

Daniel N Wilson

List of Publications by Year in descending order

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Version: 2024-02-01

185
papers

14,234
citations

14655

66
h-index

24982

109
g-index

221
all docs

221
docs citations

221
times ranked

11758
citing authors

#	ARTICLE	IF	CITATIONS
1	Putting the antibiotics chloramphenicol and linezolid into context. <i>Nature Structural and Molecular Biology</i> , 2022, 29, 79-81.	8.2	1
2	Structural basis for PoxA-mediated resistance to phenicol and oxazolidinone antibiotics. <i>Nature Communications</i> , 2022, 13, 1860.	12.8	25
3	Total Synthesis and Biological Evaluation of Paenilamicins from the Honey Bee Pathogen <i>Paenibacillus larvae</i> . <i>Journal of the American Chemical Society</i> , 2022, 144, 288-296.	13.7	6
4	The cyclic octapeptide antibiotic argyrin B inhibits translation by trapping EF-G on the ribosome during translocation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2114214119.	7.1	8
5	Structural Basis for Bacterial Ribosome-Associated Quality Control by RqcH and RqcP. <i>Molecular Cell</i> , 2021, 81, 115-126.e7.	9.7	41
6	Yeast translation elongation factor eEF3 promotes late stages of tRNA translocation. <i>EMBO Journal</i> , 2021, 40, e106449.	7.8	19
7	Ribosome Rescue Pathways in Bacteria. <i>Frontiers in Microbiology</i> , 2021, 12, 652980.	3.5	46
8	Structure of Gcn1 bound to stalled and colliding 80S ribosomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	79
9	Context-specific action of macrolide antibiotics on the eukaryotic ribosome. <i>Nature Communications</i> , 2021, 12, 2803.	12.8	18
10	Repurposing tRNAs for nonsense suppression. <i>Nature Communications</i> , 2021, 12, 3850.	12.8	22
11	Structural basis of ABCF-mediated resistance to pleuromutilin, lincosamide, and streptogramin A antibiotics in Gram-positive pathogens. <i>Nature Communications</i> , 2021, 12, 3577.	12.8	40
12	RqcH and RqcP catalyze processive poly-alanine synthesis in a reconstituted ribosome-associated quality control system. <i>Nucleic Acids Research</i> , 2021, 49, 8355-8369.	14.5	11
13	Structural and mechanistic basis for translation inhibition by macrolide and ketolide antibiotics. <i>Nature Communications</i> , 2021, 12, 4466.	12.8	43
14	Structural Basis for Regulation of the Opposing (p)ppGpp Synthetase and Hydrolase within the Stringent Response Orchestrator Rel. <i>Cell Reports</i> , 2020, 32, 108157.	6.4	39
15	Bifunctional Nitrone-Conjugated Secondary Metabolite Targeting the Ribosome. <i>Journal of the American Chemical Society</i> , 2020, 142, 18369-18377.	13.7	7
16	Coupling of 5S RNP rotation with maturation of functional centers during large ribosomal subunit assembly. <i>Nature Communications</i> , 2020, 11, 3751.	12.8	24
17	Peptide Inhibitors of Bacterial Protein Synthesis with Broad Spectrum and SbmA-Independent Bactericidal Activity against Clinical Pathogens. <i>Journal of Medicinal Chemistry</i> , 2020, 63, 9590-9602.	6.4	24
18	Characterization of Cetacean Proline-Rich Antimicrobial Peptides Displaying Activity against ESKAPE Pathogens. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7367.	4.1	8

