

Charles J Dorman

List of Publications by Year in descending order

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140
papers

10,118
citations

34016

52
h-index

38300

95
g-index

236
all docs

236
docs citations

236
times ranked

5806
citing authors

#	ARTICLE	IF	CITATIONS
1	Bacterial nucleoid-associated proteins, nucleoid structure and gene expression. <i>Nature Reviews Microbiology</i> , 2010, 8, 185-195.	13.6	755
2	A physiological role for DNA supercoiling in the osmotic regulation of gene expression in <i>S. typhimurium</i> and <i>E. coli</i> . <i>Cell</i> , 1988, 52, 569-584.	13.5	736
3	H-NS: a universal regulator for a dynamic genome. <i>Nature Reviews Microbiology</i> , 2004, 2, 391-400.	13.6	485
4	The transcriptional landscape and small RNAs of <i>Salmonella enterica</i> serovar Typhimurium. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, E1277-86.	3.3	373
5	H-NS, the genome sentinel. <i>Nature Reviews Microbiology</i> , 2007, 5, 157-161.	13.6	314
6	DNA supercoiling and the anaerobic and growth phase regulation of tonB gene expression. <i>Journal of Bacteriology</i> , 1988, 170, 2816-2826.	1.0	259
7	DNA supercoiling and environmental regulation of virulence gene expression in <i>Shigella flexneri</i> . <i>Nature</i> , 1990, 344, 789-792.	13.7	249
8	Characterization of porin and ompR mutants of a virulent strain of <i>Salmonella typhimurium</i> : ompR mutants are attenuated in vivo. <i>Infection and Immunity</i> , 1989, 57, 2136-2140.	1.0	249
9	Human Health and Ocean Pollution. <i>Annals of Global Health</i> , 2020, 86, 151.	0.8	240
10	Anti-silencing: overcoming H-NS-mediated repression of transcription in Gram-negative enteric bacteria. <i>Microbiology (United Kingdom)</i> , 2008, 154, 2533-2545.	0.7	232
11	Regulation of gene expression by histone-like proteins in bacteria. <i>Current Opinion in Genetics and Development</i> , 2003, 13, 179-184.	1.5	216
12	An H-NS-like Stealth Protein Aids Horizontal DNA Transmission in Bacteria. <i>Science</i> , 2007, 315, 251-252.	6.0	204
13	Fimbrial phase variation in <i>Escherichia coli</i> : dependence on integration host factor and homologies with other site-specific recombinases. <i>Journal of Bacteriology</i> , 1987, 169, 3840-3843.	1.0	179
14	A global role for Fis in the transcriptional control of metabolism and type III secretion in <i>Salmonella enterica</i> serovar Typhimurium. <i>Microbiology (United Kingdom)</i> , 2004, 150, 2037-2053.	0.7	175
15	Genome architecture and global gene regulation in bacteria: making progress towards a unified model?. <i>Nature Reviews Microbiology</i> , 2013, 11, 349-355.	13.6	172
16	An overlap between osmotic and anaerobic stress responses: a potential role for DNA supercoiling in the coordinate regulation of gene expression. <i>Molecular Microbiology</i> , 1989, 3, 933-942.	1.2	159
17	The integration host factor (IHF) integrates stationary-phase and virulence gene expression in <i>Salmonella enterica</i> serovar Typhimurium. <i>Molecular Microbiology</i> , 2006, 59, 1831-1847.	1.2	159
18	DNA supercoiling and environmental regulation of gene expression in pathogenic bacteria. <i>Infection and Immunity</i> , 1991, 59, 745-749.	1.0	148

