

# Marat M Yusupov

## List of Publications by Year in descending order

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97  
papers

10,440  
citations

70961

41  
h-index

45213

90  
g-index

99  
all docs

99  
docs citations

99  
times ranked

7941  
citing authors

#	ARTICLE	IF	CITATIONS
1	Crystal Structure of the Ribosome at 5.5 Å Resolution. <i>Science</i> , 2001, 292, 883-896.	6.0	1,789
2	The Structure of the Eukaryotic Ribosome at 3.0 Å Resolution. <i>Science</i> , 2011, 334, 1524-1529.	6.0	1,006
3	X-ray Crystal Structures of 70S Ribosome Functional Complexes. <i>Science</i> , 1999, 285, 2095-2104.	6.0	567
4	The Path of Messenger RNA through the Ribosome. <i>Cell</i> , 2001, 106, 233-241.	13.5	554
5	A new system for naming ribosomal proteins. <i>Current Opinion in Structural Biology</i> , 2014, 24, 165-169.	2.6	481
6	Structural basis for the inhibition of the eukaryotic ribosome. <i>Nature</i> , 2014, 513, 517-522.	13.7	434
7	Crystal Structure of the Eukaryotic Ribosome. <i>Science</i> , 2010, 330, 1203-1209.	6.0	370
8	One core, two shells: bacterial and eukaryotic ribosomes. <i>Nature Structural and Molecular Biology</i> , 2012, 19, 560-567.	3.6	345
9	A new understanding of the decoding principle on the ribosome. <i>Nature</i> , 2012, 484, 256-259.	13.7	293
10	Structural aspects of messenger RNA reading frame maintenance by the ribosome. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 555-560.	3.6	276
11	Structural basis for messenger RNA movement on the ribosome. <i>Nature</i> , 2006, 444, 391-394.	13.7	245
12	Ribosomal position and contacts of mRNA in eukaryotic translation initiation complexes. <i>EMBO Journal</i> , 2008, 27, 1609-1621.	3.5	202
13	Evidence for rRNA 2'-O-methylation plasticity: Control of intrinsic translational capabilities of human ribosomes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 12934-12939.	3.3	197
14	Structure of the 30S translation initiation complex. <i>Nature</i> , 2008, 455, 416-420.	13.7	194
15	The crystal structure of the dimerization initiation site of genomic HIV-1 RNA reveals an extended duplex with two adenine bulges. <i>Structure</i> , 1999, 7, 1439-1449.	1.6	157
16	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.	3.3	152
17	Aminoglycoside interactions and impacts on the eukaryotic ribosome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E10899-E10908.	3.3	148
18	Novel base-pairing interactions at the tRNA wobble position crucial for accurate reading of the genetic code. <i>Nature Communications</i> , 2016, 7, 10457.	5.8	141

