRS-Dependent Gene Expression in <i>Bacillus subtilis</i> : Identification of Inhibitor Proteins, Regulator Binding Sites, and Target Genes of a Conserved Cell Envelope Stress-Sensing Two-Component System. <i>Journal of Bacteriology</i> , 2006, 188, 5153-5166.	2.2	189
8	Cultivation and functional characterization of 79 planctomycetes uncovers their unique biology. <i>Nature Microbiology</i> , 2020, 5, 126-140.	13.3	164
9	Bacitracin sensing in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2008, 68, 768-785.	2.5	162
10	Intramembrane-sensing histidine kinases: a new family of cell envelope stress sensors in Firmicutes bacteria. <i>FEMS Microbiology Letters</i> , 2006, 264, 133-144.	1.8	143
11	The <i>Streptococcus pneumoniae</i> σ Regulon: CiaR Target Sites and Transcription Profile Analysis. <i>Journal of Bacteriology</i> , 2003, 185, 60-70.	2.2	142
12	Signaling diversity and evolution of extracytoplasmic function (ECF) σ factors. <i>Current Opinion in Microbiology</i> , 2013, 16, 148-155.	5.1	138
13	Coevolution of ABC Transporters and Two-Component Regulatory Systems as Resistance Modules against Antimicrobial Peptides in Firmicutes Bacteria. <i>Journal of Bacteriology</i> , 2011, 193, 3851-3862.	2.2	135
14	Regulatory Overlap and Functional Redundancy among <i>Bacillus subtilis</i> Extracytoplasmic Function σ Factors. <i>Journal of Bacteriology</i> , 2007, 189, 6919-6927.	2.2	112
15	Identification of Proteins Likely To Be Involved in Morphogenesis, Cell Division, and Signal Transduction in Planctomycetes by Comparative Genomics. <i>Journal of Bacteriology</i> , 2012, 194, 6419-6430.	2.2	110
16	In-Depth Profiling of the LiaR Response of <i>Bacillus subtilis</i> . <i>Journal of Bacteriology</i> , 2010, 192, 4680-4693.	2.2	107
17	Peptide Antibiotic Sensing and Detoxification Modules of <i>Bacillus subtilis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 515-525.	3.2	105
18	The CiaRH System of <i>Streptococcus pneumoniae</i> Prevents Lysis during Stress Induced by Treatment with Cell Wall Inhibitors and by Mutations in <i>pbp2x</i> Involved in β -Lactam Resistance. <i>Journal of Bacteriology</i> , 2006, 188, 1959-1968.	2.2	99

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19	Daptomycin versus Friulimicin B: In-Depth Profiling of <i>Bacillus subtilis</i> Cell Envelope Stress Responses. <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 1619-1623.	3.2	87
20	The Bacillus BioBrick Box 2.0: expanding the genetic toolbox for the standardized work with <i>Bacillus subtilis</i> . <i>Scientific Reports</i> , 2017, 7, 15058.	3.3	82
21	Anatomy of the bacitracin resistance network in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2016, 100, 607-620.	2.5	67
22	Antibiotic research in the age of omics: from expression profiles to interspecies communication. <i>Journal of Antimicrobial Chemotherapy</i> , 2011, 66, 2689-2704.	3.0	60
23	A New Way of Sensing: Need-Based Activation of Antibiotic Resistance by a Flux-Sensing Mechanism. <i>MBio</i> , 2015, 6, e00975.	4.1	60
24	Bacterial (intramembrane-sensing) histidine kinases: signal transfer rather than stimulus perception. <i>Trends in Microbiology</i> , 2014, 22, 559-565.	7.7	59
25	The cell envelope stress response of <i>Bacillus subtilis</i> : from static signaling devices to dynamic regulatory network. <i>Current Genetics</i> , 2017, 63, 79-90.	1.7	58
26	Defence against antimicrobial peptides: different strategies in <i>Bifidobacterium</i> . <i>Environmental Microbiology</i> , 2014, 16, 1225-1237.	3.8	54
27	Environmental Sensing in Actinobacteria: a Comprehensive Survey on the Signaling Capacity of This Phylum. <i>Journal of Bacteriology</i> , 2015, 197, 2517-2535.	2.2	54
28	The LIKE system, a novel protein expression toolbox for <i>Bacillus subtilis</i> based on the <i>lial</i> promoter. <i>Microbial Cell Factories</i> , 2012, 11, 143.	4.0	53
29	The Pneumococcal Cell Envelope Stress-Sensing System LiaFSR Is Activated by Murein Hydrolases and Lipid II-Interacting Antibiotics. <i>Journal of Bacteriology</i> , 2010, 192, 1761-1773.	2.2	50
30	General stress response in α -proteobacteria: PhyR and beyond. <i>Molecular Microbiology</i> , 2010, 78, 271-277.	2.5	48
31	Antimicrobial peptide sensing and detoxification modules: unravelling the regulatory circuitry of <i>Staphylococcus aureus</i> . <i>Molecular Microbiology</i> , 2011, 81, 581-587.	2.5	48
