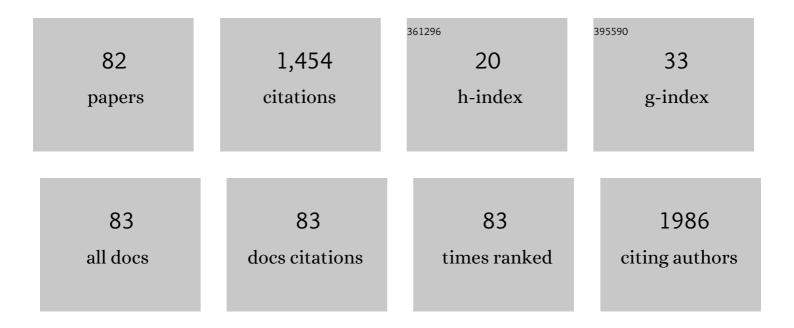
Bhartendu Nath Mishra

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Phytochemical and Pharmacological Properties of <i>Gymnema sylvestre</i> : An Important Medicinal Plant. BioMed Research International, 2014, 2014, 1-18.	0.9	126
2	Growth of hairyâ€root cultures in various bioreactors for the production of secondary metabolites. Biotechnology and Applied Biochemistry, 2008, 49, 1-10.	1.4	91
3	Prospective of biodiesel production utilizing microalgae as the cell factories: A comprehensive discussion. African Journal of Biotechnology, 2010, 9, 1402-1411.	0.3	79
4	Detection of aerolysin gene in Aeromonas hydrophila isolated from fish and pond water. Indian Journal of Microbiology, 2008, 48, 453-458.	1.5	61
5	Transcriptional regulation and signalling of cold stress response in plants: An overview of current understanding. Environmental and Experimental Botany, 2020, 180, 104243.	2.0	59
6	Future applications of electrospun nanofibers in pressure driven water treatment: A brief review and research update. Journal of Environmental Chemical Engineering, 2021, 9, 105107.	3.3	54
7	Interaction of cytochrome P4501A1 genotypes with other risk factors and susceptibility to lung cancer. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2008, 639, 1-10.	0.4	46
8	Major histocompatibility complex linked databases and prediction tools for designing vaccines. Human Immunology, 2016, 77, 295-306.	1.2	43
9	3D-QSAR studies on quinazoline antifolate thymidylate synthase inhibitors by CoMFA and CoMSIA models. European Journal of Medicinal Chemistry, 2010, 45, 1560-1571.	2.6	41
10	Association of functionally important polymorphisms in cytochrome P4501B1 with lung cancer. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2008, 643, 4-10.	0.4	34
11	Genetic transformation studies and scale up of hairy root culture of Glycyrrhiza glabra in bioreactor. Electronic Journal of Biotechnology, 2008, 11, 0-0.	1.2	34
12	Gene cloning, expression and homology modeling of hemolysin gene from Aeromonas hydrophila. Protein Expression and Purification, 2009, 65, 1-7.	0.6	32
13	A neural network approach for the prediction of in vitro culture parameters for maximum biomass yields in hairy root cultures. Journal of Theoretical Biology, 2010, 265, 579-585.	0.8	32
14	RSM–GA Based Optimization of Bacterial PHA Production and In Silico Modulation of Citrate Synthase for Enhancing PHA Production. Biomolecules, 2019, 9, 872.	1.8	31
15	Molecular docking studies on DMDP derivatives as human DHFR inhibitors. Bioinformation, 2008, 3, 180-188.	0.2	31
16	Acacetin promotes healthy aging by altering stress response in <i>Caenorhabditis elegans</i> . Free Radical Research, 2016, 50, 861-874.	1.5	29
17	Polymorphism in cytochrome P450 2A6 and glutathione S-transferase P1 modifies head and neck cancer risk and treatment outcome. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2009, 669, 36-41.	0.4	26
18	Development of diagnostic and vaccine markers through cloning, expression, and regulation of putative virulence-protein-encoding genes of Aeromonas hydrophila. Journal of Microbiology, 2013, 51, 275-282.	1.3	25

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19	Pharmacogenetic association between <i>NAT2</i> gene polymorphisms and isoniazid induced hepatotoxicity: trial sequence meta-analysis as evidence. Bioscience Reports, 2019, 39, .	1.1	23
20	Gene Cloning, Expression, and Characterization of Recombinant Aerolysin from Aeromonas hydrophila. Applied Biochemistry and Biotechnology, 2010, 160, 1985-1991.	1.4	22