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19	Structural rearrangements of the ribosome at the tRNA proofreading step. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 1072-1078.	3.6	135
20	Conformational transition of initiation factor 2 from the GTP- to GDP-bound state visualized on the ribosome. <i>Nature Structural and Molecular Biology</i> , 2005, 12, 1145-1149.	3.6	130
21	Structured mRNAs Regulate Translation Initiation by Binding to the Platform of the Ribosome. <i>Cell</i> , 2007, 130, 1019-1031.	13.5	129
22	Translocation of tRNA during protein synthesis. <i>FEBS Letters</i> , 2002, 514, 11-16.	1.3	128
23	Importance of potassium ions for ribosome structure and function revealed by long-wavelength X-ray diffraction. <i>Nature Communications</i> , 2019, 10, 2519.	5.8	124
24	A structural view of translation initiation in bacteria. <i>Cellular and Molecular Life Sciences</i> , 2009, 66, 423-436.	2.4	123
25	Crystal structure of the 80S yeast ribosome. <i>Current Opinion in Structural Biology</i> , 2012, 22, 759-767.	2.6	120
26	High-Resolution Structure of the Eukaryotic 80S Ribosome. <i>Annual Review of Biochemistry</i> , 2014, 83, 467-486.	5.0	110
27	The fidelity of translation initiation: reciprocal activities of eIF1, IF3 and YciH. <i>EMBO Journal</i> , 2006, 25, 196-210.	3.5	105
28	Structural insights into the translational infidelity mechanism. <i>Nature Communications</i> , 2015, 6, 7251.	5.8	100
29	Translational Operator of mRNA on the Ribosome: How Repressor Proteins Exclude Ribosome Binding. <i>Science</i> , 2005, 308, 120-123.	6.0	99
30	Crystallization of 70 S ribosomes and 30 S ribosomal subunits from <i>Thermus thermophilus</i> . <i>FEBS Letters</i> , 1987, 220, 319-322.	1.3	86
31	Identification of an RNA-Protein Bridge Spanning the Ribosomal Subunit Interface. <i>Science</i> , 1999, 285, 2133-2135.	6.0	82
32	Molecular insights into protein synthesis with proline residues. <i>EMBO Reports</i> , 2016, 17, 1776-1784.	2.0	73
33	New Structural Insights into Translational Miscoding. <i>Trends in Biochemical Sciences</i> , 2016, 41, 798-814.	3.7	64
34	Structures and dynamics of hibernating ribosomes from <i>Staphylococcus aureus</i> mediated by intermolecular interactions of H <sub>41</sub> HPF. <i>EMBO Journal</i> , 2017, 36, 2073-2087.	3.5	62
35	The ribosome prohibits the G-C-U wobble geometry at the first position of the codon-anticodon helix. <i>Nucleic Acids Research</i> , 2016, 44, gkw431.	6.5	59
36	Messenger RNA conformations in the ribosomal E site revealed by X-ray crystallography. <i>EMBO Reports</i> , 2007, 8, 846-850.	2.0	58

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37	Crystal Structure of Hypusine-Containing Translation Factor eIF5A Bound to a Rotated Eukaryotic Ribosome. <i>Journal of Molecular Biology</i> , 2016, 428, 3570-3576.	2.0	53
38	The Amaryllidaceae Alkaloid Haemanthamine Binds the Eukaryotic Ribosome to Repress Cancer Cell Growth. <i>Structure</i> , 2018, 26, 416-425.e4.	1.6	51
39	New structural insights into the decoding mechanism: Translation infidelity via a G <sup>A</sup> -U pair with Watson-Crick geometry. <i>FEBS Letters</i> , 2013, 587, 1848-1857.	1.3	50
40	Bulk-solvent correction in large macromolecular structures. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2005, 61, 1299-1301.	2.5	48
41	Recognition of Watson-Crick base pairs: constraints and limits due to geometric selection and tautomerism. <i>F1000prime Reports</i> , 2014, 6, 19.	5.9	47
42	Interactions of the ribosome with mRNA and tRNA. <i>Current Opinion in Structural Biology</i> , 2010, 20, 325-332.	2.6	45
43	Preliminary X-ray investigation of 70 S ribosome crystals from <i>Thermus thermophilus</i> . <i>Journal of Molecular Biology</i> , 1989, 209, 327-328.	2.0	43
44	Amicoumacin A induces cancer cell death by targeting the eukaryotic ribosome. <i>Scientific Reports</i> , 2016, 6, 27720.	1.6	42
45	Inhibition of Eukaryotic Translation by the Antitumor Natural Product Agelastatin A. <i>Cell Chemical Biology</i> , 2017, 24, 605-613.e5.	2.5	41
46	Structure of the 70S ribosome from human pathogen <i>Staphylococcus aureus</i> . <i>Nucleic Acids Research</i> , 2016, 44, gkw933.	6.5	39
47	Structural Insights into the Role of Diphthamide on Elongation Factor 2 in mRNA Reading-Frame Maintenance. <i>Journal of Molecular Biology</i> , 2018, 430, 2677-2687.	2.0	38
48	Synthesis facilitates an understanding of the structural basis for translation inhibition by the lissoclimides. <i>Nature Chemistry</i> , 2017, 9, 1140-1149.	6.6	36
49	Ribosomal Initiation Complexes Probed by Toeprinting and Effect of trans-Acting Translational Regulators in Bacteria. <i>Methods in Molecular Biology</i> , 2009, 540, 247-263.	0.4	35
50	Crystal structure of eukaryotic ribosome and its complexes with inhibitors. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2017, 372, 20160184.	1.8	34
51	Cryo-EM structure of the hibernating <i>Thermus thermophilus</i> 100S ribosome reveals a protein-mediated dimerization mechanism. <i>Nature Communications</i> , 2018, 9, 4179.	5.8	34
52	The multiple flavors of GoU pairs in RNA. <i>Journal of Molecular Recognition</i> , 2019, 32, e2782.	1.1	30
53	Mechanism of ribosome shutdown by RsfS in <i>Staphylococcus aureus</i> revealed by integrative structural biology approach. <i>Nature Communications</i> , 2020, 11, 1656.	5.8	30
54	Tautomeric G <sup>A</sup> -C <sup>A</sup> U pairs within the molecular ribosomal grip and fidelity of decoding in bacteria. <i>Nucleic Acids Research</i> , 2018, 46, 7425-7435.	6.5	29

