

# E Gerhart H Wagner

## List of Publications by Year in descending order

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49  
papers

4,921  
citations

136740

32  
h-index

205818

48  
g-index

52  
all docs

52  
docs citations

52  
times ranked

3675  
citing authors

#	ARTICLE	IF	CITATIONS
1	Small RNAs in Bacteria and Archaea. <i>Advances in Genetics</i> , 2015, 90, 133-208.	0.8	462
2	Antisense RNA Control in Bacteria, Phages, and Plasmids. <i>Annual Review of Microbiology</i> , 1994, 48, 713-742.	2.9	427
3	RNomics in <i>Escherichia coli</i> detects new sRNA species and indicates parallel transcriptional output in bacteria. <i>Nucleic Acids Research</i> , 2003, 31, 6435-6443.	6.5	388
4	Hfq-dependent regulation of OmpA synthesis is mediated by an antisense RNA. <i>Genes and Development</i> , 2005, 19, 2355-2366.	2.7	271
5	The Small RNA IstR Inhibits Synthesis of an SOS-Induced Toxic Peptide. <i>Current Biology</i> , 2004, 14, 2271-2276.	1.8	241
6	An Antisense RNA Inhibits Translation by Competing with Standby Ribosomes. <i>Molecular Cell</i> , 2007, 26, 381-392.	4.5	224
7	H $\epsilon$ S-mediated repression of CRISPR-based immunity in <i>Escherichia coli</i> K12 can be relieved by the transcription activator LeuO. <i>Molecular Microbiology</i> , 2010, 77, 1380-1393.	1.2	220
8	RNA antitoxins. <i>Current Opinion in Microbiology</i> , 2007, 10, 117-124.	2.3	216
9	12 Antisense RNAs in bacteria and their genetic elements. <i>Advances in Genetics</i> , 2002, 46, 361-398.	0.8	213
10	The role of RNAs in the regulation of virulence-gene expression. <i>Current Opinion in Microbiology</i> , 2006, 9, 229-236.	2.3	174
11	Target identification of small noncoding RNAs in bacteria. <i>Current Opinion in Microbiology</i> , 2007, 10, 262-270.	2.3	165
12	A small SOS-induced toxin is targeted against the inner membrane in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2008, 70, 258-270.	1.2	161
13	RNAs actively cycle on the Sm-like protein Hfq. <i>Genes and Development</i> , 2010, 24, 2621-2626.	2.7	143
14	Impact of bacterial sRNAs in stress responses. <i>Biochemical Society Transactions</i> , 2017, 45, 1203-1212.	1.6	133
15	Cycling of RNAs on Hfq. <i>RNA Biology</i> , 2013, 10, 619-626.	1.5	103
16	Sigma E controls biogenesis of the antisense RNA MicA. <i>Nucleic Acids Research</i> , 2007, 35, 1279-1288.	6.5	101
17	PcnB is required for the rapid degradation of RNAI, the antisense RNA that controls the copy number of ColE1-related plasmids. <i>Molecular Microbiology</i> , 1993, 9, 1131-1142.	1.2	93
18	Antisense RNAs everywhere?. <i>Trends in Genetics</i> , 2002, 18, 223-226.	2.9	91

