

Graham C Walker

List of Publications by Year in descending order

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349
papers

32,862
citations

3731

89
h-index

5394

164
g-index

362
all docs

362
docs citations

362
times ranked

17566
citing authors

#	ARTICLE	IF	CITATIONS
1	Mutagenesis and inducible responses to deoxyribonucleic acid damage in <i>Escherichia coli</i> . <i>Microbiological Reviews</i> , 1984, 48, 60-93.	10.1	1,645
2	Mechanisms of DNA damage, repair, and mutagenesis. <i>Environmental and Molecular Mutagenesis</i> , 2017, 58, 235-263.	2.2	1,129
3	A genetic basis for <i>Pseudomonas aeruginosa</i> biofilm antibiotic resistance. <i>Nature</i> , 2003, 426, 306-310.	27.8	1,036
4	Mutagenesis and inducible responses to deoxyribonucleic acid damage in <i>Escherichia coli</i> . <i>Microbiological Reviews</i> , 1984, 48, 60-93.	10.1	895
5	The Y-Family of DNA Polymerases. <i>Molecular Cell</i> , 2001, 8, 7-8.	9.7	798
6	How rhizobial symbionts invade plants: the <i>Sinorhizobium</i> – <i>Medicago</i> model. <i>Nature Reviews Microbiology</i> , 2007, 5, 619-633.	28.6	781
7	Antibiotics induce redox-related physiological alterations as part of their lethality. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E2100-9.	7.1	698
8	Exopolysaccharide-deficient mutants of <i>Rhizobium meliloti</i> that form ineffective nodules.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1985, 82, 6231-6235.	7.1	596
9	DNA Repair and Mutagenesis. , 2005, , .		591
10	Inducible DNA Repair Systems. <i>Annual Review of Biochemistry</i> , 1985, 54, 425-457.	11.1	588
11	DNA-damaging agents stimulate gene expression at specific loci in <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1980, 77, 2819-2823.	7.1	527
12	Eukaryotic Translesion Polymerases and Their Roles and Regulation in DNA Damage Tolerance. <i>Microbiology and Molecular Biology Reviews</i> , 2009, 73, 134-154.	6.6	502
13	Succinoglycan Is Required for Initiation and Elongation of Infection Threads during Nodulation of Alfalfa by <i>Rhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 1998, 180, 5183-5191.	2.2	448
14	Oxidation of the Guanine Nucleotide Pool Underlies Cell Death by Bactericidal Antibiotics. <i>Science</i> , 2012, 336, 315-319.	12.6	400
15	Bactericidal Antibiotics Induce Toxic Metabolic Perturbations that Lead to Cellular Damage. <i>Cell Reports</i> , 2015, 13, 968-980.	6.4	393
16	RecA-mediated cleavage activates UmuD for mutagenesis: mechanistic relationship between transcriptional derepression and posttranslational activation.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1988, 85, 1816-1820.	7.1	383
17	Symbiotic mutants of <i>rhizobium meliloti</i> that uncouple plant from bacterial differentiation. <i>Cell</i> , 1985, 40, 869-877.	28.9	348
18	Molecular Determinants of a Symbiotic Chronic Infection. <i>Annual Review of Genetics</i> , 2008, 42, 413-441.	7.6	326

