## Kundlik Gadhave

List of Publications by Year in descending order

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759233 677142 24 557 12 22 citations h-index g-index papers 31 31 31 864 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The ubiquitin proteasomal system: a potential target for the management of Alzheimer's disease. Journal of Cellular and Molecular Medicine, 2016, 20, 1392-1407.	3.6	101
2	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.	5.4	92
3	Insulin signaling: An opportunistic target to minify the risk of Alzheimer's disease. Psychoneuroendocrinology, 2017, 83, 159-171.	2.7	51
4	A multitude of signaling pathways associated with Alzheimer's disease and their roles in AD pathogenesis and therapy. Medicinal Research Reviews, 2021, 41, 2689-2745.	10.5	26
5	Conformational dynamics of 13 amino acids long NSP11 of SARS-CoV-2 under membrane mimetics and different solvent conditions. Microbial Pathogenesis, 2021, 158, 105041.	2.9	26
6	The dark proteome of cancer: Intrinsic disorderedness and functionality of HIF- $1\hat{l}\pm$ along with its interacting proteins. Progress in Molecular Biology and Translational Science, 2019, 166, 371-403.	1.7	25
7	The dark side of Alzheimer's disease: unstructured biology of proteins from the amyloid cascade signaling pathway. Cellular and Molecular Life Sciences, 2020, 77, 4163-4208.	5 <b>.</b> 4	23
8	Comprehensive analysis of the catalytic and structural properties of a mu-class glutathione s-transferase from Fasciola gigantica. Scientific Reports, 2017, 7, 17547.	3.3	20
9	Docosahexaenoic Acid Increases the Potency of Soluble Epoxide Hydrolase Inhibitor in Alleviating Streptozotocin-Induced Alzheimer's Disease-Like Complications of Diabetes. Frontiers in Pharmacology, 2019, 10, 288.	3 <b>.</b> 5	20
10	Japanese encephalitis virus – exploring the dark proteome and disorder–function paradigm. FEBS Journal, 2020, 287, 3751-3776.	4.7	18
11	Mammalian antimicrobial peptide protegrinâ€4 self assembles and forms amyloidâ€like aggregates: Assessment of its functional relevance. Journal of Peptide Science, 2019, 25, e3151.	1.4	17
12	Unstructured Biology of Proteins from Ubiquitin-Proteasome System: Roles in Cancer and Neurodegenerative Diseases. Biomolecules, 2020, 10, 796.	4.0	17
13	Conformational dynamics of p53ÂN-terminal TAD2 region under different solvent conditions. Archives of Biochemistry and Biophysics, 2020, 689, 108459.	3.0	14
14	Amyloid formation by intrinsically disordered trans-activation domain of cMyb. Biochemical and Biophysical Research Communications, 2020, 524, 446-452.	2.1	13
15	Insulin-copper quantum clusters preparation and receptor targeted bioimaging. Colloids and Surfaces B: Biointerfaces, 2020, 188, 110785.	5.0	11
16	Zika virus capsid anchor forms cytotoxic amyloid-like fibrils. Virology, 2021, 560, 8-16.	2.4	11
17	Bacterioboat—A novel tool to increase the half-life period of the orally administered drug. Science Advances, 2022, 8, eabh1419.	10.3	10
18	Intrinsic disorder in proteins associated with oxidative stress-induced JNK signaling. Cellular and Molecular Life Sciences, 2022, 79, 202.	5 <b>.</b> 4	9

#	Article	IF	CITATION
19	Analysis of the dark proteome of Chandipura virus reveals maximum propensity for intrinsic disorder in phosphoprotein. Scientific Reports, 2021, 11, 13253.	3.3	8
20	The signal peptide of the amyloid precursor protein forms amyloid-like aggregates and enhances $\hat{A^2}42$ aggregation. Cell Reports Physical Science, 2021, 2, 100599.	5.6	5
21	Role of the glutaredoxin domain and FAD in the stabilization of thioredoxin glutathione reductase. Archives of Biochemistry and Biophysics, 2018, 656, 38-45.	3.0	3
22	Probing the interaction of glutathione with different shape of silver-nanoparticles by optical spectroscopy. Materials Today Communications, 2021, 26, 102137.	1.9	3
23	The mechanism of phosphatidylcholineâ€induced interference of PAP (248â€286) aggregation. Journal of Peptide Science, 2019, 25, e3152.	1.4	2
24	Transactivation domain of Adenovirus Early Region 1A (E1A): Investigating folding dynamics and aggregation. Current Research in Structural Biology, 2022, 4, 29-40.	2.2	1