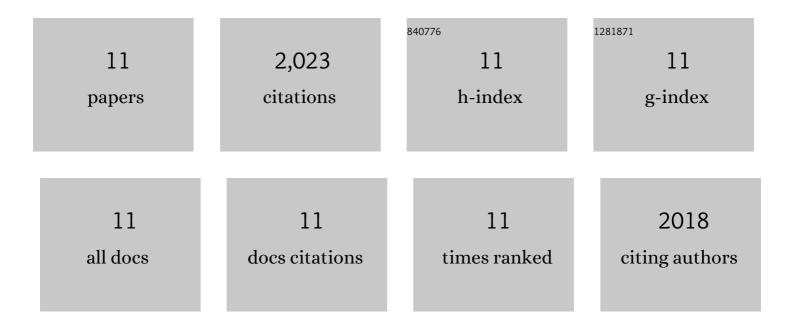
Rahul K Das

List of Publications by Year in descending order

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<u>ΡλΗΠΙ Κ Πλς</u>

#	Article	IF	CITATIONS
1	Ion Mobility Mass Spectrometry Uncovers the Impact of the Patterning of Oppositely Charged Residues on the Conformational Distributions of Intrinsically Disordered Proteins. Journal of the American Chemical Society, 2019, 141, 4908-4918.	13.7	62
2	A High-Throughput Mutational Scan of an Intrinsically Disordered Acidic Transcriptional Activation Domain. Cell Systems, 2018, 6, 444-455.e6.	6.2	135
3	CIDER: Resources to Analyze Sequence-Ensemble Relationships of Intrinsically Disordered Proteins. Biophysical Journal, 2017, 112, 16-21.	0.5	356
4	Control of transcriptional activity by design of charge patterning in the intrinsically disordered RAM region of the Notch receptor. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E9243-E9252.	7.1	95
5	Cryptic sequence features within the disordered protein p27 ^{Kip1} regulate cell cycle signaling. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 5616-5621.	7.1	109
6	Fuzzy regions in an intrinsically disordered protein impair protein–protein interactions. FEBS Journal, 2016, 283, 576-594.	4.7	43
7	Relating sequence encoded information to form and function of intrinsically disordered proteins. Current Opinion in Structural Biology, 2015, 32, 102-112.	5.7	335
8	Conformations of intrinsically disordered proteins are influenced by linear sequence distributions of oppositely charged residues. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 13392-13397.	7.1	745
9	A quantitative measure for protein conformational heterogeneity. Journal of Chemical Physics, 2013, 139, 121907.	3.0	65
10	N-Terminal Segments Modulate the α-Helical Propensities of the Intrinsically Disordered Basic Regions of bZIP Proteins. Journal of Molecular Biology, 2012, 416, 287-299.	4.2	56
11	Unmasking Functional Motifs Within Disordered Regions of Proteins. Science Signaling, 2012, 5, pe17.	3.6	22