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19	Mechanism of ribosome rescue by alternative ribosome-rescue factor B. <i>Nature Communications</i> , 2020, 11, 4106.	12.8	26
20	The 23S Ribosomal RNA From <i>Pyrococcus furiosus</i> Is Circularly Permuted. <i>Frontiers in Microbiology</i> , 2020, 11, 582022.	3.5	9
21	The alarmones (p)ppGpp are part of the heat shock response of <i>Bacillus subtilis</i> . <i>PLoS Genetics</i> , 2020, 16, e1008275.	3.5	52
22	Tetracenomycin X inhibits translation by binding within the ribosomal exit tunnel. <i>Nature Chemical Biology</i> , 2020, 16, 1071-1077.	8.0	43
23	Target protection as a key antibiotic resistance mechanism. <i>Nature Reviews Microbiology</i> , 2020, 18, 637-648.	28.6	100
24	Proline-Rich Peptides with Improved Antimicrobial Activity against <i>E. coli</i> , <i>K. pneumoniae</i> , and <i>A. baumannii</i> . <i>ChemMedChem</i> , 2019, 14, 2025-2033.	3.2	35
25	Intracellular Antimicrobial Peptides Targeting the Protein Synthesis Machinery. <i>Advances in Experimental Medicine and Biology</i> , 2019, 1117, 73-89.	1.6	63
26	The Natural Product Elegaphenone Potentiates Antibiotic Effects against <i>Pseudomonas aeruginosa</i> . <i>Angewandte Chemie - International Edition</i> , 2019, 58, 8581-8584.	13.8	13
27	Der Naturstoff Elegaphenon verstärkt antibiotische Effekte gegen <i>Pseudomonas aeruginosa</i> . <i>Angewandte Chemie</i> , 2019, 131, 8670-8674.	2.0	2
28	A role for the <i>Saccharomyces cerevisiae</i> ABCF protein New1 in translation termination/recycling. <i>Nucleic Acids Research</i> , 2019, 47, 8807-8820.	14.5	26
29	Release factor-dependent ribosome rescue by BrfA in the Gram-positive bacterium <i>Bacillus subtilis</i> . <i>Nature Communications</i> , 2019, 10, 5397.	12.8	32
30	The Dolphin Proline-Rich Antimicrobial Peptide Tur1A Inhibits Protein Synthesis by Targeting the Bacterial Ribosome. <i>Cell Chemical Biology</i> , 2018, 25, 530-539.e7.	5.2	90
31	Hierarchical recruitment of ribosomal proteins and assembly factors remodels nucleolar pre-60S ribosomes. <i>Journal of Cell Biology</i> , 2018, 217, 2503-2518.	5.2	33
32	Structural basis for (p)ppGpp-mediated inhibition of the GTPase RbgA. <i>Journal of Biological Chemistry</i> , 2018, 293, 19699-19709.	3.4	41
33	Structure of a hibernating 100S ribosome reveals an inactive conformation of the ribosomal protein S1. <i>Nature Microbiology</i> , 2018, 3, 1115-1121.	13.3	92
34	Structural basis for antibiotic resistance mediated by the <i>Bacillus subtilis</i> ABCF ATPase VmlR. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 8978-8983.	7.1	78
35	Fragments of the Nonlytic Proline-Rich Antimicrobial Peptide Bac5 Kill <i>Escherichia coli</i> Cells by Inhibiting Protein Synthesis. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	3.2	44
36	Visualization of translation termination intermediates trapped by the Apidaecin137 peptide during RF3-mediated recycling of RF1. <i>Nature Communications</i> , 2018, 9, 3053.	12.8	48