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19	The Shigella virulence gene regulatory cascade: a paradigm of bacterial gene control mechanisms. <i>Molecular Microbiology</i> , 1998, 29, 677-684.	1.2	143
20	Domain organization and oligomerization among H-NS-like nucleoid-associated proteins in bacteria. <i>Trends in Microbiology</i> , 1999, 7, 124-128.	3.5	137
21	TonB protein of <i>Salmonella typhimurium</i> . <i>Journal of Molecular Biology</i> , 1990, 216, 897-910.	2.0	133
22	A Fundamental Regulatory Mechanism Operating through OmpR and DNA Topology Controls Expression of <i>Salmonella</i> Pathogenicity Islands SPI-1 and SPI-2. <i>PLoS Genetics</i> , 2012, 8, e1002615.	1.5	119
23	Structure and function of X-Pro dipeptide repeats in the TonB proteins of <i>Salmonella typhimurium</i> and <i>Escherichia coli</i> . <i>Journal of Molecular Biology</i> , 1990, 216, 883-895.	2.0	104
24	Chapter 2 Nucleoid-Associated Proteins and Bacterial Physiology. <i>Advances in Applied Microbiology</i> , 2009, 67, 47-64.	1.3	104
25	An extended role for the nucleoid structuring protein H-NS in the virulence gene regulatory cascade of <i>Shigella flexneri</i> . <i>Molecular Microbiology</i> , 2003, 47, 825-838.	1.2	95
26	Sequence-imposed structural constraints in the TonB protein of <i>E. coli</i> . <i>FEBS Letters</i> , 1986, 208, 211-216.	1.3	92
27	Bacterial Regulon Evolution: Distinct Responses and Roles for the Identical OmpR Proteins of <i>Salmonella Typhimurium</i> and <i>Escherichia coli</i> in the Acid Stress Response. <i>PLoS Genetics</i> , 2014, 10, e1004215.	1.5	90
28	DNA supercoiling is a fundamental regulatory principle in the control of bacterial gene expression. <i>Biophysical Reviews</i> , 2016, 8, 89-100.	1.5	89
29	DNA supercoiling is differentially regulated by environmental factors and FIS in <i>Escherichia coli</i> and <i>Salmonella enterica</i> . <i>Molecular Microbiology</i> , 2011, 80, 85-101.	1.2	86
30	DNA supercoiling and transcription in bacteria: a two-way street. <i>BMC Molecular and Cell Biology</i> , 2019, 20, 26.	1.0	86
31	Competitive interaction of the OxyR DNA-binding protein and the Dam methylase at the antigen 43 gene regulatory region in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2002, 44, 509-520.	1.2	84
32	Roles for DNA supercoiling and the Fis protein in modulating expression of virulence genes during intracellular growth of <i>Salmonella enterica</i> serovar <i>Typhimurium</i> . <i>Molecular Microbiology</i> , 2006, 62, 869-882.	1.2	84
33	Compensatory Evolution of Gene Regulation in Response to Stress by <i>Escherichia coli</i> Lacking RpoS. <i>PLoS Genetics</i> , 2009, 5, e1000671.	1.5	84
34	Genome-wide analysis of the H-NS and Sfh regulatory networks in <i>Salmonella Typhimurium</i> identifies a plasmid-encoded transcription silencing mechanism. <i>Molecular Microbiology</i> , 2010, 76, 1250-1265.	1.2	84
35	H-NS-like nucleoid-associated proteins, mobile genetic elements and horizontal gene transfer in bacteria. <i>Plasmid</i> , 2014, 75, 1-11.	0.4	78
36	DNA supercoiling is a fundamental regulatory principle in the control of bacterial gene expression. <i>Biophysical Reviews</i> , 2016, 8, 209-220.	1.5	73