32	Stoichiometry and perturbation studies of the LiaFSR system of <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2013, 87, 769-788.	2.5	47
33	Subcellular localization, interactions and dynamics of the phage shock protein-like Lia response in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2014, 92, 716-732.	2.5	45
34	Cell Envelope Stress Response in Cell Wall-Deficient L-Forms of <i>Bacillus subtilis</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 5907-5915.	3.2	44
35	Substitution of the native <i>srfA</i> promoter by constitutive P in two <i>B. subtilis</i> strains and evaluation of the effect on Surfactin production. <i>Journal of Biotechnology</i> , 2016, 224, 14-17.	3.8	44
36	Characterization of a Regulatory Network of Peptide Antibiotic Detoxification Modules in <i>Lactobacillus casei</i> BL23. <i>Applied and Environmental Microbiology</i> , 2013, 79, 3160-3170.	3.1	41

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37	LiaRS-dependent gene expression is embedded in transition state regulation in <i>Bacillus subtilis</i> . <i>Microbiology (United Kingdom)</i> , 2007, 153, 2530-2540.	1.8	40
38	Extracytoplasmic function σ factors of the widely distributed group ECF41 contain a fused regulatory domain. <i>MicrobiologyOpen</i> , 2012, 1, 194-213.	3.0	40
39	A dynamin-like protein involved in bacterial cell membrane surveillance under environmental stress. <i>Environmental Microbiology</i> , 2016, 18, 2705-2720.	3.8	40
40	Regulation of the <i>Bacillus subtilis</i> Extracytoplasmic Function Protein σ Y and Its Target Promoters. <i>Journal of Bacteriology</i> , 2003, 185, 4883-4890.	2.2	37
41	The applied side of antimicrobial peptide-inducible promoters from Firmicutes bacteria: expression systems and whole-cell biosensors. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 4817-4829.	3.6	34
42	Cannibalism stress response in <i>Bacillus subtilis</i> . <i>Microbiology (United Kingdom)</i> , 2016, 162, 164-176.	1.8	34
43	Cell Envelope Stress Response in <i>Bacillus licheniformis</i> : Integrating Comparative Genomics, Transcriptional Profiling, and Regulon Mining To Decipher a Complex Regulatory Network. <i>Journal of Bacteriology</i> , 2006, 188, 7500-7511.	2.2	33
44	The peroxide stress response of <i>Bacillus licheniformis</i> . <i>Proteomics</i> , 2011, 11, 2851-2866.	2.2	32
45	The <i>Bacillus subtilis</i> GntR Family Repressor YtrA Responds to Cell Wall Antibiotics. <i>Journal of Bacteriology</i> , 2011, 193, 5793-5801.	2.2	32
46	Engineering orthogonal synthetic timer circuits based on extracytoplasmic function σ factors. <i>Nucleic Acids Research</i> , 2018, 46, 7450-7464.	14.5	32
47	Expansion and re-classification of the extracytoplasmic function (ECF) σ factor family. <i>Nucleic Acids Research</i> , 2021, 49, 986-1005.	14.5	32
48	The <i>ciaR/ciaH</i> regulatory network of <i>Streptococcus pneumoniae</i> . <i>Journal of Molecular Microbiology and Biotechnology</i> , 2002, 4, 211-6.	1.0	31
49	Coordinated Cell Death in Isogenic Bacterial Populations: Sacrificing Some for the Benefit of Many?. <i>Journal of Molecular Biology</i> , 2019, 431, 4656-4669.	4.2	30
50	The Epeptide YydF Intrinsically Triggers the Cell Envelope Stress Response of <i>Bacillus subtilis</i> and Causes Severe Membrane Perturbations. <i>Frontiers in Microbiology</i> , 2020, 11, 151.	3.5	29
51	(Actino)Bacterial "intelligence" using comparative genomics to unravel the information processing capacities of microbes. <i>Current Genetics</i> , 2016, 62, 487-498.	1.7	27
52	Defining the regulon of genes controlled by σ^E , a key regulator of the cell envelope stress response in <i>Streptomyces coelicolor</i> . <i>Molecular Microbiology</i> , 2019, 112, 461-481.	2.5	27
53	Membrane chaperoning by members of the PspA/IM30 protein family. <i>Communicative and Integrative Biology</i> , 2017, 10, e1264546.	1.4	25
54	ECF σ factors with regulatory extensions: the one-component systems of the σ universe. <i>Molecular Microbiology</i> , 2019, 112, 399-409.	2.5	23

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55	Bacillus SEVA siblings: A Golden Gate-based toolbox to create personalized integrative vectors for <i>Bacillus subtilis</i> . <i>Scientific Reports</i> , 2017, 7, 14134.	3.3	22
56	Immediate and Heterogeneous Response of the LiaFSR Two-Component System of <i>Bacillus subtilis</i> to the Peptide Antibiotic Bacitracin. <i>PLoS ONE</i> , 2013, 8, e53457.	2.5	20