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37	The Mechanisms of Action of Ribosome-Targeting Peptide Antibiotics. <i>Frontiers in Molecular Biosciences</i> , 2018, 5, 48.	3.5	84
38	Totalsynthese und Strukturkorrektur des antibiotisch wirksamen Tetrapeptids GE81112A. <i>Angewandte Chemie</i> , 2018, 130, 12334-12338.	2.0	3
39	Total Synthesis and Structural Revision of the Antibiotic Tetrapeptide GE81112A. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 12157-12161.	13.8	12
40	Structure of the <i>Bacillus subtilis</i> hibernating 100S ribosome reveals the basis for 70S dimerization. <i>EMBO Journal</i> , 2017, 36, 2061-2072.	7.8	74
41	Structural basis for ArfA-mediated translation termination on mRNAs lacking stop codons. <i>Nature</i> , 2017, 541, 546-549.	27.8	39
42	Structural Basis for Ribosome Rescue in Bacteria. <i>Trends in Biochemical Sciences</i> , 2017, 42, 669-680.	7.5	53
43	Proline-rich antimicrobial peptides targeting protein synthesis. <i>Natural Product Reports</i> , 2017, 34, 702-711.	10.3	132
44	Structural Basis for Polyproline-Mediated Ribosome Stalling and Rescue by the Translation Elongation Factor EF-P. <i>Molecular Cell</i> , 2017, 68, 515-527.e6.	9.7	118
45	An antimicrobial peptide that inhibits translation by trapping release factors on the ribosome. <i>Nature Structural and Molecular Biology</i> , 2017, 24, 752-757.	8.2	123
46	Myticalins: A Novel Multigenic Family of Linear, Cationic Antimicrobial Peptides from Marine Mussels (<i>Mytilus</i> spp.). <i>Marine Drugs</i> , 2017, 15, 261.	4.6	54
47	The force-sensing peptide VemP employs extreme compaction and secondary structure formation to induce ribosomal stalling. <i>ELife</i> , 2017, 6, .	6.0	81
48	Wnt/ β -catenin and LIF/Stat3 signaling pathways converge on Sp5 to promote mouse embryonic stem cell self-renewal. <i>Journal of Cell Science</i> , 2016, 129, 269-76.	2.0	43
49	Deciphering the Translation Initiation Factor 5A Modification Pathway in Halophilic Archaea. <i>Archaea</i> , 2016, 2016, 1-14.	2.3	24
50	Cryo-EM structure of the spinach chloroplast ribosome reveals the location of plastid-specific ribosomal proteins and extensions. <i>Nucleic Acids Research</i> , 2016, 45, gkw1272.	14.5	33
51	A combined cryo-EM and molecular dynamics approach reveals the mechanism of ErmBL-mediated translation arrest. <i>Nature Communications</i> , 2016, 7, 12026.	12.8	103
52	ErmBL Translation on the Ribosome in the Presence of Erythromycin is Stalled by Inhibition of Peptide Bond Formation. <i>Biophysical Journal</i> , 2016, 110, 234a.	0.5	0
53	The stringent factor RelA adopts an open conformation on the ribosome to stimulate ppGpp synthesis. <i>Nucleic Acids Research</i> , 2016, 44, 6471-6481.	14.5	129
54	Ribosomes Structure and Mechanisms in Regulation of Protein Synthesis Part I. <i>Journal of Molecular Biology</i> , 2016, 428, 2133.	4.2	0

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55	The ABC of Ribosome-Related Antibiotic Resistance. <i>MBio</i> , 2016, 7, .	4.1	62
56	Bacterial Protein Synthesis as a Target for Antibiotic Inhibition. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2016, 6, a025361.	6.2	94
57	Editorial. <i>Journal of Molecular Biology</i> , 2016, 428, 3557.	4.2	0
58	Knud Hermann Nierhaus 1941â€“2016. <i>Nature Structural and Molecular Biology</i> , 2016, 23, 503-504.	8.2	0
59	Structures of the orthosomycin antibiotics avilamycin and evernimicin in complex with the bacterial 70S ribosome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 7527-7532.	7.1	45
60	Structure of the hypusinylated eukaryotic translation factor eIF-5A bound to the ribosome. <i>Nucleic Acids Research</i> , 2016, 44, 1944-1951.	14.5	106
61	Stall no more at polyproline stretches with the translation elongation factors EFâ€P and IFâ€5A. <i>Molecular Microbiology</i> , 2016, 99, 219-235.	2.5	70
62	Translation regulation via nascent polypeptide-mediated ribosome stalling. <i>Current Opinion in Structural Biology</i> , 2016, 37, 123-133.	5.7	137
63	Structure of the mammalian antimicrobial peptide Bac7(1â€“16) bound within the exit tunnel of a bacterial ribosome. <i>Nucleic Acids Research</i> , 2016, 44, 2429-2438.	14.5	89
64	Blast from the Past: Reassessing Forgotten Translation Inhibitors, Antibiotic Selectivity, and Resistance Mechanisms to Aid Drug Development. <i>Molecular Cell</i> , 2016, 61, 3-14.	9.7	60
65	Wnt/ β -catenin and LIFâ€“Stat3 signaling pathways converge on Sp5 to promote mouse embryonic stem cell self-renewal. <i>Development (Cambridge)</i> , 2016, 143, e1.1-e1.1.	2.5	1
66	Distinct tRNA Accommodation Intermediates Observed on the Ribosome with the Antibiotics Hygromycin A and A201A. <i>Molecular Cell</i> , 2015, 58, 832-844.	9.7	79
67	Structural basis for the interaction of protein S1 with the Escherichia coli ribosome. <i>Nucleic Acids Research</i> , 2015, 43, 661-673.	14.5	56
68	Arginine-rhamnosylation as new strategy to activate translation elongation factor P. <i>Nature Chemical Biology</i> , 2015, 11, 266-270.	8.0	116
69	Cryo-EM structure of the tetracycline resistance protein TetM in complex with a translating ribosome at 3.9-Å... resolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 5401-5406.	7.1	58
70	The proline-rich antimicrobial peptide Onc112 inhibits translation by blocking and destabilizing the initiation complex. <i>Nature Structural and Molecular Biology</i> , 2015, 22, 470-475.	8.2	148
71	Structure of the Bacillus subtilis 70S ribosome reveals the basis for species-specific stalling. <i>Nature Communications</i> , 2015, 6, 6941.	12.8	105
72	Translational arrest by a prokaryotic signal recognition particle is mediated by RNA interactions. <i>Nature Structural and Molecular Biology</i> , 2015, 22, 767-773.	8.2	29