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55	Insights into the origin of the nuclear localization signals in conserved ribosomal proteins. <i>Nature Communications</i> , 2015, 6, 7382.	5.8	26
56	Accuracy mechanism of eukaryotic ribosome translocation. <i>Nature</i> , 2021, 600, 543-546.	13.7	26
57	Are there proteins between the ribosomal subunits?. <i>FEBS Letters</i> , 1986, 197, 229-233.	1.3	25
58	Primer Selection by HIV-1 Reverse Transcriptase on RNA tRNA <sup>3</sup> Lys and DNA tRNA <sup>3</sup> LysHybrids. <i>Journal of Molecular Biology</i> , 1996, 261, 315-321.	2.0	21
59	Structure of the Ribosome at 5.5 Å Resolution and Its Interactions with Functional Ligands. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2001, 66, 57-66.	2.0	21
60	Proteins of the <i>Thermus thermophilus</i> ribosome Purification of several individual proteins and crystallization of protein TL7. <i>FEBS Letters</i> , 1987, 220, 227-230.	1.3	20
61	Sequence of tRNA <sup>asp</sup> from <i>Thermus thermophilus</i> HB8. <i>Nucleic Acids Research</i> , 1993, 21, 4399-4399.	6.5	20
62	The location of protein S8 and surrounding elements of 16S rRNA in the 70S ribosome from combined use of directed hydroxyl radical probing and X-ray crystallography. <i>Rna</i> , 2000, 6, 717-729.	1.6	20
63	<i>Thermus thermophilus</i> ribosomes for crystallographic studies. <i>Biochimie</i> , 1991, 73, 887-897.	1.3	18
64	Crystal Structures of the uL3 Mutant Ribosome: Illustration of the Importance of Ribosomal Proteins for Translation Efficiency. <i>Journal of Molecular Biology</i> , 2016, 428, 2195-2202.	2.0	17
65	A new crystalline form of 30 S ribosomal subunits from <i>Thermus thermophilus</i> . <i>FEBS Letters</i> , 1988, 238, 113-115.	1.3	16
66	Understanding the role of intermolecular interactions between lissoclimides and the eukaryotic ribosome. <i>Nucleic Acids Research</i> , 2019, 47, 3223-3232.	6.5	15
67	Ribosome biochemistry in crystal structure determination. <i>Rna</i> , 2015, 21, 771-773.	1.6	14
68	Cryo-EM structure of the ribosome functional complex of the human pathogen <i>Staphylococcus aureus</i> at 3.2 Å resolution. <i>FEBS Letters</i> , 2020, 594, 3551-3567.	1.3	14
69	Synthesis and ribosome binding properties of model mRNAs modified with undecagold cluster. <i>Bioconjugate Chemistry</i> , 1993, 4, 549-553.	1.8	10
70	E-site drug specificity of the human pathogen <i>Candida albicans</i> ribosome. <i>Science Advances</i> , 2022, 8, .	4.7	10
71	Purification and crystallization of components of the protein-synthesizing system from <i>Thermus thermophilus</i> . <i>Journal of Crystal Growth</i> , 1991, 110, 228-236.	0.7	9
72	Stabilization of Ribosomal RNA of the Small Subunit by Spermidine in <i>Staphylococcus aureus</i> . <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 738752.	1.6	7