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19	Proteins required for ultraviolet light and chemical mutagenesis. <i>Journal of Molecular Biology</i> , 1983, 164, 175-192.	4.2	306
20	Enhancing tumor cell response to chemotherapy through nanoparticle-mediated codelivery of siRNA and cisplatin prodrug. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 18638-18643.	7.1	302
21	Biosynthesis of succinoglycan, a symbiotically important exopolysaccharide of <i>Rhizobium meliloti</i> . <i>Cell</i> , 1993, 74, 269-280.	28.9	296
22	A novel exopolysaccharide can function in place of the Calcofluor-binding exopolysaccharide in nodulation of alfalfa by <i>Rhizobium meliloti</i> . <i>Cell</i> , 1989, 56, 661-672.	28.9	295
23	The Sos Response: Recent Insights into umuDC-Dependent Mutagenesis and DNA Damage Tolerance. <i>Annual Review of Genetics</i> , 2000, 34, 479-497.	7.6	288
24	Inducibility of a gene product required for UV and chemical mutagenesis in <i>Escherichia coli</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1981, 78, 5749-5753.	7.1	278
25	General transduction in <i>Rhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 1984, 159, 120-124.	2.2	274
26	Degradation of carboxy-terminal-tagged cytoplasmic proteins by the <i>Escherichia coli</i> protease HflB (FtsH). <i>Genes and Development</i> , 1998, 12, 1348-1355.	5.9	255
27	<i>Rhizobium meliloti</i> mutants that fail to succinylate their Calcofluor-binding exopolysaccharide are defective in nodule invasion. <i>Cell</i> , 1987, 51, 579-587.	28.9	243
28	<i>Escherichia coli</i> dnaK null mutants are inviable at high temperature. <i>Journal of Bacteriology</i> , 1987, 169, 283-290.	2.2	241
29	New recA mutations that dissociate the various RecA protein activities in <i>Escherichia coli</i> provide evidence for an additional role for RecA protein in UV mutagenesis. <i>Journal of Bacteriology</i> , 1989, 171, 2415-2423.	2.2	240
30	Cellular defects caused by deletion of the <i>Escherichia coli</i> dnaK gene indicate roles for heat shock protein in normal metabolism. <i>Journal of Bacteriology</i> , 1989, 171, 2337-2346.	2.2	236
31	A single amino acid governs enhanced activity of DinB DNA polymerases on damaged templates. <i>Nature</i> , 2006, 439, 225-228.	27.8	227
32	A White-Box Machine Learning Approach for Revealing Antibiotic Mechanisms of Action. <i>Cell</i> , 2019, 177, 1649-1661.e9.	28.9	227
33	umuDC and mucAB operons whose products are required for UV light- and chemical-induced mutagenesis: UmuD, MucA, and LexA proteins share homology. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1985, 82, 4331-4335.	7.1	222
34	Unraveling the Physiological Complexities of Antibiotic Lethality. <i>Annual Review of Pharmacology and Toxicology</i> , 2015, 55, 313-332.	9.4	222
35	DnaK as a thermometer: threonine-199 is site of autophosphorylation and is critical for ATPase activity. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1991, 88, 9513-9517.	7.1	216
36	Expression of the <i>E. coli</i> uvrA gene is inducible. <i>Nature</i> , 1981, 289, 808-810.	27.8	210

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37	Genes needed for the modification, polymerization, export, and processing of succinoglycan by <i>Rhizobium meliloti</i> : a model for succinoglycan biosynthesis. <i>Journal of Bacteriology</i> , 1993, 175, 7045-7055.	2.2	197
38	Plasmid (pKM101)-mediated enhancement of repair and mutagenesis: Dependence on chromosomal genes in <i>Escherichia coli</i> K-12. <i>Molecular Genetics and Genomics</i> , 1977, 152, 93-103.	2.4	190
39	Alfalfa Root Nodule Invasion Efficiency Is Dependent on <i>Sinorhizobium meliloti</i> Polysaccharides. <i>Journal of Bacteriology</i> , 2000, 182, 4310-4318.	2.2	190
40	Changing the Culture of Science Education at Research Universities. <i>Science</i> , 2011, 331, 152-153.	12.6	188
41	Identification of plasmid (pKM101)-coded proteins involved in mutagenesis and UV resistance. <i>Nature</i> , 1982, 300, 278-281.	27.8	186
42	Differential response of the plant <i>Medicago truncatula</i> to its symbiont <i>Sinorhizobium meliloti</i> or an exopolysaccharide-deficient mutant. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 704-709.	7.1	185
43	Error-prone translesion synthesis mediates acquired chemoresistance. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 20792-20797.	7.1	183
44	Low molecular weight EPS II of <i>Rhizobium meliloti</i> allows nodule invasion in <i>Medicago sativa</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1996, 93, 8636-8641.	7.1	179
45	Detailed structural characterization of succinoglycan, the major exopolysaccharide of <i>Rhizobium meliloti</i> Rm1021. <i>Journal of Bacteriology</i> , 1994, 176, 1997-2002.	2.2	175
46	Managing DNA polymerases: Coordinating DNA replication, DNA repair, and DNA recombination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 8342-8349.	7.1	170
47	Genetic analysis of a cluster of genes required for synthesis of the calcofluor-binding exopolysaccharide of <i>Rhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 1988, 170, 4239-4248.	2.2	169
48	Characterization of a Novel Pyranopyridine Inhibitor of the AcrAB Efflux Pump of <i>Escherichia coli</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 722-733.	3.2	169
49	Hydroxyurea Induces Hydroxyl Radical-Mediated Cell Death in <i>Escherichia coli</i> . <i>Molecular Cell</i> , 2009, 36, 845-860.	9.7	168
50	A <i>Rhizobium meliloti</i> homolog of the <i>Escherichia coli</i> peptide-antibiotic transport protein SbmA is essential for bacteroid development. <i>Genes and Development</i> , 1993, 7, 1485-1497.	5.9	167
51	Exopolysaccharides of <i>Rhizobium</i> : synthesis, regulation and symbiotic function. <i>Trends in Genetics</i> , 1994, 10, 63-67.	6.7	165
52	<i>Rhizobium meliloti</i> mutants that overproduce the <i>R. meliloti</i> acidic calcofluor-binding exopolysaccharide. <i>Journal of Bacteriology</i> , 1988, 170, 4249-4256.	2.2	164
53	A LuxR Homolog Controls Production of Symbiotically Active Extracellular Polysaccharide II by <i>Sinorhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 2002, 184, 5067-5076.	2.2	164
54	Similar Requirements of a Plant Symbiont and a Mammalian Pathogen for Prolonged Intracellular Survival. <i>Science</i> , 2000, 287, 2492-2493.	12.6	162