21	Artificial Intelligence versus Statistical Modeling and Optimization of Cholesterol Oxidase Production by using Streptomyces Sp PLoS ONE, 2015, 10, e0137268.	1.1	21
22	Cloning, Expression, Activity and Folding Studies of Serine Hydroxymethyltransferase: A Target Enzyme for Cancer Chemotherapy. Journal of Molecular Microbiology and Biotechnology, 2003, 6, 67-75.	1.0	19
23	Computational characterization of Plasmodium falciparum proteomic data for screening of potential vaccine candidates. Human Immunology, 2010, 71, 136-143.	1.2	19
24	Molecular Cloning and Biochemical Characterization of a Recombinant Sterol 3-O-Glucosyltransferase from <i>Gymnema sylvestre</i> R.Br. Catalyzing Biosynthesis of Steryl Glucosides. BioMed Research International, 2014, 2014, 1-14.	0.9	19
25	Efficiency of neural networks for prediction of inÂvitro culture conditions and inoculum properties for optimum productivity. Plant Cell, Tissue and Organ Culture, 2008, 95, 29-35.	1.2	18
26	Role of epigenetics in carcinogenesis: Recent advancements in anticancer therapy. Seminars in Cancer Biology, 2022, 83, 441-451.	4.3	18
27	Machine Learning Techniques in Plant Biology. , 2015, , 731-754.		17
28	In-Silico Identification of Drug Lead Molecule Against Pesticide Exposed-neurodevelopmental Disorders Through Network-Based Computational Model Approach. Current Bioinformatics, 2019, 14, 460-467.	0.7	17
29	Genome-wide identification of novel vaccine candidates for Plasmodium falciparum malaria using integrative bioinformatics approaches. 3 Biotech, 2017, 7, 318.	1.1	16
30	2D-QSAR model development and analysis on variant groups of anti -tuberculosis drugs. Bioinformation, 2011, 7, 82-90.	0.2	16
31	Inhibitory effect of clove oil nanoemulsion on fumonisin isolated from maize kernels. LWT - Food Science and Technology, 2020, 134, 110237.	2.5	15
32	3D Bioprinting in Plant Science: An Interdisciplinary Approach. Trends in Plant Science, 2020, 25, 9-13.	4.3	14
33	MIF -173 G > C (rs755622) Gene Polymorphism Modulates Tuberculosis Risk: Evidence from a Meta-analysis and Trial Sequential Analysis. Scientific Reports, 2017, 7, 17003.	1.6	13
34	Microbial dysbiosis and epigenetics modulation in cancer development – A chemopreventive approach. Seminars in Cancer Biology, 2022, 86, 666-681.	4.3	13
35	Prediction of MHC binding peptide using Gibbs motif sampler, weight matrix and artificial neural network. Bioinformation, 2008, 3, 150-155.	0.2	13
36	Genome-Wide Prediction of Vaccine Candidates for <i>Leishmania major</i> : An Integrated Approach. Journal of Tropical Medicine, 2015, 2015, 1-14.	0.6	12

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37	Silybin B and Cianidanol Inhibit Mpro and Spike Protein of SARS-CoV-2: Evidence from in silico Molecular Docking Studies. Current Pharmaceutical Design, 2021, 27, 3476-3489.	0.9	12
38	<i>In Silico</i> Studies Reveal Antiviral Effects of Traditional Indian Spices on COVID-19. Current Pharmaceutical Design, 2021, 27, 3462-3475.	0.9	12
39	Association of MBL2 gene polymorphisms with pulmonary tuberculosis susceptibility: trial sequence meta-analysis as evidence. Infection and Drug Resistance, 2019, Volume 12, 185-210.	1.1	11
40	Preparation and Evaluation of the ZnO NP–Ampicillin/Sulbactam Nanoantibiotic: Optimization of Formulation Variables Using RSM Coupled GA Method and Antibacterial Activities. Biomolecules, 2019, 9, 764.	1.8	11
41	Whey protein based electrosprayed nanospheres for encapsulation and controlled release of bioactive compounds from Tinospora cordifolia extract. Innovative Food Science and Emerging Technologies, 2021, 69, 102671.	2.7	11
42	CoMFA and CoMSIA 3D-QSAR analysis of DMDP derivatives as. Bioinformation, 2008, 2, 384-391.	0.2	11