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37	Regulation of virulence gene expression in <i>Shigella flexneri</i> , a facultative intracellular pathogen. <i>International Journal of Medical Microbiology</i> , 2001, 291, 89-96.	1.5	72
38	Three-way interactions among the Sfh, StpA and H-NS nucleoid-structuring proteins of <i>Shigella flexneri</i> 2a strain 2457T. <i>Molecular Microbiology</i> , 2003, 48, 1401-1416.	1.2	72
39	Coupling of <i>Escherichia coli</i> hns mRNA levels to DNA synthesis by autoregulation: implications for growth phase control. <i>Molecular Microbiology</i> , 1995, 18, 101-113.	1.2	68
40	A role for the <i>Escherichia coli</i> H-NS-like protein StpA in <i>OmpF</i> porin expression through modulation of <i>micF</i> RNA stability. <i>Molecular Microbiology</i> , 2000, 38, 126-139.	1.2	68
41	LeuO is a global regulator of gene expression in <i>Salmonella enterica</i> serovar Typhimurium. <i>Molecular Microbiology</i> , 2012, 85, 1072-1089.	1.2	68
42	Osmotic and growth-phase dependent regulation of the <i>eta</i> gene of <i>Staphylococcus aureus</i> : a role for DNA supercoiling. <i>Molecular Genetics and Genomics</i> , 1992, 232, 49-57.	2.4	67
43	Dna Supercoiling and Bacterial Gene Expression. <i>Science Progress</i> , 2006, 89, 151-166.	1.0	67
44	H-NS Antagonism in <i>Shigella flexneri</i> by VirB, a Virulence Gene Transcription Regulator That Is Closely Related to Plasmid Partition Factors. <i>Journal of Bacteriology</i> , 2007, 189, 3403-3413.	1.0	67
45	Bacterial DNA topology and infectious disease. <i>Nucleic Acids Research</i> , 2009, 37, 672-678.	6.5	67
46	DNA topology and the global control of bacterial gene expression: implications for the regulation of virulence gene expression. <i>Microbiology (United Kingdom)</i> , 1995, 141, 1271-1280.	0.7	67
47	<i>Shigella flexneri</i> 2a strain 2457T expresses three members of the H-NS-like protein family: characterization of the Sfh protein. <i>Molecular Genetics and Genomics</i> , 2003, 270, 66-77.	1.0	65
48	Nucleoid-associated protein HU controls three regulons that coordinate virulence, response to stress and general physiology in <i>Salmonella enterica</i> serovar Typhimurium. <i>Microbiology (United Kingdom)</i> , 2007, 151, 1072-1089.	1.0	64
49	DNA bridging and antibridging: a role for bacterial nucleoid-associated proteins in regulating the expression of laterally acquired genes. <i>FEMS Microbiology Reviews</i> , 2009, 33, 587-592.	3.9	63
50	DNA relaxation-dependent phase biasing of the <i>fim</i> genetic switch in <i>Escherichia coli</i> depends on the interplay of H-NS, IHF and LRP. <i>Molecular Microbiology</i> , 2009, 74, 1071-1082.	1.2	62
51	Differential regulation of the plasmid-encoded genes in the <i>Shigella flexneri</i> virulence regulon. <i>Molecular Genetics and Genomics</i> , 1997, 256, 93-103.	2.4	60
52	Molecular Dissection of VirB, a Key Regulator of the Virulence Cascade of <i>Shigella flexneri</i> . <i>Journal of Biological Chemistry</i> , 2002, 277, 15333-15344.	1.6	57
53	Role of hns in the virulence phenotype of pathogenic salmonellae. <i>Molecular Microbiology</i> , 1994, 13, 133-140.	1.2	56
54	Hierarchical gene regulators adapt to its host milieu. <i>International Journal of Medical Microbiology</i> , 2005, 294, 487-502.	1.5	56

