

Alexey Amunts

List of Publications by Year in descending order

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Version: 2024-02-01

43
papers

3,842
citations

218381

26
h-index

264894

42
g-index

113
all docs

113
docs citations

113
times ranked

4220
citing authors

#	ARTICLE	IF	CITATIONS
1	The revolution evolution. <i>Nature Plants</i> , 2022, 8, 14-17.	4.7	4
2	Cryo-EM structure and rRNA modification sites of a plant ribosome. <i>Plant Communications</i> , 2022, 3, 100342.	3.6	15
3	Mechanism of mitoribosomal small subunit biogenesis and preinitiation. <i>Nature</i> , 2022, 606, 603-608.	13.7	32
4	Interconnected assembly factors regulate the biogenesis of mitoribosomal large subunit. <i>EMBO Journal</i> , 2021, 40, e106292.	3.5	36
5	Mechanism of membrane-tethered mitochondrial protein synthesis. <i>Science</i> , 2021, 371, 846-849.	6.0	76
6	Inhibition of mitochondrial translation suppresses glioblastoma stem cell growth. <i>Cell Reports</i> , 2021, 35, 109024.	2.9	33
7	ATP synthase hexamer assemblies shape cristae of <i>Toxoplasma</i> mitochondria. <i>Nature Communications</i> , 2021, 12, 120.	5.8	64
8	Application of Cryo-EM for Visualization of Mitoribosomes. <i>Methods in Molecular Biology</i> , 2021, 2192, 197-210.	0.4	1
9	Analysis of translating mitoribosome reveals functional characteristics of translation in mitochondria of fungi. <i>Nature Communications</i> , 2020, 11, 5187.	5.8	34
10	Type III ATP synthase is a symmetry-deviated dimer that induces membrane curvature through tetramerization. <i>Nature Communications</i> , 2020, 11, 5342.	5.8	37
11	Distinct pre-initiation steps in human mitochondrial translation. <i>Nature Communications</i> , 2020, 11, 2932.	5.8	45
12	Distinct structural modulation of photosystem I and lipid environment stabilizes its tetrameric assembly. <i>Nature Plants</i> , 2020, 6, 314-320.	4.7	30
13	Structure of a minimal photosystem I from the green alga <i>Dunaliella salina</i> . <i>Nature Plants</i> , 2020, 6, 321-327.	4.7	40
14	Structural basis of mitochondrial translation. <i>ELife</i> , 2020, 9, .	2.8	71
15	Ciliate mitoribosome illuminates evolutionary steps of mitochondrial translation. <i>ELife</i> , 2020, 9, .	2.8	35
16	Cryo-EM structure of the activated RET signaling complex reveals the importance of its cysteine-rich domain. <i>Science Advances</i> , 2019, 5, eaau4202.	4.7	23
17	Structure-based mechanism for activation of the AAA+ GTPase McrB by the endonuclease McrC. <i>Nature Communications</i> , 2019, 10, 3058.	5.8	19
18	Zinc depletion does not necessarily induce ribosome hibernation in mycobacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 2395-2397.	3.3	10

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19	MRPS25 mutations impair mitochondrial translation and cause encephalomyopathy. <i>Human Molecular Genetics</i> , 2019, 28, 2711-2719.	1.4	33
20	Structural Patching Fosters Divergence of Mitochondrial Ribosomes. <i>Molecular Biology and Evolution</i> , 2019, 36, 207-219.	3.5	56
21	Structure of a mitochondrial ATP synthase with bound native cardiolipin. <i>ELife</i> , 2019, 8, .	2.8	69
22	Structure of the chloroplast ribosome with chl-RRF and hibernation-promoting factor. <i>Nature Plants</i> , 2018, 4, 212-217.	4.7	61
23	Rapid Isolation of the Mitoribosome from HEK Cells. <i>Journal of Visualized Experiments</i> , 2018, , .	0.2	16
24	The structure of the yeast mitochondrial ribosome. <i>Science</i> , 2017, 355, 528-531.	6.0	161
25	The cryo-EM structure of hibernating 100S ribosome dimer from pathogenic <i>Staphylococcus aureus</i> . <i>Nature Communications</i> , 2017, 8, 723.	5.8	69
26	Structures of the human mitochondrial ribosome in native states of assembly. <i>Nature Structural and Molecular Biology</i> , 2017, 24, 866-869.	3.6	140
27	Ribosome origami. <i>Nature Structural and Molecular Biology</i> , 2017, 24, 879-881.	3.6	2
28	Cryo-EM reconstruction of the chlororibosome to 3.2Å resolution within 24h. <i>IUCr</i> , 2017, 4, 723-727.	1.0	3
29	Organization and Regulation of Mitochondrial Protein Synthesis. <i>Annual Review of Biochemistry</i> , 2016, 85, 77-101.	5.0	221
30	Mitochondrial ribosome assembly in health and disease. <i>Cell Cycle</i> , 2015, 14, 2226-2250.	1.3	157
31	Bactobolin A Binds to a Site on the 70S Ribosome Distinct from Previously Seen Antibiotics. <i>Journal of Molecular Biology</i> , 2015, 427, 753-755.	2.0	48
32	The structure of the human mitochondrial ribosome. <i>Science</i> , 2015, 348, 95-98.	6.0	432
33	Structure of the Yeast Mitochondrial Large Ribosomal Subunit. <i>Science</i> , 2014, 343, 1485-1489.	6.0	521
34	Structure of the large ribosomal subunit from human mitochondria. <i>Science</i> , 2014, 346, 718-722.	6.0	260
35	Structure of the Yeast Mitochondrial Large Ribosomal Subunit. <i>Microscopy and Microanalysis</i> , 2014, 20, 1252-1253.	0.2	1
36	Parameters of the Protein Energy Landscapes of Several Light-Harvesting Complexes Probed via Spectral Hole Growth Kinetics Measurements. <i>Journal of Physical Chemistry B</i> , 2011, 115, 2737-2747.	1.2	16

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37	Structure Determination and Improved Model of Plant Photosystem I. Journal of Biological Chemistry, 2010, 285, 3478-3486.	1.6	238
38	Plant Photosystem I Design in the Light of Evolution. Structure, 2009, 17, 637-650.	1.6	89
39	Picosecond Fluorescence Of Intact And Dissolved PSI-LHCI Crystals. Biophysical Journal, 2009, 96, 524a.	0.2	0
40	Functional organization of a plant Photosystem I: Evolution of a highly efficient photochemical machine. Plant Physiology and Biochemistry, 2008, 46, 228-237.	2.8	55
41	Picosecond Fluorescence of Intact and Dissolved PSI-LHCI Crystals. Biophysical Journal, 2008, 95, 5851-5861.	0.2	85
42	The structure of a plant photosystem I supercomplex at 3.4Å resolution. Nature, 2007, 447, 58-63.	13.7	443
43	Solving the structure of plant photosystem I”biochemistry is vital. Photochemical and Photobiological Sciences, 2005, 4, 1011.	1.6	19