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73	Entropic Contribution of Elongation Factor P to Proline Positioning at the Catalytic Center of the Ribosome. <i>Journal of the American Chemical Society</i> , 2015, 137, 12997-13006.	13.7	88
74	Antimicrobial peptides target ribosomes. <i>Oncotarget</i> , 2015, 6, 16826-16827.	1.8	4
75	A Conserved Proline Triplet in Val-tRNA Synthetase and the Origin of Elongation Factor P. <i>Cell Reports</i> , 2014, 9, 476-483.	6.4	41
76	Translational stalling at polyproline stretches is modulated by the sequence context upstream of the stall site. <i>Nucleic Acids Research</i> , 2014, 42, 10711-10719.	14.5	88
77	Entrapment of DNA in an intersubunit tunnel system of a single-stranded DNA-binding protein. <i>Nucleic Acids Research</i> , 2014, 42, 6698-6708.	14.5	15
78	A new system for naming ribosomal proteins. <i>Current Opinion in Structural Biology</i> , 2014, 24, 165-169.	5.7	481
79	Ribosome Rescue, Nearing the End. <i>Cell</i> , 2014, 156, 866-867.	28.9	2
80	Ribosome-targeting antibiotics and mechanisms of bacterial resistance. <i>Nature Reviews Microbiology</i> , 2014, 12, 35-48.	28.6	790
81	Insights into the Mode of Action of Novel Fluoroketolides, Potent Inhibitors of Bacterial Protein Synthesis. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 472-480.	3.2	17
82	Drug Sensing by the Ribosome Induces Translational Arrest via Active Site Perturbation. <i>Molecular Cell</i> , 2014, 56, 446-452.	9.7	104
83	Molecular basis for erythromycin-dependent ribosome stalling during translation of the ErmBL leader peptide. <i>Nature Communications</i> , 2014, 5, 3501.	12.8	115
84	Tetracycline antibiotics and resistance mechanisms. <i>Biological Chemistry</i> , 2014, 395, 559-575.	2.5	324
85	The bacterial translation stress response. <i>FEMS Microbiology Reviews</i> , 2014, 38, 1172-1201.	8.6	165
86	Structures of Nascent Polypeptide Chain-Dependent-Stalled Ribosome Complexes. , 2014, , 45-59.		1
87	Translation Elongation Factor EF-P Alleviates Ribosome Stalling at Polyproline Stretches. <i>Science</i> , 2013, 339, 82-85.	12.6	393
88	Structural basis for potent inhibitory activity of the antibiotic tigecycline during protein synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 3812-3816.	7.1	152
89	Nascent peptides that block protein synthesis in bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E878-87.	7.1	137
90	Structures of the human and Drosophila 80S ribosome. <i>Nature</i> , 2013, 497, 80-85.	27.8	474