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55	The site-specific recombination system regulating expression of the Type 1 fimbrial subunit gene of <i>Escherichia coli</i> is sensitive to changes in DNA supercoiling. <i>Molecular Microbiology</i> , 1994, 14, 975-988.	1.2	54
56	Isolation and characterization of a <i>topA</i> mutant of <i>Shigella flexneri</i> . <i>Molecular Microbiology</i> , 1993, 7, 351-358.	1.2	52
57	Control of <i>Escherichia coli</i> type 1 fimbrial gene expression in stationary phase: a negative role for RpoS. <i>Molecular Genetics and Genomics</i> , 1997, 254, 13-20.	2.4	52
58	<i>Escherichia coli</i> <i>tyrT</i> gene transcription is sensitive to DNA supercoiling in its native chromosomal context: effect of DNA topoisomerase IV overexpression on <i>tyrT</i> promoter function. <i>Molecular Microbiology</i> , 1994, 14, 151-161.	1.2	50
59	In vivo analysis of the interactions of the LysR-like regulator SpvR with the operator sequences of the <i>spvA</i> and <i>spvR</i> virulence genes of <i>Salmonella typhimurium</i> . <i>Molecular Microbiology</i> , 1998, 30, 91-105.	1.2	49
60	The Leucine-Responsive Regulatory Protein, Lrp, Activates Transcription of the <i>fim</i> Operon in <i>Salmonella enterica</i> Serovar Typhimurium via the <i>fimZ</i> Regulatory Gene. <i>Journal of Bacteriology</i> , 2008, 190, 602-612.	1.0	49
61	Co-operative roles for DNA supercoiling and nucleoid-associated proteins in the regulation of bacterial transcription. <i>Biochemical Society Transactions</i> , 2013, 41, 542-547.	1.6	49
62	When is a transcription factor a NAP?. <i>Current Opinion in Microbiology</i> , 2020, 55, 26-33.	2.3	48
63	Overexpression of the <i>Shigella flexneri</i> genes coding for DNA topoisomerase IV compensates for loss of DNA topoisomerase I: effect on virulence gene expression. <i>Molecular Microbiology</i> , 1995, 15, 507-517.	1.2	47
64	The <i>gyr</i> genes of <i>Salmonella enterica</i> serovar Typhimurium are repressed by the factor for inversion stimulation, Fis. <i>Molecular Genetics and Genomics</i> , 2003, 270, 56-65.	1.0	47
65	DNA Supercoiling and the Lrp Protein Determine the Directionality of <i>fim</i> Switch DNA Inversion in <i>Escherichia coli</i> K-12. <i>Journal of Bacteriology</i> , 2006, 188, 5356-5363.	1.0	47
66	Function of Nucleoid-Associated Proteins in Chromosome Structuring and Transcriptional Regulation. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2014, 24, 316-331.	1.0	47
67	A role for the leucine-responsive regulatory protein and integration host factor in the regulation of the <i>Salmonella</i> plasmid virulence (<i>spv</i>) locus in <i>Salmonella typhimurium</i> . <i>Molecular Microbiology</i> , 1999, 34, 134-145.	1.2	46
68	Effects of local transcription and H-NS on inversion of the <i>fim</i> switch of <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2000, 36, 457-466.	1.2	46
69	The StpA Protein Functions as a Molecular Adapter To Mediate Repression of the <i>bgl</i> Operon by Truncated H-NS in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 1998, 180, 994-997.	1.0	45
70	Requirement for the molecular adapter function of StpA at the <i>Escherichia coli</i> <i>bgl</i> promoter depends upon the level of truncated H-NS protein. <i>Molecular Microbiology</i> , 2001, 42, 903-917.	1.2	44
71	Transcription of the <i>Salmonella typhimurium</i> <i>spv</i> virulence locus is regulated negatively by the nucleoid-associated protein H-NS. <i>FEMS Microbiology Letters</i> , 1994, 121, 99-105.	0.7	42
72	In Vivo DNA-Binding and Oligomerization Properties of the <i>Shigella flexneri</i> AraC-Like Transcriptional Regulator VirF as Identified by Random and Site-Specific Mutagenesis. <i>Journal of Bacteriology</i> , 2002, 184, 531-539.	1.0	41