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55	BluB cannibalizes flavin to form the lower ligand of vitamin B12. <i>Nature</i> , 2007, 446, 449-453.	27.8	160
56	Suppression of Rev3, the catalytic subunit of Pol η , sensitizes drug-resistant lung tumors to chemotherapy. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 20786-20791.	7.1	160
57	A model for a umuDC-dependent prokaryotic DNA damage checkpoint. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 9218-9223.	7.1	158
58	The critical mutagenic translesion DNA polymerase Rev1 is highly expressed during G2/M phase rather than S phase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 8971-8976.	7.1	158
59	Mutagenesis and repair deficiencies of <i>Escherichia coli</i> umuC mutants are suppressed by the plasmid pKM101. <i>Molecular Genetics and Genomics</i> , 1979, 172, 17-24.	2.4	147
60	Succinoglycan Production by <i>Rhizobium meliloti</i> Is Regulated through the ExoS-ChvI Two-Component Regulatory System. <i>Journal of Bacteriology</i> , 1998, 180, 20-26.	2.2	146
61	Mutations altering heat shock specific subunit of RNA polymerase suppress major cellular defects of <i>E. coli</i> mutants lacking the DnaK chaperone.. <i>EMBO Journal</i> , 1990, 9, 4027-4036.	7.8	140
62	Family of glycosyl transferases needed for the synthesis of succinoglycan by <i>Rhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 1993, 175, 7033-7044.	2.2	140
63	Mutagenesis and More: umuDC and the <i>Escherichia coli</i> SOS Response. <i>Genetics</i> , 1998, 148, 1599-1610.	2.9	140
64	<i>Rhizobium meliloti</i> exopolysaccharides: Synthesis and symbiotic function. <i>Gene</i> , 1996, 179, 141-146.	2.2	139
65	The 32-kilobase exp gene cluster of <i>Rhizobium meliloti</i> directing the biosynthesis of galactoglucan: genetic organization and properties of the encoded gene products. <i>Journal of Bacteriology</i> , 1997, 179, 1375-1384.	2.2	139
66	Y-family DNA polymerases in <i>Escherichia coli</i> . <i>Trends in Microbiology</i> , 2007, 15, 70-77.	7.7	137
67	The SOS Regulatory Network. <i>EcoSal Plus</i> , 2008, 3, .	5.4	134
68	Host plant peptides elicit a transcriptional response to control the <i>Sinorhizobium meliloti</i> cell cycle during symbiosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 3561-3566.	7.1	134
69	Chronic intracellular infection of alfalfa nodules by <i>Sinorhizobium meliloti</i> requires correct lipopolysaccharide core. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 3938-3943.	7.1	129
70	Conserved Bacterial RNase YbeY Plays Key Roles in 70S Ribosome Quality Control and 16S rRNA Maturation. <i>Molecular Cell</i> , 2013, 49, 427-438.	9.7	127
71	A Small Molecule Targeting Mutagenic Translesion Synthesis Improves Chemotherapy. <i>Cell</i> , 2019, 178, 152-159.e11.	28.9	126
72	Delta dnaK52 mutants of <i>Escherichia coli</i> have defects in chromosome segregation and plasmid maintenance at normal growth temperatures. <i>Journal of Bacteriology</i> , 1989, 171, 6030-6038.	2.2	124