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73	A glimpse on Staphylococcus aureus translation machinery and its control. <i>Molecular Biology</i> , 2016, 50, 477-488.	0.4	5
74	NMR assignments of the N-terminal domain of Staphylococcus aureus hibernation promoting factor (SaHPF). <i>Biomolecular NMR Assignments</i> , 2018, 12, 85-89.	0.4	5
75	Isolation and crystallization of a chimeric Q $\beta$ replicase containing <i>Thermus thermophilus</i> EF-Ts. <i>Biochemistry (Moscow)</i> , 2010, 75, 989-994.	0.7	4
76	Backbone and side chain NMR assignments for the ribosome Elongation Factor P (EF-P) from <i>Staphylococcus aureus</i> . <i>Biomolecular NMR Assignments</i> , 2018, 12, 351-355.	0.4	4
77	Dimerization of long hibernation promoting factor from <i>Staphylococcus aureus</i> : Structural analysis and biochemical characterization. <i>Journal of Structural Biology</i> , 2020, 209, 107408.	1.3	4
78	Inhibition of the Eukaryotic 80S Ribosome as a Potential Anticancer Therapy: A Structural Perspective. <i>Cancers</i> , 2021, 13, 4392.	1.7	4
79	Crystals of <i>Thermus thermophilus</i> tRNA <sup>Asp</sup> Complexed with its Cognate Aspartyl-tRNA Synthetase Have a Solvent Content of 75%. Comparison with Other Aminoacylation Systems. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1998, 54, 1382-1386.	2.5	3
80	Reconstitution of Functionally Active <i>Thermus thermophilus</i> 30S Ribosomal Subunit from Ribosomal 16S RNA and Ribosomal Proteins. <i>Methods in Molecular Biology</i> , 2016, 1320, 303-314.	0.4	3
81	Elongation Factor P: New Mechanisms of Function and an Evolutionary Diversity of Translation Regulation. <i>Molecular Biology</i> , 2019, 53, 501-512.	0.4	3
82	Solution structure of the N-terminal domain of the <i>Staphylococcus aureus</i> hibernation promoting factor. <i>Journal of Biomolecular NMR</i> , 2019, 73, 223-227.	1.6	3
83	Structural dynamics of a spinlabeled ribosome elongation factor P (EF-P) from <i>Staphylococcus aureus</i> by EPR spectroscopy. <i>SN Applied Sciences</i> , 2019, 1, 1.	1.5	3
84	Backbone and side chain NMR assignments for the ribosome binding factor A (RbfA) from <i>Staphylococcus aureus</i> . <i>Biomolecular NMR Assignments</i> , 2019, 13, 27-30.	0.4	3
85	A Path to the Atomic-Resolution Structures of Prokaryotic and Eukaryotic Ribosomes. <i>Biochemistry (Moscow)</i> , 2021, 86, 926-941.	0.7	3
86	NMR and crystallographic structural studies of the Elongation factor P from <i>Staphylococcus aureus</i> . <i>European Biophysics Journal</i> , 2020, 49, 223-230.	1.2	2
87	Interaction of bacterial ribosomes with mRNA and tRNA as studied by X-ray crystallographic analysis. , 2011, , 45-55.		2
88	Posttranslational modification of Elongation Factor P from <i>Staphylococcus aureus</i> . <i>FEBS Open Bio</i> , 2020, 10, 1342-1347.	1.0	2
89	Crystallization of the dimerization-initiation site of genomic HIV-1 RNA: preliminary crystallographic results. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1999, 55, 281-284.	2.5	1
90	Crystal structure of the eukaryotic 80S ribosome. , 2011, , 75-81.		1

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91	Studies on the Structure and Function of Ribosomes by Combined Use of Chemical Probing and X-Ray Crystallography. , 0, , 127-150.		1
92	Is RsfS a Hibernation Factor or a Ribosome Biogenesis Factor?. Biochemistry (Moscow), 2022, 87, 500-510.	0.7	1
93	Messenger RNA movement on the ribosome. Molecular Biology, 2007, 41, 240-249.	0.4	0
94	Ribosomes Structure and Mechanisms in Regulation of Protein Synthesis Part I. Journal of Molecular Biology, 2016, 428, 2133.	2.0	0
95	Editorial. Journal of Molecular Biology, 2016, 428, 3557.	2.0	0
96	X-Ray Analysis of Prokaryotic and Eukaryotic Ribosomes. , 2012, , 1-25.		0
97	Recent Progress in Ribosome Structure Studies. , 2014, , 23-43.		0