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73	DNA topology and bacterial virulence gene regulation. <i>Trends in Microbiology</i> , 1993, 1, 92-99.	3.5	40
74	Functional analysis of the FimE integrase of <i>Escherichia coli</i> K-12: isolation of mutant derivatives with altered DNA inversion preferences. <i>Molecular Microbiology</i> , 1999, 34, 965-979.	1.2	40
75	Expression of the Fis protein is sustained in late exponential and stationary phase cultures of <i>Salmonella enterica</i> serovar Typhimurium grown in the absence of aeration. <i>Molecular Microbiology</i> , 2007, 66, 237-251.	1.2	40
76	The Effect of Mobile Element IS10 on Experimental Regulatory Evolution in <i>Escherichia coli</i> . <i>Molecular Biology and Evolution</i> , 2010, 27, 2105-2112.	3.5	40
77	Thermal regulation of <i>fimA</i> , the <i>Escherichia coli</i> gene coding for the type 1 fimbrial subunit protein. <i>FEMS Microbiology Letters</i> , 1992, 99, 125-130.	0.7	34
78	The DNA supercoiling-sensitive expression of the <i>Salmonella typhimurium his</i> operon requires the <i>his</i> attenuator and is modulated by anaerobiosis and by osmolarity. <i>Molecular Microbiology</i> , 1992, 6, 2467-2476.	1.2	33
79	DNA sequence heterogeneity in Fim tyrosine integrase recombinase binding elements and functional motif asymmetries determine the directionality of the <i>fim</i> genetic switch in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2008, 67, 171-187.	1.2	33
80	Regulatory Hierarchies Controlling Virulence Gene Expression in <i>Shigella flexneri</i> and <i>Vibrio cholerae</i> . <i>Frontiers in Microbiology</i> , 2018, 9, 2686.	1.5	31
81	Multicopy <i>fimB</i> gene expression in <i>Escherichia coli</i> : binding to inverted repeats in vivo, effect on <i>fimA</i> gene transcription and DNA inversion. <i>Molecular Microbiology</i> , 1996, 21, 1161-1173.	1.2	28
82	DNA topology and adaptation of <i>salmonella typhimurium</i> to an intracellular environment. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2000, 355, 565-574.	1.8	28
83	A Rho-dependent phase-variable transcription terminator controls expression of the FimE recombinase in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2002, 45, 1107-1117.	1.2	28
84	Rational Design of an Artificial Genetic Switch: Co-Option of the H-NS-Repressed <i>proU</i> Operon by the VirB Virulence Master Regulator. <i>Journal of Bacteriology</i> , 2011, 193, 5950-5960.	1.0	28
85	In vitro DNA-binding properties of VirB, the <i>Shigella flexneri</i> virulence regulatory protein. <i>FEBS Letters</i> , 2003, 545, 183-187.	1.3	27
86	VirB-Mediated Positive Feedback Control of the Virulence Gene Regulatory Cascade of <i>Shigella flexneri</i> . <i>Journal of Bacteriology</i> , 2012, 194, 5264-5273.	1.0	27
87	A novel role for antibiotic resistance plasmids in facilitating <i>Salmonella</i> adaptation to non-host environments. <i>Environmental Microbiology</i> , 2014, 16, 950-962.	1.8	26
88	Control of virulence gene transcription by indirect readout in <i>Vibrio cholerae</i> and <i>Salmonella enterica</i> serovar Typhimurium. <i>Environmental Microbiology</i> , 2017, 19, 3834-3845.	1.8	26
89	CRISPR-Cas, DNA Supercoiling, and Nucleoid-Associated Proteins. <i>Trends in Microbiology</i> , 2020, 28, 19-27.	3.5	26
90	Regulation of transcription by DNA supercoiling in <i>Mycoplasma genitalium</i> : global control in the smallest known self-replicating genome. <i>Molecular Microbiology</i> , 2011, 81, 302-304.	1.2	25