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91	On the use of the antibiotic chloramphenicol to target polypeptide chain mimics to the ribosomal exit tunnel. <i>Biochimie</i> , 2013, 95, 1765-1772.	2.6	14
92	Amythiamicin and Related Thiopeptides as Inhibitors of the Bacterial Elongation Factor EF-Tu: Modification of the Amino Acid at Carbon Atom C2 of Ring C Dramatically Influences Activity. <i>ChemMedChem</i> , 2013, 8, 1954-1962.	3.2	18
93	Promiscuous behaviour of archaeal ribosomal proteins: Implications for eukaryotic ribosome evolution. <i>Nucleic Acids Research</i> , 2013, 41, 1284-1293.	14.5	59
94	Distinct XPPX sequence motifs induce ribosome stalling, which is rescued by the translation elongation factor EF-P. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 15265-15270.	7.1	167
95	The DARC site: a database of aligned ribosomal complexes. <i>Nucleic Acids Research</i> , 2012, 40, D495-D500.	14.5	7
96	Target- and Resistance-Based Mechanistic Studies with TP-434, a Novel Fluorocycline Antibiotic. <i>Antimicrobial Agents and Chemotherapy</i> , 2012, 56, 2559-2564.	3.2	132
97	Structural basis for TetM-mediated tetracycline resistance. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 16900-16905.	7.1	151
98	Lys34 of translation elongation factor EF-P is hydroxylated by YfcM. <i>Nature Chemical Biology</i> , 2012, 8, 695-697.	8.0	81
99	Mechanisms of SecM-Mediated Stalling in the Ribosome. <i>Biophysical Journal</i> , 2012, 103, 331-341.	0.5	82
100	The Structure and Function of the Eukaryotic Ribosome. <i>Cold Spring Harbor Perspectives in Biology</i> , 2012, 4, a011536-a011536.	5.5	330
101	Peptides in the Ribosomal Tunnel Talk Back. <i>Molecular Cell</i> , 2011, 41, 247-248.	9.7	7
102	Proteomic Characterization of Archaeal Ribosomes Reveals the Presence of Novel Archaeal-Specific Ribosomal Proteins. <i>Journal of Molecular Biology</i> , 2011, 405, 1215-1232.	4.2	28
103	Antibiotic-induced ribosomal assembly defects result from changes in the synthesis of ribosomal proteins. <i>Molecular Microbiology</i> , 2011, 80, 54-67.	2.5	31
104	On the specificity of antibiotics targeting the large ribosomal subunit. <i>Annals of the New York Academy of Sciences</i> , 2011, 1241, 1-16.	3.8	57
105	The ribosomal tunnel as a functional environment for nascent polypeptide folding and translational stalling. <i>Current Opinion in Structural Biology</i> , 2011, 21, 274-282.	5.7	179
106	Differential Effects of Thiopeptide and Orthosomycin Antibiotics on Translational GTPases. <i>Chemistry and Biology</i> , 2011, 18, 589-600.	6.0	37
107	Molecular Basis for the Selectivity of Antituberculosis Compounds Capreomycin and Viomycin. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 4712-4717.	3.2	36
108	SecM-Stalled Ribosomes Adopt an Altered Geometry at the Peptidyl Transferase Center. <i>PLoS Biology</i> , 2011, 9, e1000581.	5.6	132