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73	Conjugal transfer system of the IncN plasmid pKM101. <i>Journal of Bacteriology</i> , 1985, 161, 402-410.	2.2	123
74	[19] Genetic techniques in <i>Rhizobium meliloti</i> . <i>Methods in Enzymology</i> , 1991, 204, 398-418.	1.0	122
75	Structure of the Endonuclease Domain of MutL: Unlicensed to Cut. <i>Molecular Cell</i> , 2010, 39, 145-151.	9.7	122
76	Structural Characterization of the Symbiotically Important Low-Molecular-Weight Succinoglycan of <i>Sinorhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 1999, 181, 6788-6796.	2.2	120
77	Symbiotic loci of <i>Rhizobium meliloti</i> identified by random TnphoA mutagenesis. <i>Journal of Bacteriology</i> , 1988, 170, 4257-4265.	2.2	119
78	Localization of the plasmid (pKM101) gene(s) involved in recA + lexA +-dependent mutagenesis. <i>Molecular Genetics and Genomics</i> , 1980, 179, 289-297.	2.4	118
79	Dominant negative umuD mutations decreasing RecA-mediated cleavage suggest roles for intact UmuD in modulation of SOS mutagenesis.. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1990, 87, 7190-7194.	7.1	118
80	Cold sensitivity induced by overproduction of UmuDC in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 1985, 162, 155-161.	2.2	117
81	Functional organization of plasmid pKM101. <i>Journal of Bacteriology</i> , 1981, 145, 1310-1316.	2.2	108
82	Cell Cycle Control by the Master Regulator CtrA in <i>Sinorhizobium meliloti</i> . <i>PLoS Genetics</i> , 2015, 11, e1005232.	3.5	105
83	Genetic manipulations in <i>Rhizobium meliloti</i> utilizing two new transposon Tn5 derivatives. <i>Molecular Genetics and Genomics</i> , 1986, 204, 485-491.	2.4	104
84	The <i>Escherichia coli</i> SOS mutagenesis proteins UmuD and UmuD' interact physically with the replicative DNA polymerase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 12373-12378.	7.1	100
85	Î² Clamp Directs Localization of Mismatch Repair in <i>Bacillus subtilis</i> . <i>Molecular Cell</i> , 2008, 29, 291-301.	9.7	100
86	Biosynthetic control of molecular weight in the polymerization of the octasaccharide subunits of succinoglycan, a symbiotically important exopolysaccharide of <i>Rhizobium meliloti</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 13477-13482.	7.1	99
87	Similarity to peroxisomal-membrane protein family reveals that <i>Sinorhizobium</i> and <i>Brucella</i> BacA affect lipid-A fatty acids. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004, 101, 5012-5017.	7.1	99
88	UmuD and RecA Directly Modulate the Mutagenic Potential of the Y Family DNA Polymerase DinB. <i>Molecular Cell</i> , 2007, 28, 1058-1070.	9.7	99
89	Functional characterization of bacterial sRNAs using a network biology approach. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 15522-15527.	7.1	99
90	Structural Basis of Rev1-mediated Assembly of a Quaternary Vertebrate Translesion Polymerase Complex Consisting of Rev1, Heterodimeric Polymerase (Pol) Î¶, and Pol Î². <i>Journal of Biological Chemistry</i> , 2012, 287, 33836-33846.	3.4	98