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91	Two highly related regulatory proteins, Shigella flexneri VirF and enterotoxigenic Escherichia coli Rns, have common and distinct regulatory properties. FEMS Microbiology Letters, 1998, 162, 303-309.	0.7	23
92	Characterization of the Detachable Rho-Dependent Transcription Terminator of the fimE Gene in Escherichia coli K-12. Journal of Bacteriology, 2005, 187, 8256-8266.	1.0	23
93	Interaction of the FimB Integrase with the fimS Invertible DNA Element in Escherichia coli In Vivo and In Vitro. Journal of Bacteriology, 2000, 182, 2953-2959.	1.0	22
94	Use of the stationary phase inducible promoters, spv and dps, to drive heterologous antigen expression in Salmonella vaccine strains. Vaccine, 2000, 18, 1298-1306.	1.7	22
95	Re-engineering cellular physiology by rewiring high-level global regulatory genes. Scientific Reports, 2016, 5, 17653.	1.6	22
96	Environmentally constrained mutation and adaptive evolution in Salmonella. Current Biology, 1999, 9, 1477-1481.	1.8	20
97	H-NS Silences <i>gfp</i> , the Green Fluorescent Protein Gene: ^{TCD} Is a Genetically Remastered <i>gfp</i> Gene with Reduced Susceptibility to H-NS-Mediated Transcription Silencing and with Enhanced Translation. Journal of Bacteriology, 2010, 192, 4790-4793.	1.0	20
98	Horizontally acquired homologues of the nucleoid-associated protein H-NS: implications for gene regulation. Molecular Microbiology, 2010, 75, 264-267.	1.2	19
99	Transmission of an Oxygen Availability Signal at the Salmonella enterica Serovar Typhimurium <i>fis</i> Promoter. PLoS ONE, 2013, 8, e84382.	1.1	19
100	Negative supercoiling of DNA by gyrase is inhibited in <i>Salmonella enterica</i> serovar Typhimurium during adaptation to acid stress. Molecular Microbiology, 2018, 107, 734-746.	1.2	18
101	Nucleotide sequence of the R26 chloramphenicol resistance determinant and identification of its gene product. Gene, 1986, 41, 349-353.	1.0	17
102	Thinking and decision making, bacterial style: Bacterial Neural Networks, Obernai, France, 7th-12th June 2002. Molecular Microbiology, 2003, 47, 583-593.	1.2	17
103	Thermal regulation of <i>fimA</i> , the Escherichia coli gene coding for the type 1 fimbrial subunit protein. FEMS Microbiology Letters, 1992, 78, 125-30.	0.7	17
104	Reciprocal Transcriptional and Posttranscriptional Growth-Phase-Dependent Expression of <i>sfh</i> , a Gene That Encodes a Parologue of the Nucleoid-Associated Protein H-NS. Journal of Bacteriology, 2006, 188, 7581-7591.	1.0	16
105	Regulatory integration of horizontally-transferred genes in bacteria. Frontiers in Bioscience, 2009, 14, 4103-12.	0.8	16
106	Global regulators and environmental adaptation in Gram-negative pathogens. Clinical Microbiology and Infection, 2009, 15, 47-50.	2.8	14
107	Autoregulated expression of the gene coding for the leucine-responsive protein, Lrp, a global regulator in Salmonella enterica serovar Typhimurium. Microbiology (United Kingdom), 2008, 154, 2008-2016.	0.7	13
108	Probing bacterial nucleoid structure with optical tweezers. BioEssays, 2007, 29, 212-216.	1.2	12