57	Sporobeads: The Utilization of the <i>Bacillus subtilis</i> Endospore Crust as a Protein Display Platform. <i>ACS Synthetic Biology</i> , 2018, 7, 452-461.	3.8	20
58	The <i>Bacillus subtilis</i> endospore crust: protein interaction network, architecture and glycosylation state of a potential glycoprotein layer. <i>Molecular Microbiology</i> , 2019, 112, 1576-1592.	2.5	19
59	The role of C-terminal extensions in controlling ECF σ factor activity in the widely conserved groups ECF41 and ECF42. <i>Molecular Microbiology</i> , 2019, 112, 498-514.	2.5	19
60	Application of a <i>Bacillus subtilis</i> Whole-Cell Biosensor (Pial-lux) for the Identification of Cell Wall Active Antibacterial Compounds. <i>Methods in Molecular Biology</i> , 2017, 1520, 121-131.	0.9	17
61	Environmental Sensing and the Role of Extracytoplasmic Function Sigma Factors. , 2008, , 233-261.		16
62	The Essential UPP Phosphatase Pair BcrC and UppP Connects Cell Wall Homeostasis during Growth and Sporulation with Cell Envelope Stress Response in <i>Bacillus subtilis</i> . <i>Frontiers in Microbiology</i> , 2017, 8, 2403.	3.5	16
63	Characterization of the Widely Distributed Novel ECF42 Group of Extracytoplasmic Function σ Factors in <i>Streptomyces venezuelae</i> . <i>Journal of Bacteriology</i> , 2018, 200, .	2.2	16
64	The rhamnolipid stress response of <i>Bacillus subtilis</i> . <i>FEMS Microbiology Letters</i> , 2011, 323, 113-123.	1.8	15
65	The three-component system EsiSR regulates a cell envelope stress response in <i>Corynebacterium glutamicum</i> . <i>Molecular Microbiology</i> , 2017, 106, 719-741.	2.5	15
66	Amphotericin B Specifically Induces the Two-Component System LnrJK: Development of a Novel Whole-Cell Biosensor for the Detection of Amphotericin-Like Polyenes. <i>Frontiers in Microbiology</i> , 2020, 11, 2022.	3.5	13
67	The Epipeptide Biosynthesis Locus <i>epsXEPAB</i> Is Widely Distributed in <i>Firmicutes</i> and Triggers Intrinsic Cell Envelope Stress. <i>Microbial Physiology</i> , 2021, 31, 306-318.	2.4	13
68	Development of a novel heterologous β -lactam-specific whole-cell biosensor in <i>Bacillus subtilis</i> . <i>Journal of Biological Engineering</i> , 2020, 14, 21.	4.7	12
69	Phyletic Distribution and Diversification of the Phage Shock Protein Stress Response System in Bacteria and Archaea. <i>MSystems</i> , 2022, 7, .	3.8	11
70	A balancing act times two: sensing and regulating cell envelope homeostasis in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2014, 94, 1201-1207.	2.5	10
71	<i>Bacillus subtilis</i> as a Platform for Molecular Characterisation of Regulatory Mechanisms of <i>Enterococcus faecalis</i> Resistance against Cell Wall Antibiotics. <i>PLoS ONE</i> , 2014, 9, e93169.	2.5	9
72	Silver (I) N-Heterocyclic Carbenes Carbosilane Dendritic Systems and Their Imidazolium-Terminated Analogues as Antibacterial Agents: Study of Their Mode of Action. <i>Pharmaceutics</i> , 2020, 12, 968.	4.5	9

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73	Extracytoplasmic Function ħ Factors Can Be Implemented as Robust Heterologous Genetic Switches in <i>Bacillus subtilis</i> . <i>IScience</i> , 2019, 13, 380-390.	4.1	8
74	From Modules to Networks: a Systems-Level Analysis of the Bacitracin Stress Response in <i>Bacillus subtilis</i> . <i>MSystems</i> , 2020, 5, .	3.8	8
75	Insulation and wiring specificity of BceR-like response regulators and their target promoters in <i>Bacillus subtilis</i> . <i>Molecular Microbiology</i> , 2017, 104, 16-31.	2.5	7
76	Extracytoplasmic Function ħ Factors Come of Age. <i>Microbe Magazine</i> , 2010, 5, 164-170.	0.4	7
77	Regulation of heterologous subtilin production in <i>Bacillus subtilis</i> W168. <i>Microbial Cell Factories</i> , 2022, 21, 57.	4.0	7
78	Regulatory Characteristics of <i>Bacillus pumilus</i> Protease Promoters. <i>Current Microbiology</i> , 2017, 74, 550-559.	2.2	4
79	Promoter RNA sequencing (PRSeq) for the massive and quantitative promoter analysis in vitro. <i>Scientific Reports</i> , 2019, 9, 3118.	3.3	3
80	ABC Transporter DerAB of <i>Lactobacillus casei</i> Mediates Resistance against Insect-Derived Defensins. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	3
81	Low phosphatase activity of LiaS and strong LiaR-DNA affinity explain the unusual LiaS to LiaR in vivo stoichiometry. <i>BMC Microbiology</i> , 2020, 20, 104.	3.3	3
82	The REACT Suite: A Software Toolkit for Microbial REgulon Annotation and Comparative Transcriptomics. , 0, , .		0