

# Regine Hengge

## List of Publications by Year in descending order

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73  
papers

7,880  
citations

71102

41  
h-index

82547

72  
g-index

81  
all docs

81  
docs citations

81  
times ranked

7064  
citing authors

#	ARTICLE	IF	CITATIONS
1	High-specificity local and global c-di-GMP signaling. Trends in Microbiology, 2021, 29, 993-1003.	7.7	74
2	Bacterial Multicellularity: The Biology of <i>Escherichia coli</i> Building Large-Scale Biofilm Communities. Annual Review of Microbiology, 2021, 75, 269-290.	7.3	36
3	Adaptation of <i>Escherichia coli</i> Biofilm Growth, Morphology, and Mechanical Properties to Substrate Water Content. ACS Biomaterials Science and Engineering, 2021, 7, 5315-5325.	5.2	14
4	Crosstalking second messengers. Nature Microbiology, 2021, 6, 9-10.	13.3	3
5	A Novel Locally c-di-GMP-Controlled Exopolysaccharide Synthase Required for Bacteriophage N4 Infection of <i>Escherichia coli</i> . MBio, 2021, 12, e0324921.	4.1	14
6	Common plant flavonoids prevent the assembly of amyloid curli fibres and can interfere with bacterial biofilm formation. Environmental Microbiology, 2020, 22, 5280-5299.	3.8	28
7	Local c-di-GMP Signaling in the Control of Synthesis of the E. coli Biofilm Exopolysaccharide pEtN-Cellulose. Journal of Molecular Biology, 2020, 432, 4576-4595.	4.2	53
8	Non-lethal exposure to H2O2 boosts bacterial survival and evolvability against oxidative stress. PLoS Genetics, 2020, 16, e1008649.	3.5	59
9	Linking bacterial growth, survival, and multicellularity – small signaling molecules as triggers and drivers. Current Opinion in Microbiology, 2020, 55, 57-66.	5.1	59
10	Targeting Bacterial Biofilms by the Green Tea Polyphenol EGCG. Molecules, 2019, 24, 2403.	3.8	60
11	Cellulose in Bacterial Biofilms. Biologically-inspired Systems, 2019, , 355-392.	0.2	17
12	The <i>Escherichia coli</i> MarA protein regulates the <i>ycgZ</i> – <i>ymgABC</i> operon to inhibit biofilm formation. Molecular Microbiology, 2019, 112, 1609-1625.	2.5	17
13	Recent Advances and Current Trends in Nucleotide Second Messenger Signaling in Bacteria. Journal of Molecular Biology, 2019, 431, 908-927.	4.2	41
14	Genetic dissection of <i>Escherichia coli</i> 's master diguanylate cyclase DgcE: Role of the N-terminal MASE1 domain and direct signal input from a GTPase partner system. PLoS Genetics, 2019, 15, e1008059.	3.5	28
15	A c-di-GMP-Based Switch Controls Local Heterogeneity of Extracellular Matrix Synthesis which Is Crucial for Integrity and Morphogenesis of <i>Escherichia coli</i> Macrocolony Biofilms. Journal of Molecular Biology, 2019, 431, 4775-4793.	4.2	41
16	Transmembrane redox control and proteolysis of PdeC, a novel type of c-di-GMP phosphodiesterase. EMBO Journal, 2018, 37, .	7.8	37
17	Phosphoethanolamine cellulose: A naturally produced chemically modified cellulose. Science, 2018, 359, 334-338.	12.6	208
18	Discovery of Phosphoethanolamine Cellulose and the Genetic Basis for its Biosynthesis in E. coli Biofilms. Biophysical Journal, 2018, 114, 158a.	0.5	0

