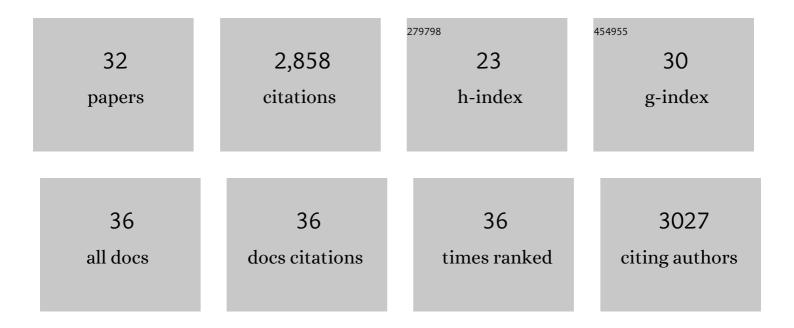
## Sean R Connell

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Histone mRNA is subject to 3′ uridylation and reâ€adenylation in <i>Aspergillus nidulans</i> . Molecular Microbiology, 2021, 115, 238-254.	2.5	3
2	A conserved rRNA switch is central to decoding site maturation on the small ribosomal subunit. Science Advances, 2021, 7, .	10.3	23
3	Backbone and sidechain NMR assignments for the ribosome maturation factor RbfA from Escherichia coli. Biomolecular NMR Assignments, 2020, 14, 317-321.	0.8	1
4	Backbone and sidechain NMR assignments for the ribosome maturation factor RimP from Escherichia coli. Biomolecular NMR Assignments, 2020, 14, 189-193.	0.8	4
5	Structure of a 30S pre-initiation complex stalled by GE81112 reveals structural parallels in bacterial and eukaryotic protein synthesis initiation pathways. Nucleic Acids Research, 2017, 45, gkw1251.	14.5	23
6	RsgA couples the maturation state of the 30S ribosomal decoding center to activation of its GTPase pocket. Nucleic Acids Research, 2017, 45, 6945-6959.	14.5	29
7	The Novel Aminomethylcycline Omadacycline Has High Specificity for the Primary Tetracycline-Binding Site on the Bacterial Ribosome. Antibiotics, 2016, 5, 32.	3.7	33
8	Inhibition of translation initiation complex formation by GE81112 unravels a 16S rRNA structural switch involved in P-site decoding. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E2286-95.	7.1	28
9	Crystallographic characterization of the ribosomal binding site and molecular mechanism of action of Hygromycin A. Nucleic Acids Research, 2015, 43, gkv975.	14.5	15
10	Structural Characterization of an Alternative Mode of Tigecycline Binding to the Bacterial Ribosome. Antimicrobial Agents and Chemotherapy, 2015, 59, 2849-2854.	3.2	32
11	Solid-state NMR enhanced by dynamic nuclear polarization as a novel tool for ribosome structural biology. Journal of Biomolecular NMR, 2013, 56, 85-93.	2.8	59
12	Structure and Conformational Variability of the Mycobacterium tuberculosis Fatty Acid Synthase Multienzyme Complex. Structure, 2013, 21, 1251-1257.	3.3	39
13	Fatty acid synthase: insights in the substrate shuttling mechanism by cryo-EM. Acta Crystallographica Section A: Foundations and Advances, 2012, 68, s31-s31.	0.3	0
14	Head swivel on the ribosome facilitates translocation by means of intra-subunit tRNA hybrid sites. Nature, 2010, 468, 713-716.	27.8	336
15	GTPase activation of elongation factor EF-Tu by the ribosome during decoding. EMBO Journal, 2009, 28, 755-765.	7.8	175
16	A new tRNA intermediate revealed on the ribosome during EF4-mediated back-translocation. Nature Structural and Molecular Biology, 2008, 15, 910-915.	8.2	65
17	Translational Regulation via L11: Molecular Switches on the Ribosome Turned On and Off by Thiostrepton and Micrococcin. Molecular Cell, 2008, 30, 26-38.	9.7	269
18	The oxazolidinone antibiotics perturb the ribosomal peptidyl-transferase center and effect tRNA positioning. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 13339-13344.	7.1	285

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19	Structural Basis for Interaction of the Ribosome with the Switch Regions of GTP-Bound Elongation Factors. Molecular Cell, 2007, 25, 751-764.	9.7	168
20	A dedicated translation factor controls the synthesis of the global regulator Fis. EMBO Journal, 2007, 26, 4607-4607.	7.8	0
21	Structure of the ribosome-bound cricket paralysis virus IRES RNA. Nature Structural and Molecular Biology, 2006, 13, 1092-1096.	8.2	177
22	16S rRNA Mutations That Confer Tetracycline Resistance in Helicobacter pylori Decrease Drug Binding in Escherichia coli Ribosomes. Journal of Bacteriology, 2005, 187, 3708-3712.	2.2	49
23	Incidence of Antibiotic Resistance in Campylobacter jejuni Isolated in Alberta, Canada, from 1999 to 2002, with Special Reference to tet (O)-Mediated Tetracycline Resistance. Antimicrobial Agents and Chemotherapy, 2004, 48, 3442-3450.	3.2	139
24	Heteronuclear NMR investigations of dynamic regions of intact Escherichia coli ribosomes. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 10949-10954.	7.1	87
25	A dedicated translation factor controls the synthesis of the global regulator Fis. EMBO Journal, 2004, 23, 3375-3385.	7.8	46
26	Three-Dimensional Structures of Translating Ribosomes by Cryo-EM. Molecular Cell, 2004, 14, 57-66.	9.7	104
27	Mechanism of Tet(O)-mediated tetracycline resistance. EMBO Journal, 2003, 22, 945-953.	7.8	99
28	Ribosomal Protection Proteins and Their Mechanism of Tetracycline Resistance. Antimicrobial Agents and Chemotherapy, 2003, 47, 3675-3681.	3.2	355
29	Erythromycin, Roxithromycin, and Clarithromycin: Use of Slow-Binding Kinetics to Compare Their in Vitro Interaction with a Bacterial Ribosomal Complex Active in Peptide Bond Formation. Molecular Pharmacology, 2003, 63, 617-623.	2.3	27
30	The tetracycline resistance protein Tet( $\hat{a}$ —() perturbs the conformation of the ribosomal decoding centre. Molecular Microbiology, 2002, 45, 1463-1472.	2.5	40
31	Protein Synthesis at Atomic Resolution: Mechanistics of Translation in the Light of Highly Resolved Structures for the Ribosome. Current Protein and Peptide Science, 2002, 3, 1-53.	1.4	56
32	Localization of the Ribosomal Protection Protein Tet(O) on the Ribosome and the Mechanism of Tetracycline Resistance. Molecular Cell, 2001, 7, 1037-1045.	9.7	82