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109	Nascent polypeptide chains within the ribosomal tunnel analyzed by cryo-EM. , 2011, , 393-404.		0
110	Localization of eukaryote-specific ribosomal proteins in a 5.5-Å cryo-EM map of the 80S eukaryotic ribosome. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 19754-19759.	7.1	122
111	Interplay between the Ribosomal Tunnel, Nascent Chain, and Macrolides Influences Drug Inhibition. Chemistry and Biology, 2010, 17, 504-514.	6.0	94
112	Probing Translation with Small-Molecule Inhibitors. Chemistry and Biology, 2010, 17, 633-645.	6.0	42
113	±-Helical nascent polypeptide chains visualized within distinct regions of the ribosomal exit tunnel. Nature Structural and Molecular Biology, 2010, 17, 313-317.	8.2	187
114	Head swivel on the ribosome facilitates translocation by means of intra-subunit tRNA hybrid sites. Nature, 2010, 468, 713-716.	27.8	336
115	Cryo-EM structure and rRNA model of a translating eukaryotic 80S ribosome at 5.5-Å resolution. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 19748-19753.	7.1	196
116	PSRP1 Is Not a Ribosomal Protein, but a Ribosome-binding Factor That Is Recycled by the Ribosome-recycling Factor (RRF) and Elongation Factor G (EF-G). Journal of Biological Chemistry, 2010, 285, 4006-4014.	3.4	66
117	Structural Basis for Translational Stalling by Human Cytomegalovirus and Fungal Arginine Attenuator Peptide. Molecular Cell, 2010, 40, 138-146.	9.7	106
118	The E Site and Its Importance for Improving Accuracy and Preventing Frameshifts. Nucleic Acids and Molecular Biology, 2010, , 345-362.	0.2	2
119	Distinct Mode of Interaction of a Novel Ketolide Antibiotic That Displays Enhanced Antimicrobial Activity. Antimicrobial Agents and Chemotherapy, 2009, 53, 1411-1419.	3.2	15
120	The Final Step of Hygromycin A Biosynthesis, Oxidation of C-5 ³ -Dihydrohygromycin A, Is Linked to a Putative Proton Gradient-Dependent Efflux. Antimicrobial Agents and Chemotherapy, 2009, 53, 5163-5172.	3.2	6
121	Non-Hydrolyzable RNA-Peptide Conjugates: A Powerful Advance in the Synthesis of Mimics for 3 ² -Peptidyl tRNA Termini. Angewandte Chemie - International Edition, 2009, 48, 4056-4060.	13.8	38
122	Identification of Distinct Thiopeptide-Antibiotic Precursor Lead Compounds Using Translation Machinery Assays. Chemistry and Biology, 2009, 16, 1087-1096.	6.0	24
123	Biosynthesis of the Aminocyclitol Subunit of Hygromycin A in Streptomyces hygroscopicus NRRL 2388. Chemistry and Biology, 2009, 16, 1180-1189.	6.0	21
124	SnapShot: Antibiotic Inhibition of Protein Synthesis I. Cell, 2009, 138, 1248-1248.e1.	28.9	29
125	Enhanced SnapShot: Antibiotic Inhibition of Protein Synthesis II. Cell, 2009, 139, 212-212.e1.	28.9	20
126	Time-Resolved Binding of Azithromycin to Escherichia coli Ribosomes. Journal of Molecular Biology, 2009, 385, 1179-1192.	4.2	31

