## Basil J Greber

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Structural transitions in the GTP cap visualized by cryo-electron microscopy of catalytically inactive microtubules. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, .	7.1	28
2	2.5ÂÃresolution structure of human CDK-activating kinase bound to the clinical inhibitor ICEC0942. Biophysical Journal, 2021, 120, 677-686.	0.5	22
3	Recent insights into the structure of TFIID, its assembly, and its binding to core promoter. Current Opinion in Structural Biology, 2020, 61, 17-24.	5.7	42
4	The cryoelectron microscopy structure of the human CDK-activating kinase. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 22849-22857.	7.1	42
5	Structural basis for dimerization quality control. Nature, 2020, 586, 452-456.	27.8	36
6	High-resolution cryo-EM structures of TFIIH and their functional implications. Current Opinion in Structural Biology, 2019, 59, 188-194.	5.7	13
7	The Plasticity of Molecular Interactions Governs Bacterial Microcompartment Shell Assembly. Structure, 2019, 27, 749-763.e4.	3.3	50
8	The Structures of Eukaryotic Transcription Pre-initiation Complexes and Their Functional Implications. Sub-Cellular Biochemistry, 2019, 93, 143-192.	2.4	27
9	The complete structure of the human TFIIH core complex. ELife, 2019, 8, .	6.0	91
10	Architecture of the chromatin remodeler RSC and insights into its nucleosome engagement. ELife, 2019, 8, .	6.0	68
11	Cryo-EM structure of substrate-bound human telomerase holoenzyme. Nature, 2018, 557, 190-195.	27.8	171
12	High-resolution structures of mitochondrial ribosomes and their functional implications. Current Opinion in Structural Biology, 2018, 49, 44-53.	5.7	45
13	Structure of human TFIID and mechanism of TBP loading onto promoter DNA. Science, 2018, 362, .	12.6	123
14	Assembly principles and structure of a 6.5-MDa bacterial microcompartment shell. Science, 2017, 356, 1293-1297.	12.6	187
15	The cryo-electron microscopy structure of human transcription factor IIH. Nature, 2017, 549, 414-417.	27.8	89
16	Mechanistic insight into eukaryotic 60S ribosomal subunit biogenesis by cryo-electron microscopy. Rna, 2016, 22, 1643-1662.	3.5	54
17	Structure and Function of the Mitochondrial Ribosome. Annual Review of Biochemistry, 2016, 85, 103-132.	11.1	199
18	Insertion of the Biogenesis Factor Rei1 Probes the Ribosomal Tunnel during 60S Maturation. Cell, 2016, 164, 91-102.	28.9	97

BASIL J GREBER

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19	Mitochondrial 16S rRNA Is Methylated by tRNA Methyltransferase TRMT61B in All Vertebrates. PLoS Biology, 2016, 14, e1002557.	5.6	95
20	The complete structure of the 55 <i>S</i> mammalian mitochondrial ribosome. Science, 2015, 348, 303-308.	12.6	344
21	Archaeal aminoacyl-tRNA synthetases interact with the ribosome to recycle tRNAs. Nucleic Acids Research, 2014, 42, 5191-5201.	14.5	19
22	Architecture of the large subunit of the mammalian mitochondrial ribosome. Nature, 2014, 505, 515-519.	27.8	207
23	The complete structure of the large subunit of the mammalian mitochondrial ribosome. Nature, 2014, 515, 283-286.	27.8	231
24	Cryo-EM structures of Arx1 and maturation factors Rei1 and Jjj1 bound to the 60S ribosomal subunit. Nature Structural and Molecular Biology, 2012, 19, 1228-1233.	8.2	95
25	Cryo-EM Structure of the Archaeal 50S Ribosomal Subunit in Complex with Initiation Factor 6 and Implications for Ribosome Evolution. Journal of Molecular Biology, 2012, 418, 145-160.	4.2	42
26	A Midzone-Based Ruler Adjusts Chromosome Compaction to Anaphase Spindle Length. Science, 2011, 332, 465-468.	12.6	87
27	Mechanistic insight into co-translational protein processing, folding, targeting, and membrane insertion. , 2011, , 405-418.		1
28	YidC and Oxa1 Form Dimeric Insertion Pores on the Translating Ribosome. Molecular Cell, 2009, 34, 344-353.	9.7	117