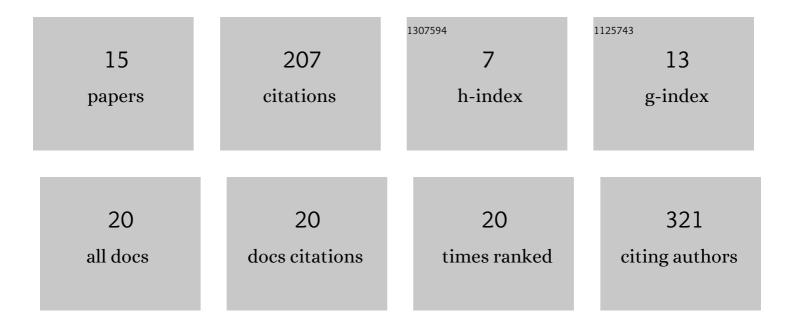
Abhishek Dey

List of Publications by Year in descending order

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Δριισμέν Πεν

#	Article	IF	CITATIONS
1	Sequence and tissue targeting specificity of ZFP36L2 reveals <i>Elavl2</i> as a novel target with co-regulation potential. Nucleic Acids Research, 2022, , .	14.5	3
2	The RNA structurome in the asexual blood stages of malaria pathogen <i>plasmodium falciparum</i> . RNA Biology, 2021, 18, 2480-2497.	3.1	6
3	To Knot or Not to Knot: Multiple Conformations of the SARS-CoV-2 Frameshifting RNA Element. Journal of the American Chemical Society, 2021, 143, 11404-11422.	13.7	35
4	<i>In vivo</i> architecture of the telomerase RNA catalytic core in <i>Trypanosoma brucei</i> . Nucleic Acids Research, 2021, 49, 12445-12466.	14.5	8
5	Innately Water-Soluble Isatoic Anhydrides with Modulated Reactivities for RNA SHAPE Analysis. Bioconjugate Chemistry, 2020, 31, 884-888.	3.6	4
6	The Non-Coding RNA Journal Club: Highlights on Recent Papers—7. Non-coding RNA, 2019, 5, 40.	2.6	2
7	Current Perspectives of Telomerase Structure and Function in Eukaryotes with Emerging Views on Telomerase in Human Parasites. International Journal of Molecular Sciences, 2018, 19, 333.	4.1	26
8	Water-Soluble Isatoic Anhydrides: A Platform for RNA-SHAPE Analysis and Protein Bioconjugation. Bioconjugate Chemistry, 2018, 29, 3196-3202.	3.6	8
9	Sequence Variation of Rare Outer Membrane Protein β-Barrel Domains in Clinical Strains Provides Insights into the Evolution of <i>Treponema pallidum</i> subsp. <i>pallidum</i> , the Syphilis Spirochete. MBio, 2018, 9, .	4.1	29
10	The major outer sheath protein forms distinct conformers and multimeric complexes in the outer membrane and periplasm of Treponema denticola. Scientific Reports, 2017, 7, 13260.	3.3	10
11	Peptide Uptake Is Essential for <i>Borrelia burgdorferi</i> Viability and Involves Structural and Regulatory Complexity of its Oligopeptide Transporter. MBio, 2017, 8, .	4.1	28
12	Crystal Structure of Mycobacterium tuberculosis H37Rv AldR (Rv2779c), a Regulator of the ald Gene. Journal of Biological Chemistry, 2016, 291, 11967-11980.	3.4	17
13	Cloning, overexpression, purification and preliminary X-ray analysis of a feast/famine regulatory protein (Rv2779c) fromMycobacterium tuberculosisH37Rv. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 97-100.	0.8	6
14	Ligand-Induced Structural Transitions, Mutational Analysis, and †Open' Quaternary Structure of the M. tuberculosis Feast/Famine Regulatory Protein (Rv3291c). Journal of Molecular Biology, 2009, 392, 1007-1019.	4.2	15
15	Quantitative prediction of variant effects on alternative splicing in MAPT using endogenous pre-messenger RNA structure probing. ELife, 0, 11, .	6.0	6