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91	SOS-regulated proteins in translesion DNA synthesis and mutagenesis. Trends in Biochemical Sciences, 1995, 20, 416-420.	7.5	97
92	Role of <i>Escherichia coli</i> YbeY, a highly conserved protein, in rRNA processing. Molecular Microbiology, 2010, 78, 506-518.	2.5	97
93	The <i>Escherichia coli</i> polB gene, which encodes DNA polymerase II, is regulated by the SOS system. Journal of Bacteriology, 1990, 172, 6268-6273.	2.2	96
94	Roles for the transcription elongation factor NusA in both DNA repair and damage tolerance pathways in <i>Escherichia coli</i> . Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 15517-15522.	7.1	96
95	Isolation and characterization of Tn5 insertion mutations in the <i>lexA</i> gene of <i>Escherichia coli</i> . Journal of Bacteriology, 1983, 153, 1368-1378.	2.2	96
96	Exogenous suppression of the symbiotic deficiencies of <i>Rhizobium meliloti</i> <i>exo</i> mutants. Journal of Bacteriology, 1992, 174, 3403-3406.	2.2	95
97	The <i>Rhizobium meliloti</i> <i>exoK</i> gene and <i>prsD</i> / <i>prsE</i> / <i>exsH</i> genes are components of independent degradative pathways which contribute to production of low molecular weight succinoglycan. Molecular Microbiology, 1997, 25, 117-134.	2.5	92
98	Construction of an <i>Escherichia coli</i> K-12 <i>ada</i> deletion by gene replacement in a <i>recD</i> strain reveals a second methyltransferase that repairs alkylated DNA. Journal of Bacteriology, 1988, 170, 3294-3296.	2.2	91
99	<i>Sinorhizobium meliloti</i> <i>bluB</i> is necessary for production of 5,6-dimethylbenzimidazole, the lower ligand of B12. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 4634-4639.	7.1	91
100	Deficiency of a <i>Sinorhizobium meliloti</i> <i>bacA</i> Mutant in Alfalfa Symbiosis Correlates with Alteration of the Cell Envelope. Journal of Bacteriology, 2002, 184, 5625-5632.	2.2	89
101	Induction and autoregulation of <i>ada</i> , a positively acting element regulating the response of <i>Escherichia coli</i> K-12 to methylating agents. Journal of Bacteriology, 1985, 161, 888-895.	2.2	89
102	Structural studies of a novel exopolysaccharide produced by a mutant of <i>Rhizobium meliloti</i> strain Rm1021. Carbohydrate Research, 1990, 198, 305-312.	2.3	88
103	Visualization of Mismatch Repair in Bacterial Cells. Molecular Cell, 2001, 8, 1197-1206.	9.7	86
104	<i>Rhizobium meliloti</i> <i>exoG</i> and <i>exoJ</i> mutations affect the <i>exoX</i> - <i>exoY</i> system for modulation of exopolysaccharide production. Journal of Bacteriology, 1991, 173, 3776-3788.	2.2	85
105	Regulation of <i>Rhizobium meliloti</i> <i>exo</i> genes in free-living cells and in planta examined by using TnphoA fusions. Journal of Bacteriology, 1991, 173, 426-434.	2.2	85
106	Global analysis of cell cycle gene expression of the legume symbiont <i>Sinorhizobium meliloti</i> . Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 3217-3224.	7.1	85
107	The <i>muc</i> genes of pKM101 are induced by DNA damage. Journal of Bacteriology, 1983, 155, 1306-1315.	2.2	83
108	Rhizobial peptidase HrrP cleaves host-encoded signaling peptides and mediates symbiotic compatibility. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 15244-15249.	7.1	82