43	Molecular docking studies on quinazoline antifolate derivatives as human thymidylate synthase inhibitors. Bioinformation, 2010, 4, 357-365.	0.2	11
44	3D-QSAR and docking studies on ursolic acid derivatives for anticancer activity based on bladder cell line T24 targeting NF-kB pathway inhibition. Journal of Biomolecular Structure and Dynamics, 2019, 37, 3822-3837.	2.0	10
45	Ranking of binding and nonbinding peptides to MHC class I molecules using inverse folding approach: Implications for vaccine design. Bioinformation, 2008, 3, 72-82.	0.2	10
46	Reconstruction and visualization of carbohydrate, N-glycosylation pathways in Pichia pastoris CBS7435 using computational and system biology approaches. Systems and Synthetic Biology, 2013, 7, 7-22.	1.0	9
47	Identification and characterization of merozoite surface protein 1 epitope. Bioinformation, 2009, 4, 1-5.	0.2	9
48	Conservation of the LexA repressor binding site in Deinococcus radiodurans. Journal of Integrative Bioinformatics, 2008, 5, 1-56.	1.0	8
49	Angiotensin-Converting Enzyme Gene I/D Polymorphism Is Associated With Systemic Lupus Erythematosus Susceptibility: An Updated Meta-Analysis and Trial Sequential Analysis. Frontiers in Physiology, 2018, 9, 1793.	1.3	8
50	Structure Activity Relationship Studies of Gymnemic Acid Analogues for Antidiabetic Activity Targeting PPARÎ ³ . Current Computer-Aided Drug Design, 2015, 11, 57-71.	0.8	8
51	Coherent ontogenic dynamics of geraniol acetyltransferase activity and geranyl acetate concentration in flowers and leaves of aroma grass Cymbopogon martinii var. Motia. Plant Growth Regulation, 2009, 57, 103-108.	1.8	7
52	Design of an ON/OFF mist duty cycle in mist bioreactors for the growth of hairy roots. Biotechnology and Bioprocess Engineering, 2009, 14, 38-45.	1.4	7
53	Characterization of <i>Plasmodium falciparum </i> Proteome at Asexual Blood Stages for Screening of Effective Vaccine Candidates: An Immunoinformatics Approach. Immunology and Immunogenetics Insights, 2015, 7, III.S24755.	1.0	7
54	Impact of TNF -308 G>A (rs1800629) gene polymorphism in modulation of leprosy risk: a reappraise meta-analysis of 14 case–control studies. Bioscience Reports, 2017, 37, .	1.1	7

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55	Biotransformation of newly synthesized coumarin derivatives by Candida albicans as potential antibacterial, antioxidant and cytotoxic agents. Process Biochemistry, 2019, 87, 138-144.	1.8	7
56	Isolation, Purification, and Characterization of Heparinase from Streptomyces variabilis MTCC 12266. Scientific Reports, 2019, 9, 6482.	1.6	7
57	A trial sequential meta-analysis of <i>TNF-</i> î± â€"308G>A (rs800629) gene polymorphism and susceptibility to colorectal cancer. Bioscience Reports, 2019, 39, .	1.1	7
58	Identification of new BACE1 inhibitors for treating Alzheimer's disease. Journal of Molecular Modeling, 2021, 27, 58.	0.8	7
59	Impact of LMP7 (rs2071543) gene polymorphism in increasing cancer risk: evidence from a meta-analysis and trial sequential analysis. Oncotarget, 2018, 9, 6572-6585.	0.8	7
60	<i>IL-10</i> -1082 A>G (rs1800896) polymorphism confers susceptibility to pulmonary tuberculosis in Caucasians but not in Asians and Africans: a meta-analysis. Bioscience Reports, 2017, 37,	1.1	6
61	Lilium philadelphicum Flower as a Novel Source of Antimicrobial Agents: A Study of Bioactivity, Phytochemical Analysis, and Partial Identification of Antimicrobial Metabolites. Sustainability, 2021, 13, 8471.	1.6	6