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19	The Intestinal Roundworm <i>Ascaris suum</i> Releases Antimicrobial Factors Which Interfere With Bacterial Growth and Biofilm Formation. <i>Frontiers in Cellular and Infection Microbiology</i> , 2018, 8, 271.	3.9	41
20	Spatial organization of different sigma factor activities and c-di-GMP signalling within the three-dimensional landscape of a bacterial biofilm. <i>Open Biology</i> , 2018, 8, .	3.6	61
21	More than Enzymes That Make or Break Cyclic Di-GMP Local Signaling in the Interactome of GGDEF/EAL Domain Proteins of <i>Escherichia coli</i> . <i>MBio</i> , 2017, 8, .	4.1	136
22	Experimental Detection and Visualization of the Extracellular Matrix in Macrocolony Biofilms. <i>Methods in Molecular Biology</i> , 2017, 1657, 133-145.	0.9	19
23	The green tea polyphenol EGCG inhibits <i>E. coli</i> biofilm formation by impairing amyloid curli fibre assembly and downregulating the biofilm regulator CsgD via the $\lambda$ -dependent sRNA RybB. <i>Molecular Microbiology</i> , 2016, 101, 136-151.	2.5	107
24	Trigger phosphodiesterases as a novel class of c-di-GMP effector proteins. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016, 371, 20150498.	4.0	71
25	Bacterial Signal Transduction by Cyclic Di-GMP and Other Nucleotide Second Messengers. <i>Journal of Bacteriology</i> , 2016, 198, 15-26.	2.2	127
26	Genome-Based Comparison of Cyclic Di-GMP Signaling in Pathogenic and Commensal <i>Escherichia coli</i> Strains. <i>Journal of Bacteriology</i> , 2016, 198, 111-126.	2.2	59
27	Systematic Nomenclature for GGDEF and EAL Domain-Containing Cyclic Di-GMP Turnover Proteins of <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2016, 198, 7-11.	2.2	96
28	Vertical stratification of matrix production is essential for physical integrity and architecture of macrocolony biofilms of <i>Escherichia coli</i> . <i>Environmental Microbiology</i> , 2015, 17, 5073-5088.	3.8	44
29	Logical-continuous modelling of post-translationally regulated bistability of curli fiber expression in <i>Escherichia coli</i> . <i>BMC Systems Biology</i> , 2015, 9, 39.	3.0	11
30	Small RNAs in the control of RpoS, CsgD, and biofilm architecture of <i>Escherichia coli</i> . <i>RNA Biology</i> , 2014, 11, 494-507.	3.1	146
31	Stress responses go three dimensional – the spatial order of physiological differentiation in bacterial macrocolony biofilms. <i>Environmental Microbiology</i> , 2014, 16, 1455-1471.	3.8	153
32	Cyclic-di-GMP signalling and biofilm-related properties of the Shiga toxin-producing 2011 German outbreak <i>Escherichia coli</i> O104:H4. <i>EMBO Molecular Medicine</i> , 2014, 6, 1622-1637.	6.9	60
33	Novel tricks played by the second messenger c-di-GMP in bacterial biofilm formation. <i>EMBO Journal</i> , 2013, 32, 322-323.	7.8	10
34	Reply to “Precedence for the Structural Role of Flagella in Biofilms”. <i>MBio</i> , 2013, 4, e00245-13.	4.1	1
35	Small Regulatory RNAs in the Control of Motility and Biofilm Formation in <i>E. coli</i> and <i>Salmonella</i> . <i>International Journal of Molecular Sciences</i> , 2013, 14, 4560-4579.	4.1	142
36	The EAL domain protein YciR acts as a trigger enzyme in a c-di-GMP signalling cascade in <i>E. coli</i> biofilm control. <i>EMBO Journal</i> , 2013, 32, 2001-2014.	7.8	157

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37	Microanatomy at Cellular Resolution and Spatial Order of Physiological Differentiation in a Bacterial Biofilm. <i>MBio</i> , 2013, 4, e00103-13.	4.1	286
38	Cellulose as an Architectural Element in Spatially Structured <i>Escherichia coli</i> Biofilms. <i>Journal of Bacteriology</i> , 2013, 195, 5540-5554.	2.2	291
39	The global repressor FlhZ antagonizes gene expression by $\sigma^S$ -containing RNA polymerase due to overlapping DNA binding specificity. <i>Nucleic Acids Research</i> , 2012, 40, 4783-4793.	14.5	38
40	"Life-style" control networks in <i>Escherichia coli</i> : Signaling by the second messenger c-di-GMP. <i>Journal of Biotechnology</i> , 2012, 160, 10-16.	3.8	94
41	The enemy within us: lessons from the 2011 European <i>Escherichia coli</i> O104:H4 outbreak. <i>EMBO Molecular Medicine</i> , 2012, 4, 841-848.	6.9	215
42	Targeting of <i>csgD</i> by the small regulatory RNA RprA links stationary phase, biofilm formation and cell envelope stress in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2012, 84, 51-65.	2.5	111
43	Molecular function and potential evolution of the biofilm-modulating blue light signalling pathway of <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2012, 85, 893-906.	2.5	46
44	Stationary-Phase Gene Regulation in <i>Escherichia coli</i> . <i>EcoSal Plus</i> , 2011, 4, .	5.4	48
45	Rare codons play a positive role in the expression of the stationary phase sigma factor RpoS ( $\sigma^S$ ) in <i>Escherichia coli</i> . <i>RNA Biology</i> , 2011, 8, 913-921.	3.1	25
46	<i>Escherichia coli</i> $\sigma^S$ senses sequence and conformation of the promoter spacer region. <i>Nucleic Acids Research</i> , 2011, 39, 5109-5118.	14.5	58
47	Cyclic-di-GMP Reaches Out into the Bacterial RNA World. <i>Science Signaling</i> , 2010, 3, pe44.	3.6	35
48	The influence of Hfq and ribonucleases on the stability of the small non-coding RNA OxyS and its target <i>rpoS</i> in <i>E. coli</i> is growth phase dependent. <i>RNA Biology</i> , 2009, 6, 584-594.	3.1	34
49	Gene expression patterns and differential input into curli fimbriae regulation of all GGDEF/EAL domain proteins in <i>Escherichia coli</i> . <i>Microbiology (United Kingdom)</i> , 2009, 155, 1318-1331.	1.8	150
50	Principles of c-di-GMP signalling in bacteria. <i>Nature Reviews Microbiology</i> , 2009, 7, 263-273.	28.6	1,320
51	The BLUF-EAL protein YcgF acts as a direct anti-repressor in a blue-light response of <i>Escherichia coli</i> . <i>Genes and Development</i> , 2009, 23, 522-534.	5.9	165
52	Proteolysis of $\sigma^S$ (RpoS) and the general stress response in <i>Escherichia coli</i> . <i>Research in Microbiology</i> , 2009, 160, 667-676.	2.1	157
53	Proteolysis in prokaryotes "from molecular machines to a systems perspective. <i>Research in Microbiology</i> , 2009, 160, 615-617.	2.1	4
54	Bacterial nucleotide-based second messengers. <i>Current Opinion in Microbiology</i> , 2009, 12, 170-176.	5.1	158