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109	Isolation and characterization of mutants of the plasmid pKM101 deficient in their ability to enhance mutagenesis and repair. <i>Journal of Bacteriology</i> , 1978, 133, 1203-1211.	2.2	82
110	groE mutants of <i>Escherichia coli</i> are defective in umuDC-dependent UV mutagenesis. <i>Journal of Bacteriology</i> , 1989, 171, 6117-6125.	2.2	80
111	Y-family DNA polymerases respond to DNA damage-independent inhibition of replication fork progression. <i>EMBO Journal</i> , 2006, 25, 868-879.	7.8	78
112	TtsI regulates symbiotic genes in <i>Rhizobium</i> species NGR234 by binding to <i>tts</i> boxes. <i>Molecular Microbiology</i> , 2008, 68, 736-748.	2.5	77
113	BacA, an ABC Transporter Involved in Maintenance of Chronic Murine Infections with <i>Mycobacterium tuberculosis</i> . <i>Journal of Bacteriology</i> , 2009, 191, 477-485.	2.2	76
114	Polymerase exchange on single DNA molecules reveals processivity clamp control of translesion synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 7647-7652.	7.1	76
115	The <i>exoR</i> gene of <i>Rhizobium meliloti</i> affects RNA levels of other <i>exo</i> genes but lacks homology to known transcriptional regulators. <i>Journal of Bacteriology</i> , 1991, 173, 3789-3794.	2.2	75
116	Importance of unusually modified lipid A in <i>Sinorhizobium</i> stress resistance and legume symbiosis. <i>Molecular Microbiology</i> , 2005, 56, 68-80.	2.5	74
117	Dimerization of the UmuD' protein in solution and its implications for regulation of SOS mutagenesis. <i>Nature Structural Biology</i> , 1997, 4, 979-982.	9.7	73
118	Identification and characterization of the <i>mutL</i> and <i>mutS</i> gene products of <i>Salmonella typhimurium</i> LT2. <i>Journal of Bacteriology</i> , 1985, 163, 1007-1015.	2.2	73
119	Striking Complexity of Lipopolysaccharide Defects in a Collection of <i>Sinorhizobium meliloti</i> Mutants. <i>Journal of Bacteriology</i> , 2003, 185, 3853-3862.	2.2	72
120	Regulation of <i>Escherichia coli</i> SOS mutagenesis by dimeric intrinsically disordered <i>umuD</i> gene products. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1152-1157.	7.1	71
121	Essential Role for the BacA Protein in the Uptake of a Truncated Eukaryotic Peptide in <i>Sinorhizobium meliloti</i> . <i>Journal of Bacteriology</i> , 2009, 191, 1519-1527.	2.2	71
122	Multifaceted Recognition of Vertebrate Rev1 by Translesion Polymerases Ψ and Ψ' . <i>Journal of Biological Chemistry</i> , 2012, 287, 26400-26408.	3.4	69
123	NMR Structure and Dynamics of the C-Terminal Domain from Human Rev1 and Its Complex with Rev1 Interacting Region of DNA Polymerase Ψ . <i>Biochemistry</i> , 2012, 51, 5506-5520.	2.5	69
124	Altering the conserved nucleotide binding motif in the <i>Salmonella typhimurium</i> MutS mismatch repair protein affects both its ATPase and mismatch binding activities. <i>EMBO Journal</i> , 1991, 10, 2707-15.	7.8	69
125	Mutations altering heat shock specific subunit of RNA polymerase suppress major cellular defects of <i>E. coli</i> mutants lacking the DnaK chaperone. <i>EMBO Journal</i> , 1990, 9, 4027-36.	7.8	69
126	DnaK mutants defective in ATPase activity are defective in negative regulation of the heat shock response: expression of mutant DnaK proteins results in filamentation. <i>Journal of Bacteriology</i> , 1994, 176, 764-780.	2.2	68

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127	Unconventional Ubiquitin Recognition by the Ubiquitin-Binding Motif within the Y Family DNA Polymerases $\hat{1}$ and Rev1. <i>Molecular Cell</i> , 2010, 37, 408-417.	9.7	68
128	The <i>DivJ</i> , <i>CbrA</i> and <i>PleC</i> system controls <i>DivK</i> phosphorylation and symbiosis in <i>Sinorhizobium meliloti</i> . <i>Molecular Microbiology</i> , 2013, 90, 54-71.	2.5	68
129	Regulation of damage-inducible genes in <i>Escherichia coli</i> . <i>Journal of Molecular Biology</i> , 1982, 160, 445-457.	4.2	67
130	A highly conserved protein of unknown function in <i>Sinorhizobium meliloti</i> affects sRNA regulation similar to Hfq. <i>Nucleic Acids Research</i> , 2011, 39, 4691-4708.	14.5	67
131	Exo-Oligosaccharides of <i>Rhizobium</i> sp. Strain NGR234 Are Required for Symbiosis with Various Legumes. <i>Journal of Bacteriology</i> , 2006, 188, 6168-6178.	2.2	65
132	Comparison of Responses to Double-Strand Breaks between <i>Escherichia coli</i> and <i>Bacillus subtilis</i> Reveals Different Requirements for SOS Induction. <i>Journal of Bacteriology</i> , 2009, 191, 1152-1161.	2.2	65
133	The Transcription Elongation Factor NusA Is Required for Stress-Induced Mutagenesis in <i>Escherichia coli</i> . <i>Current Biology</i> , 2010, 20, 80-85.	3.9	65
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