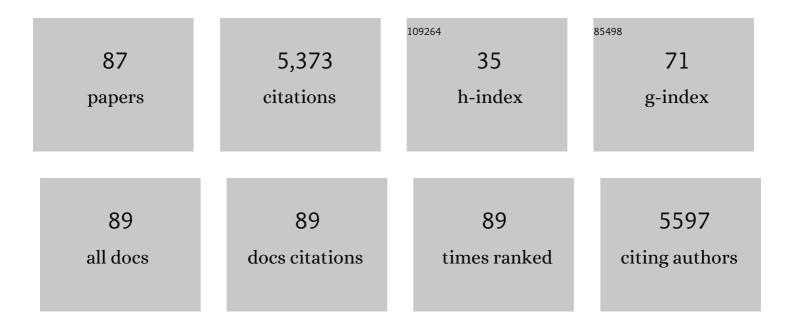
Jagjit S Yadav

List of Publications by Year in descending order

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ΙΛΟΗΤ S ΥΛΟΛΥ

#	Article	IF	CITATIONS
1	The Paleozoic Origin of Enzymatic Lignin Decomposition Reconstructed from 31 Fungal Genomes. Science, 2012, 336, 1715-1719.	6.0	1,424
2	Genome, transcriptome, and secretome analysis of wood decay fungus <i>Postia placenta</i> supports unique mechanisms of lignocellulose conversion. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 1954-1959.	3.3	530
3	Comparative genomics of <i>Ceriporiopsis subvermispora</i> and <i>Phanerochaete chrysosporium</i> provide insight into selective ligninolysis. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 5458-5463.	3.3	259
4	Comparative Genomics of Early-Diverging Mushroom-Forming Fungi Provides Insights into the Origins of Lignocellulose Decay Capabilities. Molecular Biology and Evolution, 2016, 33, 959-970.	3.5	213
5	Expansion of Signal Transduction Pathways in Fungi by Extensive Genome Duplication. Current Biology, 2016, 26, 1577-1584.	1.8	175
6	Degradation of benzene, toluene, ethylbenzene, and xylenes (BTEX) by the lignin-degrading basidiomycete Phanerochaete chrysosporium. Applied and Environmental Microbiology, 1993, 59, 756-762.	1.4	175
7	Comparative genomics of the white-rot fungi, Phanerochaete carnosa and P. chrysosporium, to elucidate the genetic basis of the distinct wood types they colonize. BMC Genomics, 2012, 13, 444.	1.2	125
8	Degradation of polychlorinated biphenyl mixtures (Aroclors 1242, 1254, and 1260) by the white rot fungus Phanerochaete chrysosporium as evidenced by congener-specific analysis. Applied and Environmental Microbiology, 1995, 61, 2560-2565.	1.4	120
9	P450 monooxygenases (P450ome) of the model white rot fungus <i>Phanerochaete chrysosporium</i> . Critical Reviews in Microbiology, 2012, 38, 339-363.	2.7	116
10	Genome-to-function characterization of novel fungal P450 monooxygenases oxidizing polycyclic aromatic hydrocarbons (PAHs). Biochemical and Biophysical Research Communications, 2010, 399, 492-497.	1.0	107
11	CYP63A2, a Catalytically Versatile Fungal P450 Monooxygenase Capable of Oxidizing Higher-Molecular-Weight Polycyclic Aromatic Hydrocarbons, Alkylphenols, and Alkanes. Applied and Environmental Microbiology, 2013, 79, 2692-2702.	1.4	93
12	Genome-wide structural and evolutionary analysis of the P450 monooxygenase genes (P450ome) in the white rot fungus Phanerochaete chrysosporium : Evidence for gene duplications and extensive gene clustering. BMC Genomics, 2005, 6, 92.	1.2	90
13	Analysis of the Phlebiopsis gigantea Genome, Transcriptome and Secretome Provides Insight into Its Pioneer Colonization Strategies of Wood. PLoS Genetics, 2014, 10, e1004759.	1.5	90
14	Development of a Single-Tube, Cell Lysis-Based, Genus-Specific PCR Method for Rapid Identification of Mycobacteria: Optimization of Cell Lysis, PCR Primers and Conditions, and Restriction Pattern Analysis. Journal of Clinical Microbiology, 2004, 42, 453-457.	1.8	74
15	Effect of electrical charges and fields on injury and viability of airborne bacteria. Biotechnology and Bioengineering, 2002, 79, 229-241.	1.7	68
16	Mineralization of 2,4-Dichlorophenoxyacetic Acid (2,4-D) and Mixtures of 2,4-D and 2,4,5-Trichlorophenoxyacetic Acid by Phanerochaete chrysosporium. Applied and Environmental Microbiology, 1993, 59, 2904-2908.	1.4	66
17	P450ome of the white rot fungus Phanerochaete chrysosporium: structure, evolution and regulation of expression of genomic P450 clusters. Biochemical Society Transactions, 2006, 34, 1165-1169.	1.6	64
18	Role of P450 Monooxygenases in the Degradation of the Endocrine-Disrupting Chemical Nonylphenol by the White Rot Fungus <i>Phanerochaete chrysosporium</i> . Applied and Environmental Microbiology, 2009, 75, 5570-5580.	1.4	59