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109	Virulence Gene Regulation in <i>Shigella</i> . <i>EcoSal Plus</i> , 2004, 1, .	2.1	11
110	The Interplay between DNA Topology and Accessory Factors in Site-Specific Recombination in Bacteria and their Bacteriophages. <i>Science Progress</i> , 2016, 99, 420-437.	1.0	11
111	Bacterial pathogen gene regulation: a DNA-structure-centred view of a protein-dominated domain. <i>Clinical Science</i> , 2016, 130, 1165-1177.	1.8	11
112	Network Rewiring: Physiological Consequences of Reciprocally Exchanging the Physical Locations and Growth-Phase-Dependent Expression Patterns of the <i>Salmonella flexneri</i> and <i>dps</i> Genes. <i>MBio</i> , 2020, 11, .	1.8	11
113	Virulence gene deletion frequency is increased in <i>Shigella flexneri</i> following conjugation, transduction, and transformation. <i>FEMS Microbiology Letters</i> , 2006, 147, 163-172.	0.7	10
114	Small molecule signaling. <i>Current Opinion in Microbiology</i> , 2009, 12, 125-128.	2.3	10
115	Bacterial Chromatin. , 2010, , .		7
116	Integrating small molecule signalling and <i>HNS</i> antagonism in <i>Vibrio cholerae</i> , a bacterium with two chromosomes. <i>Molecular Microbiology</i> , 2015, 97, 612-615.	1.2	7
117	Consequences of producing DNA gyrase from a synthetic <i>gyrBA</i> operon in <i>Salmonella enterica</i> serovar Typhimurium. <i>Molecular Microbiology</i> , 2021, 115, 1410-1429.	1.2	7
118	Broad-scale redistribution of mRNA abundance and transcriptional machinery in response to growth rate in <i>Salmonella enterica</i> serovar Typhimurium. <i>Microbial Genomics</i> , 2017, 3, e000127.	1.0	6
119	The Virulence Plasmid of <i>Salmonella typhimurium</i> Contains an Autoregulated Gene, <i>rlgA</i> , That Codes for a Resolvase-like DNA Binding Protein. <i>Plasmid</i> , 2000, 44, 24-33.	0.4	5
120	The Virulence Plasmids of <i>Shigella flexneri</i> . <i>Microbiology Monographs</i> , 2009, , 151-170.	0.3	5
121	Transcription of the <i>Salmonella typhimurium</i> <i>spv</i> virulence locus is regulated negatively by the nucleoid-associated protein H-NS. , 0, .		5
122	Regulation of Transcription in Bacteria by DNA Supercoiling. , 2008, , 155-178.		4
123	The Evolution of Gene Regulatory Mechanisms in Bacteria. <i>Grand Challenges in Biology and Biotechnology</i> , 2018, , 125-152.	2.4	4
124	Reciprocally rewiring and repositioning the Integration Host Factor (IHF) subunit genes in <i>Salmonella enterica</i> serovar Typhimurium: impacts on physiology and virulence. <i>Microbial Genomics</i> , 2022, 8, .	1.0	3
125	A novel genetic locus determines in vivo B-Z DNA structural transitions in <i>Escherichia coli</i> . <i>Trends in Biochemical Sciences</i> , 1988, 13, 130.	3.7	2
126	Chromosome rearrangements induced by recombinant coliphage λ pplacMu. <i>Gene</i> , 1990, 94, 15-22.	1.0	2

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127	Regulation of Virulence Gene Expression in Bacterial Pathogens. , 2001, , 75-132.		2
128	H-NS and genomic bridge building: lessons from the human pathogen Salmonella Typhi. Microbiology (United Kingdom), 2009, 155, 2114-2115.	0.7	2
129	DNA TOPOLOGY AND ADAPTATION OF SALMONELLA TYPHIMURIUM TO AN INTRACELLULAR ENVIRONMENT. , 2001, , .		2
130	Bacterial Chromatin and Gene Regulation. , 2010, , 245-250.		1
131	Two highly related regulatory proteins, Shigella flexneri VirF and enterotoxigenic Escherichia coli Rns, have common and distinct regulatory properties. , 0, .		1
132	7.1 Introduction. Methods in Microbiology, 1998, 27, 345-347.	0.4	0
133	Whither microbial sciences?. Nature Reviews Microbiology, 2007, 5, 828-828.	13.6	0
134	Coordination of Bacterial Virulence Gene Expression. , 2015, , 315-335.		0
135	Editorial overview: Bacterial regulatory hierarchies and networks. Current Opinion in Microbiology, 2020, 55, iii-v.	2.3	0
136	Nucleus Organization of the Bacterial Nucleoid. , 2021, , 409-414.		0
137	Nucleoid Organization of Bacterial Chromosomes. , 2004, , 115-118.		0
138	Coordination of Gene Expression in Pathogenic Salmonella typhimurium. , 1993, , 51-62.		0
139	AgnÃ's Fouet â€“ departing Editor-in-Chief. Microbiology (United Kingdom), 2015, 161, 1149-1149.	0.7	0
140	Factors governing orthologous RpoD and H-NS evolution in Salmonella enterica Serovar Typhimurium and Escherichia coli. Access Microbiology, 2020, 2, .	0.2	0