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127	Less Is More for Leaderless mRNA Translation. <i>Molecular Cell</i> , 2009, 33, 141-142.	9.7	2
128	Structural Insight into Nascent Polypeptide Chain-Mediated Translational Stalling. <i>Science</i> , 2009, 326, 1412-1415.	12.6	263
129	The A-Z of bacterial translation inhibitors. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2009, 44, 393-433.	5.2	273
130	The Role of 23S Ribosomal RNA Residue A2451 in Peptide Bond Synthesis Revealed by Atomic Mutagenesis. <i>Chemistry and Biology</i> , 2008, 15, 485-492.	6.0	88
131	A new tRNA intermediate revealed on the ribosome during EF4-mediated back-translocation. <i>Nature Structural and Molecular Biology</i> , 2008, 15, 910-915.	8.2	65
132	New Features of the Ribosome and Ribosomal Inhibitors: Non-Enzymatic Recycling, Misreading and Back-Translocation. <i>Journal of Molecular Biology</i> , 2008, 380, 193-205.	4.2	48
133	Translational Regulation via L11: Molecular Switches on the Ribosome Turned On and Off by Thiostrepton and Micrococcin. <i>Molecular Cell</i> , 2008, 30, 26-38.	9.7	269
134	Shine-Dalgarno interaction prevents incorporation of noncognate amino acids at the codon following the AUG. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 10715-10720.	7.1	24
135	The oxazolidinone antibiotics perturb the ribosomal peptidyl-transferase center and effect tRNA positioning. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 13339-13344.	7.1	285
136	Cryo-EM study of the spinach chloroplast ribosome reveals the structural and functional roles of plastid-specific ribosomal proteins. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 19315-19320.	7.1	101
137	Structural Basis for Interaction of the Ribosome with the Switch Regions of GTP-Bound Elongation Factors. <i>Molecular Cell</i> , 2007, 25, 751-764.	9.7	168
138	The Oxazolidinone Class of Drugs Find Their Orientation on the Ribosome. <i>Molecular Cell</i> , 2007, 26, 460-462.	9.7	11
139	Structural Aspects of RbfA Action during Small Ribosomal Subunit Assembly. <i>Molecular Cell</i> , 2007, 28, 434-445.	9.7	90
140	The Weird and Wonderful World of Bacterial Ribosome Regulation. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2007, 42, 187-219.	5.2	186
141	A Snapshot of the 30S Ribosomal Subunit Capturing mRNA via the Shine-Dalgarno Interaction. <i>Structure</i> , 2007, 15, 289-297.	3.3	94
142	EF-G-Dependent GTPase on the Ribosome. Conformational Change and Fusidic Acid Inhibition. <i>Biochemistry</i> , 2006, 45, 2504-2514.	2.5	73
143	The Highly Conserved LepA Is a Ribosomal Elongation Factor that Back-Translocates the Ribosome. <i>Cell</i> , 2006, 127, 721-733.	28.9	192
144	2P594 The antibiotic kasugamycin mimics mRNA nucleotides to destabilize tRNA binding and inhibit canonical translation initiation (55. Drug design and delivery, Poster Session, Abstract, Meeting) Tj ETQq0 0 0 rgBT /Ovarlock 10 Tf 50 57 T		

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145	The antibiotic kasugamycin mimics mRNA nucleotides to destabilize tRNA binding and inhibit canonical translation initiation. <i>Nature Structural and Molecular Biology</i> , 2006, 13, 871-878.	8.2	116
146	The E-site story: the importance of maintaining two tRNAs on the ribosome during protein synthesis. <i>Cellular and Molecular Life Sciences</i> , 2006, 63, 2725-2737.	5.4	58
147	The Mechanism of Recoding in Pro- and Eukaryotes. , 2006, , 397-428.		2
148	Regulation of Ribosome Biosynthesis in <i>Escherichia coli</i> . , 2006, , 429-448.		1
149	Antibiotics and the Inhibition of Ribosome Function. , 2006, , 449-527.		9
150	The Work of Chaperones. , 2006, , 529-562.		3
151	Structure of the Ribosome. , 2006, , 53-84.		2
152	tRNA and Synthetases. , 2006, , 145-184.		0
153	The Elongation Cycle. , 2006, , 323-366.		1
154	Termination and Ribosome Recycling. , 2006, , 367-395.		1
155	On the Mechanism of Action of 9-O-Arylalkyloxime Derivatives of 6-O-Mycaminylosylonolide, a New Class of 16-Membered Macrolide Antibiotics. <i>Molecular Pharmacology</i> , 2006, 70, 1271-1280.	2.3	19
156	The binding mode of the trigger factor on the ribosome: Implications for protein folding and SRP interaction. <i>FASEB Journal</i> , 2006, 20, A965.	0.5	0
157	Microarray Analysis of Postictal Transcriptional Regulation of Neuropeptides. <i>Journal of Molecular Neuroscience</i> , 2005, 25, 285-298.	2.3	48
158	RelBE or not to be. <i>Nature Structural and Molecular Biology</i> , 2005, 12, 282-284.	8.2	13
159	X-ray crystallography study on ribosome recycling: the mechanism of binding and action of RRF on the 50S ribosomal subunit. <i>EMBO Journal</i> , 2005, 24, 251-260.	7.8	104
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