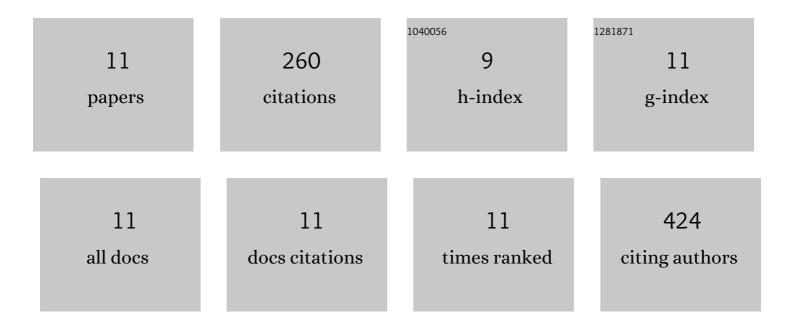
Richa Mudgal

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11368569/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	RECQL4 is essential for the transport of p53 to mitochondria in normal human cells in the absence of exogenous stress. Journal of Cell Science, 2012, 125, 2509-22.	2.0	88
2	De-DUFing the DUFs: Deciphering distant evolutionary relationships of Domains of Unknown Function using sensitive homology detection methods. Biology Direct, 2015, 10, 38.	4.6	34
3	BLM helicase stimulates the ATPase and chromatin-remodeling activities of RAD54. Journal of Cell Science, 2009, 122, 3093-3103.	2.0	30
4	Chk1-Dependent Constitutive Phosphorylation of BLM Helicase at Serine 646 Decreases after DNA Damage. Molecular Cancer Research, 2010, 8, 1234-1247.	3.4	22
5	Enhancement of c-Myc degradation by Bloom (BLM) helicase leads to delayed tumor initiation. Journal of Cell Science, 2013, 126, 3782-95.	2.0	21
6	Filling-in Void and Sparse Regions in Protein Sequence Space by Protein-Like Artificial Sequences Enables Remarkable Enhancement in Remote Homology Detection Capability. Journal of Molecular Biology, 2014, 426, 962-979.	4.2	15
7	Resolving protein structureâ€functionâ€binding site relationships from a binding site similarity network perspective. Proteins: Structure, Function and Bioinformatics, 2017, 85, 1319-1335.	2.6	14
8	NrichD database: sequence databases enriched with computationally designed protein-like sequences aid in remote homology detection. Nucleic Acids Research, 2015, 43, D300-D305.	14.5	12
9	Protein sequence design and its applications. Current Opinion in Structural Biology, 2016, 37, 71-80.	5.7	10
10	Enriching the annotation of Mycobacterium tuberculosis H37Rv proteome using remote homology detection approaches: Insights into structure and function. Tuberculosis, 2015, 95, 14-25.	1.9	9
11	Use of designed sequences in protein structure recognition. Biology Direct, 2018, 13, 8.	4.6	5