#	ARTICLE	IF	CITATIONS
19	Antisense RNA-mediated transcriptional attenuation: an in vitro study of plasmid pT181. <i>Molecular Microbiology</i> , 2002, 35, 1469-1482.	1.2	74
20	Lead(II) as a probe for investigating RNA structure in vivo. <i>Rna</i> , 2002, 8, 534-541.	1.6	70
21	A mixed double negative feedback loop between the sRNA MicF and the global regulator Lrp. <i>Molecular Microbiology</i> , 2012, 84, 414-427.	1.2	70
22	Two regulatory <sc>RNA</sc> elements affect <sc>T</sc>is<sc>B</sc>-dependent depolarization and persister formation. <i>Molecular Microbiology</i> , 2017, 103, 1020-1033.	1.2	69
23	An unusual structure formed by antisense-target RNA binding involves an extended kissing complex with a four-way junction and a side-by-side helical alignment. <i>Rna</i> , 2000, 6, 311-324.	1.6	66
24	Differential Translation Tunes Uneven Production of Operon-Encoded Proteins. <i>Cell Reports</i> , 2013, 4, 938-944.	2.9	64
25	Regulation of plasmid R1 replication: PcnB and RNase E expedite the decay of the antisense RNA, CopA. <i>Molecular Microbiology</i> , 1997, 26, 493-504.	1.2	59
26	Structural and functional analyses of the FinP antisense RNA regulatory system of the F conjugative plasmid. <i>Molecular Microbiology</i> , 1993, 10, 35-43.	1.2	45
27	An unusually long-lived antisense RNA in plasmid copy number control: in vivo RNAs encoded by the streptococcal plasmid pIP501. <i>Journal of Molecular Biology</i> , 1996, 255, 275-288.	2.0	45
28	RNA-based regulation in type I toxin-antitoxin systems and its implication for bacterial persistence. <i>Current Genetics</i> , 2017, 63, 1011-1016.	0.8	45
29	The toxin-antitoxin system <i>tisB-istR1</i> . <i>RNA Biology</i> , 2012, 9, 1513-1519.	1.5	43
30	An Antisense RNA-Mediated Transcriptional Attenuation Mechanism Functions in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2002, 184, 2740-2747.	1.0	42
31	Replication control of plasmid R1: disruption of an inhibitory RNA structure that sequesters the repA ribosome-binding site permits tap-independent RepA synthesis. <i>Molecular Microbiology</i> , 1994, 12, 49-60.	1.2	41
32	Hfq-dependent <sc>mRNA</sc> unfolding promotes <sc>sRNA</sc>-based inhibition of translation. <i>EMBO Journal</i> , 2019, 38, .	3.5	37
33	Degradation pathway of CopA, the antisense RNA that controls replication of plasmid R1. <i>Microbiology (United Kingdom)</i> , 1998, 144, 1907-1917.	0.7	34
34	One Gene and Two Proteins: a Leaderless mRNA Supports the Translation of a Shorter Form of the <i>Shigella</i> VirF Regulator. <i>MBio</i> , 2016, 7, .	1.8	34
35	Kill the messenger: bacterial antisense RNA promotes mRNA decay. <i>Nature Structural and Molecular Biology</i> , 2009, 16, 804-806.	3.6	33
36	The ribosomal protein S1-dependent standby site in <i>tisB</i> mRNA consists of a single-stranded region and a 5' structure element. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 15901-15906.	3.3	32

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37	Massive functional mapping of a 5' UTR by saturation mutagenesis, phenotypic sorting and deep sequencing. <i>Nucleic Acids Research</i> , 2013, 41, e122-e122.	6.5	27
38	Unstructured 5' tails act through ribosome standby to override inhibitory structure at ribosome binding sites. <i>Nucleic Acids Research</i> , 2018, 46, 4188-4199.	6.5	27
39	Dealing with stable structures at ribosome binding sites: Bacterial translation and ribosome standby. <i>RNA Biology</i> , 2007, 4, 113-117.	1.5	24
40	Disruption of the gene encoding the V-ATPase subunit A results in inhibition of normal growth and abolished sporulation in <i>Aspergillus nidulans</i> . <i>Microbiology (United Kingdom)</i> , 2004, 150, 743-748.	0.7	19
41	Small RNAs OmrA and OmrB promote class III flagellar gene expression by inhibiting the synthesis of anti-Sigma factor FlgM. <i>RNA Biology</i> , 2020, 17, 872-880.	1.5	18
42	Fluorescent CRISPR Adaptation Reporter for rapid quantification of spacer acquisition. <i>Scientific Reports</i> , 2017, 7, 10392.	1.6	15
43	RNA-sequence data normalization through in silico prediction of reference genes: the bacterial response to DNA damage as case study. <i>BioData Mining</i> , 2017, 10, 30.	2.2	15
44	Exploring the complex world of RNA regulation. <i>Biology of the Cell</i> , 2008, 100, e1-3.	0.7	11
45	An RNA pseudoknot is essential for standby-mediated translation of the <i>tisB</i> toxin mRNA in <i>Escherichia coli</i> . <i>Nucleic Acids Research</i> , 2020, 48, 12336-12347.	6.5	10
46	Loop Swapping in an Antisense RNA/Target RNA Pair Changes Directionality of Helix Progression. <i>Journal of Biological Chemistry</i> , 2003, 278, 35558-35563.	1.6	9
47	A method to map changes in bacterial surface composition induced by regulatory RNAs in <i>Escherichia coli</i> and <i>Staphylococcus aureus</i> . <i>Biochimie</i> , 2014, 106, 175-179.	1.3	8
48	Persistor Formation Driven by TisB-Dependent Membrane Depolarization. , 2019, , 77-97.		4
49	The Length of a DNA T-Tract Modulates Expression of a Virulence-Regulating sRNA. <i>Molecular Cell</i> , 2020, 80, 175-177.	4.5	0