Edoardo Salladini

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	DisProt in 2022: improved quality and accessibility of protein intrinsic disorder annotation. Nucleic Acids Research, 2022, 50, D480-D487.	14.5	117
2	SARSâ€CoVâ€2 variants preferentially emerge at intrinsically disordered protein sites helping immune evasion. FEBS Journal, 2022, 289, 4240-4250.	4.7	25
3	αα-hub coregulator structure and flexibility determine transcription factor binding and selection in regulatory interactomes. Journal of Biological Chemistry, 2022, 298, 101963.	3.4	5
4	Exploring Manually Curated Annotations of Intrinsically Disordered Proteins with DisProt. Current Protocols, 2022, 2, .	2.9	2
5	Structural and dynamics analysis of intrinsically disordered proteins by high-speed atomic force microscopy. Nature Nanotechnology, 2021, 16, 181-189.	31.5	69
6	Connecting the $\hat{l}\pm\hat{l}\pm$ -hubs: same fold, disordered ligands, new functions. Cell Communication and Signaling, 2021, 19, 2.	6.5	9
7	αα-Hub domains and intrinsically disordered proteins: A decisive combo. Journal of Biological Chemistry, 2021, 296, 100226.	3.4	16
8	Identification of a Region in the Common Amino-terminal Domain of Hendra Virus P, V, and W Proteins Responsible for Phase Transition and Amyloid Formation. Biomolecules, 2021, 11, 1324.	4.0	20
9	PED in 2021: a major update of the protein ensemble database for intrinsically disordered proteins. Nucleic Acids Research, 2021, 49, D404-D411.	14.5	95
10	Ensemble description of the intrinsically disordered N-terminal domain of the Nipah virus P/V protein from combined NMR and SAXS. Scientific Reports, 2020, 10, 19574.	3.3	13
11	Intrinsic Disorder in Plant Transcription Factor Systems: Functional Implications. International Journal of Molecular Sciences, 2020, 21, 9755.	4.1	14
12	Conformational response to charge clustering in synthetic intrinsically disordered proteins. Biochimica Et Biophysica Acta - General Subjects, 2018, 1862, 2204-2214.	2.4	16
13	DisProt 7.0: a major update of the database of disordered proteins. Nucleic Acids Research, 2017, 45, D219-D227.	14.5	242
14	The Henipavirus V protein is a prevalently unfolded protein with a zinc-finger domain involved in binding to DDB1. Molecular BioSystems, 2017, 13, 2254-2267.	2.9	18
15	Structural analysis of the interaction between Jaburetox, an intrinsically disordered protein, and membrane models. Colloids and Surfaces B: Biointerfaces, 2017, 159, 849-860.	5.0	10
16	Structural disorder and induced folding within two cereal, ABA stress and ripening (ASR) proteins. Scientific Reports, 2017, 7, 15544.	3.3	47