Niels Fischer

List of Publications by Year in descending order

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NIFLS FISCHED

#	Article	lF	CITATIONS
1	Conformational rearrangements upon start codon recognition in human 48S translation initiation complex. Nucleic Acids Research, 2022, 50, 5282-5298.	14.5	15
2	Structural mechanism of GTPase-powered ribosome-tRNA movement. Nature Communications, 2021, 12, 5933.	12.8	33
3	Atomic-resolution protein structure determination by cryo-EM. Nature, 2020, 587, 157-161.	27.8	454
4	Mechanism of ribosome rescue by alternative ribosome-rescue factor B. Nature Communications, 2020, 11, 4106.	12.8	26
5	Ribosome dynamics during decoding. Philosophical Transactions of the Royal Society B: Biological Sciences, 2017, 372, 20160182.	4.0	76
6	The pathway to GTPase activation of elongation factor SelB on the ribosome. Nature, 2016, 540, 80-85.	27.8	93
7	Structure of the E. coli ribosome–EF-Tu complex at <3ÂÃ… resolution by Cs-corrected cryo-EM. Nature, 2015, 520, 567-570.	27.8	338
8	ProteoPlex: stability optimization of macromolecular complexes by sparse-matrix screening of chemical space. Nature Methods, 2015, 12, 859-865.	19.0	87
9	Fluctuations between multiple EF-G-induced chimeric tRNA states during translocation on the ribosome. Nature Communications, 2015, 6, 7442.	12.8	55
10	GraDeR: Membrane Protein Complex Preparation for Single-Particle Cryo-EM. Structure, 2015, 23, 1769-1775.	3.3	96
11	Energy barriers and driving forces in tRNA translocation through the ribosome. Nature Structural and Molecular Biology, 2013, 20, 1390-1396.	8.2	150
12	Functions of elongation factor G in translocation and ribosome recycling. , 2011, , 329-338.		8
13	Ribosome dynamics and tRNA movement by time-resolved electron cryomicroscopy. Nature, 2010, 466, 329-333.	27.8	400
14	GraFix: sample preparation for single-particle electron cryomicroscopy. Nature Methods, 2008, 5, 53-55.	19.0	476
15	Towards understanding selenocysteine incorporation into bacterial proteins. Biological Chemistry, 2007, 388, 1061-1067.	2.5	16
16	Spontaneous reverse movement of mRNA-bound tRNA through the ribosome. Nature Structural and Molecular Biology, 2007, 14, 318-324.	8.2	87
17	Structural Basis for the Function of the Ribosomal L7/12 Stalk in Factor Binding and GTPase Activation. Cell, 2005, 121, 991-1004.	28.9	354
18	Experimental Identification of Downhill Protein Folding. Science, 2002, 298, 2191-2195.	12.6	290