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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Understanding COVID-19 via comparative analysis of dark proteomes of SARS-CoV-2, human SARS and bat SARS-like coronaviruses. Cellular and Molecular Life Sciences, 2021, 78, 1655-1688.	2.4	92
2	DescribePROT: database of amino acid-level protein structure and function predictions. Nucleic Acids Research, 2021, 49, D298-D308.	6.5	46
3	Accurate Sequence-Based Prediction of Deleterious nsSNPs with Multiple Sequence Profiles and Putative Binding Residues. Biomolecules, 2021, 11, 1337.	1.8	1
4	Codon selection reduces GC content bias in nucleic acids encoding for intrinsically disordered proteins. Cellular and Molecular Life Sciences, 2020, 77, 149-160.	2.4	8
5	DISOselect: Disorder predictor selection at the protein level. Protein Science, 2020, 29, 184-200.	3.1	10
6	Accuracy of protein-level disorder predictions. Briefings in Bioinformatics, 2020, 21, 1509-1522.	3.2	36
7	Identification of Intrinsic Disorder in Complexes from the Protein Data Bank. ACS Omega, 2020, 5, 17883-17891.	1.6	17
8	Disordered RNA-Binding Region Prediction with DisoRDPbind. Methods in Molecular Biology, 2020, 2106, 225-239.	0.4	16
9	Prediction of Intrinsic Disorder with Quality Assessment Using QUARTER. Methods in Molecular Biology, 2020, 2165, 83-101.	0.4	3
10	Computational Prediction of Intrinsic Disorder in Protein Sequences with the disCoP Meta-predictor. Methods in Molecular Biology, 2020, 2141, 21-35.	0.4	4
11	Intrinsically disordered domains: Sequence âž" disorder âž" function relationships. Protein Science, 2019, 28, 1652-1663.	3.1	31
12	Computational Prediction of Secondary and Supersecondary Structures from Protein Sequences. Methods in Molecular Biology, 2019, 1958, 73-100.	0.4	11
13	Quality assessment for the putative intrinsic disorder in proteins. Bioinformatics, 2019, 35, 1692-1700.	1.8	20
14	Predicting Functions of Disordered Proteins with MoRFpred. Methods in Molecular Biology, 2019, 1851, 337-352.	0.4	14
15	Disordered Function Conjunction: On the in-silico function annotation of intrinsically disordered regions. , 2019, , .		1
16	Intrinsically Disordered Proteome of Human Membrane‣ess Organelles. Proteomics, 2018, 18, e1700193.	1.3	151
17	Identification of intrinsic disorder in complexes from Protein Data Bank. , 2018, , .		0
18	Evidence for a Strong Correlation Between Transcription Factor Protein Disorder and Organismic Complexity. Genome Biology and Evolution, 2017, 9, 1248-1265.	1.1	49

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19	DisProt 7.0: a major update of the database of disordered proteins. Nucleic Acids Research, 2017, 45, D219-D227.	6.5	242
20	Back to the Future: Nuclear Magnetic Resonance and Bioinformatics Studies on Intrinsically Disordered Proteins. Advances in Experimental Medicine and Biology, 2015, 870, 1-34.	0.8	18
21	A creature with a hundred waggly tails: intrinsically disordered proteins in the ribosome. Cellular and Molecular Life Sciences, 2014, 71, 1477-1504.	2.4	119
22	Classification of Intrinsically Disordered Regions and Proteins. Chemical Reviews, 2014, 114, 6589-6631.	23.0	1,618
23	Intrinsically Disordered Proteins and Intrinsically Disordered Protein Regions. Annual Review of Biochemistry, 2014, 83, 553-584.	5.0	850
24	Improving protein order-disorder classification using charge-hydropathy plots. BMC Bioinformatics, 2014, 15, S4.	1.2	63
25	Utilization of protein intrinsic disorder knowledge in structural proteomics. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 487-498.	1.1	58
26	MoRFpred, a computational tool for sequence-based prediction and characterization of short disorder-to-order transitioning binding regions in proteins. Bioinformatics, 2012, 28, i75-i83.	1.8	311
27	CDF it all: Consensus prediction of intrinsically disordered proteins based on various cumulative distribution functions. FEBS Letters, 2009, 583, 1469-1474.	1.3	123
28	Close encounters of the third kind: disordered domains and the interactions of proteins. BioEssays, 2009, 31, 328-335.	1.2	229
29	Functional Anthology of Intrinsic Disorder. 1. Biological Processes and Functions of Proteins with Long Disordered Regions. Journal of Proteome Research, 2007, 6, 1882-1898.	1.8	525
30	Functional Anthology of Intrinsic Disorder. 2. Cellular Components, Domains, Technical Terms, Developmental Processes, and Coding Sequence Diversities Correlated with Long Disordered Regions. Journal of Proteome Research, 2007, 6, 1899-1916.	1.8	244
31	Analysis of Molecular Recognition Features (MoRFs). Journal of Molecular Biology, 2006, 362, 1043-1059.	2.0	672
32	Showing your ID: intrinsic disorder as an ID for recognition, regulation and cell signaling. Journal of Molecular Recognition, 2005, 18, 343-384.	1.1	762
33	Comparing and Combining Predictors of Mostly Disordered Proteins. Biochemistry, 2005, 44, 1989-2000.	1.2	485
34	Evolutionary Rate Heterogeneity in Proteins with Long Disordered Regions. Journal of Molecular Evolution, 2002, 55, 104-110.	0.8	398
35	Intrinsically disordered protein. Journal of Molecular Graphics and Modelling, 2001, 19, 26-59.	1.3	2,005