62	Improved Prediction of MHC Class I Binders/Non-Binders Peptides Through Artificial Neural Network Using Variable Learning Rate: SARS Corona Virus, a Case Study. Advances in Experimental Medicine and Biology, 2011, 696, 223-229.	0.8	6
63	Assessing Genetic Differentiation in Geographic Populations of Labeo calbasu Using Allozyme Markers. Biochemical Genetics, 2010, 48, 760-778.	0.8	5
64	A trial sequential meta-analysis of IFN-γ +874 A>T (rs2430561) gene polymorphism and extrapulmonary tuberculosis risk. Microbial Pathogenesis, 2019, 130, 1-9.	1.3	5
65	IFN-γ +874 A>T (rs2430561) gene polymorphism and risk of pulmonary tuberculosis: aÂmeta-analysis. Archives of Medical Science, 2021, 17, 177-188.	0.4	5
66	Sustained operation of nutrient mist reactor to grow hairy roots. Asia-Pacific Journal of Chemical Engineering, 2011, 6, 23-28.	0.8	4
67	Biodegradation of pyridine raffinate using bacterial laccase isolated from garden soil. Biocatalysis and Agricultural Biotechnology, 2019, 17, 32-35.	1.5	4
68	Prediction of MHC class I binding peptides using probability distribution functions. Bioinformation, 2009, 3, 403-408.	0.2	4
69	Vitamin D Receptor Apal (rs7975232) Polymorphism Confers Decreased Risk of Pulmonary Tuberculosis in Overall and African Population, but not in Asians: Evidence from a Meta-analysis. Annals of Clinical and Laboratory Science, 2017, 47, 628-637.	0.2	4
70	GENOME WIDE IDENTIFICATION OF DNA BINDING MOTIFS OF NodD-FACTOR IN SINORHIZOBIUM MELILOTI AND MESORHIZOBIUM LOTI. Journal of Bioinformatics and Computational Biology, 2005, 03, 773-801.	0.3	3
71	Quantitative structure-activity relationship and molecular docking studies on human proteasome inhibitors for anticancer activity targeting NF-κB signaling pathway. Journal of Biomolecular Structure and Dynamics, 2020, 38, 3621-3632.	2.0	3
72	Protein Network Analysis to Prioritize Key Genes and Pathway for Stress-Mediated Neurodegeneration. Open Bioinformatics Journal, 2018, 11, 240-251.	1.0	3

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73	Prediction Model of MHC Class-II Binding Peptide Motifs Using Sequence Weighting Method for Vaccine Design. , 2012, , .		2
74	Potency of inhibitors depends upon the accessibility of their aromatic rings within the hydrophobic specificity pocket: a novel avenue for future aldose reductase inhibitor design. Journal of Biomolecular Structure and Dynamics, 2021, 39, 1512-1518.	2.0	2
75	3D QSAR Based Study of Potent Growth Inhibitors of Terpenes as Antimycobacterial Agents. The Open Nutraceuticals Journal, 2011, 4, 119-124.	0.2	2
76	5,7-Dihydroxy-4-Methoxyflavone a bioactive flavonoid delays amyloid beta-induced paralysis and attenuates oxidative stress in transgenic Caenorhabditis elegans. Pharmacognosy Magazine, 2018, 14, 57.	0.3	2
77	An Overview of Morpho-Physiological, Biochemical, and Molecular Responses of Sorghum Towards Heavy Metal Stress. Reviews of Environmental Contamination and Toxicology, 2020, 256, 155-177.	0.7	2
78	Gibbs Motif Sampler, Weight Matrix and Artificial Neural Network for the Prediction of MHC Class-II Binding Peptides. Communications in Computer and Information Science, 2009, , 503-509.	0.4	1
79	Affinity purification and partial characterization of IgM-like immunoglobulins of African catfish, Clarias gariepinus (Burchell, 1822). Indian Journal of Experimental Biology, 2006, 44, 1018-21.	0.5	1
80	Conservation of the LexA repressor binding site in Deinococcus radiodurans. Journal of Integrative Bioinformatics, 2008, 5, .	1.0	1
81	Genetics of Yellow Mosaic Virus Resistance in Mung bean. , 2010, , .		0
82	Identification of MHC Class II Binders/ Non-binders using Negative Selection Algorithm. Nature Precedings, 2011, , .	0.1	0