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55	A role for Lon protease in the control of the acid resistance genes of <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2008, 69, 534-547.	2.5	35
56	Inverse regulatory coordination of motility and curli-mediated adhesion in <i>Escherichia coli</i> . <i>Genes and Development</i> , 2008, 22, 2434-2446.	5.9	299
57	The $\sigma^S$ subunit of RNA polymerase as a signal integrator and network master regulator in the general stress response in <i>Escherichia coli</i> . <i>Science Progress</i> , 2007, 90, 103-127.	1.9	65
58	Stationary phase reorganisation of the <i>Escherichia coli</i> transcription machinery by Crl protein, a fine-tuner of $\sigma^S$ activity and levels. <i>EMBO Journal</i> , 2007, 26, 1569-1578.	7.8	107
59	The -35 sequence location and the Fis- $\sigma$ factor interface determine $\sigma$ -selectivity of the proP (P2) promoter in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2007, 63, 780-96.	2.5	28
60	The molecular basis of selective promoter activation by the $\sigma$ subunit of RNA polymerase. <i>Molecular Microbiology</i> , 2007, 63, 1296-1306.	2.5	147
61	Cellular levels and activity of the flagellar sigma factor FliA of <i>Escherichia coli</i> are controlled by FlgM-modulated proteolysis. <i>Molecular Microbiology</i> , 2007, 65, 76-89.	2.5	75
62	Role of the spacer between the -35 and -10 regions in sigma promoter selectivity in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2006, 59, 1037-1051.	2.5	73
63	Poly(A)-polymerase I links transcription with mRNA degradation via $\sigma^S$ proteolysis. <i>Molecular Microbiology</i> , 2006, 60, 177-188.	2.5	24
64	Cyclic-di-GMP-mediated signalling within the $\sigma$ network of <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2006, 62, 1014-1034.	2.5	250
65	Genome-Wide Analysis of the General Stress Response Network in <i>Escherichia coli</i> : $\sigma^S$ -Dependent Genes, Promoters, and Sigma Factor Selectivity. <i>Journal of Bacteriology</i> , 2005, 187, 1591-1603.	2.2	743
66	A two-component phosphotransfer network involving ArcB, ArcA, and RssB coordinates synthesis and proteolysis of $\sigma^S$ (RpoS) in <i>E. coli</i> . <i>Genes and Development</i> , 2005, 19, 2770-2781.	5.9	169
67	Differential ability of $\sigma^S$ and $\sigma^{70}$ of <i>Escherichia coli</i> to utilize promoters containing half or full UP-element sites. <i>Molecular Microbiology</i> , 2004, 55, 250-260.	2.5	37
68	Sequential recognition of two distinct sites in $\sigma^S$ by the proteolytic targeting factor RssB and ClpX. <i>EMBO Journal</i> , 2003, 22, 4111-4120.	7.8	91
69	Dynamic control of Dps protein levels by ClpXP and ClpAP proteases in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2003, 49, 1605-1614.	2.5	70
70	Multiple stress signal integration in the regulation of the complex $\sigma^S$ -dependent <i>csiD-ygaF-gabDTP</i> operon in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2003, 51, 799-811.	2.5	62
71	The General Stress Response in Gram-Negative Bacteria. , 0, , 251-289.		41
72	Role of Cyclic Di-GMP in the Regulatory Networks of <i>Escherichia coli</i> . , 0, , 230-252.		9

#	ARTICLE	IF	CITATIONS
73	General Stress Response in <i>Bacillus subtilis</i> and Related Gram-Positive Bacteria. , 0, , 301-318.		23