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19	Biocidal Activity of Formaldehyde and Nonformaldehyde Biocides toward Mycobacterium immunogenum and Pseudomonas fluorescens in Pure and Mixed Suspensions in Synthetic Metalworking Fluid and Saline. Applied and Environmental Microbiology, 2005, 71, 542-546.	1.4	58
20	Differential regulation and xenobiotic induction of tandem P450 monooxygenase genes pc-1 (CYP63A1) and pc-2 (CYP63A2) in the white-rot fungus Phanerochaete chrysosporium. Applied Microbiology and Biotechnology, 2004, 65, 559-65.	1.7	56
21	Real-time PCR assays for genus-specific detection and quantification of culturable and non-culturable mycobacteria and pseudomonads in metalworking fluids. Molecular and Cellular Probes, 2004, 18, 67-73.	0.9	52
22	Mineralization of mono- and dichlorobenzenes and simultaneous degradation of chloro- and methyl-substituted benzenes by the white rot fungus Phanerochaete chrysosporium. Applied and Environmental Microbiology, 1995, 61, 677-680.	1.4	47
23	Immunoproteomic Identification of Secretory and Subcellular Protein Antigens and Functional Evaluation of the Secretome Fraction of <i>Mycobacterium immunogenum</i> , a Newly Recognized Species of the <i>Mycobacterium chelonaeâ^'Mycobacterium abscessus</i> Group. Journal of Proteome Research. 2009. 8. 2319-2330.	1.8	46
24	Gut microbiome diversity influenced more by the Westernized dietary regime than the body mass index as assessed using effect size statistic. MicrobiologyOpen, 2017, 6, e00476.	1.2	46
25	Non-involvement of lignin peroxidases and manganese peroxidases in 2,4,5-trichlorophenoxyacetic acid degradation byPhanerochaete chrysosporium. Biotechnology Letters, 1992, 14, 1089-1092.	1.1	45
26	A novel P450-initiated biphasic process for sustainable biodegradation of benzo[a]pyrene in soil under nutrient-sufficient conditions by the white rot fungus Phanerochaete chrysosporium. Journal of Hazardous Materials, 2013, 261, 675-683.	6.5	45
27	Microarray-based global differential expression profiling of P450 monooxygenases and regulatory proteins for signal transduction pathways in the white rot fungus Phanerochaete chrysosporium. Molecular Genetics and Genomics, 2005, 274, 454-466.	1.0	43
28	Multiple P450alk (cytochrome P450 alkane hydroxylase) genes from the halotolerant yeast Debaryomyces hansenii. Gene, 1999, 226, 139-146.	1.0	41
29	Biotransformation of linear alkylbenzene sulfonate (LAS) by Phanerochaete chrysosporium: oxidation of alkyl side-chain. Biodegradation, 2001, 12, 443-453.	1.5	40
30	Cytochrome b5 reductase–cytochrome b5 as an active P450 redox enzyme system in Phanerochaete chrysosporium: Atypical properties and in vivo evidence of electron transfer capability to CYP63A2. Archives of Biochemistry and Biophysics, 2011, 509, 26-32.	1.4	40
31	A Fungal P450 (CYP5136A3) Capable of Oxidizing Polycyclic Aromatic Hydrocarbons and Endocrine Disrupting Alkylphenols: Role of Trp129 and Leu324. PLoS ONE, 2011, 6, e28286.	1.1	40
32	Genomewide annotation and comparative genomics of cytochrome P450 monooxygenases (P450s) in the polypore species <i>Bjerkandera adusta</i> , <i>Ganoderma</i> sp. and <i>Phlebia brevispora</i> . Mycologia, 2013, 105, 1445-1455.	0.8	40
33	Cytochrome P450 oxidoreductase gene and its differentially terminated cDNAs from the white rot fungus Phanerochaete chrysosporium. Current Genetics, 2000, 37, 65-73.	0.8	39
34	Modulation of in vitro phagocytic uptake and immunogenicity potential of modified Herceptin®-conjugated PLGA-PEG nanoparticles for drug delivery. Colloids and Surfaces B: Biointerfaces, 2018, 162, 271-278.	2.5	38
35	Comparative toxicity reduction potential of UV/sodium percarbonate and UV/hydrogen peroxide treatments for bisphenol A in water: An integrated analysis using chemical, computational, biological, and metabolomic approaches. Water Research, 2021, 190, 116755.	5.3	37
36	Occurrence and characterization of multiple novel genotypes of Mycobacterium immunogenum and Mycobacterium chelonae in metalworking fluids. FEMS Microbiology Ecology, 2005, 54, 329-338.	1.3	35

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37	Physiological Regulation, Xenobiotic Induction, and Heterologous Expression of P450 Monooxygenase Gene pc-3 (CYP63A3), a New Member of the CYP63 Gene Cluster in the White-rot FungusPhanerochaete chrysosporium. Current Microbiology, 2005, 50, 292-298.	1.0	34
38	A new method for species identification and differentiation of Mycobacterium chelonae complex based on amplified hsp65 restriction analysis (AHSPRA). Molecular and Cellular Probes, 2005, 19, 93-99.	0.9	34
39	Genome Sequence of the Chestnut Blight Fungus <i>Cryphonectria parasitica</i> EP155: A Fundamental Resource for an Archetypical Invasive Plant Pathogen. Phytopathology, 2020, 110, 1180-1188.	1.1	34
40	Tandem cytochrome P450 monooxygenase genes and splice variants in the white rot fungus Phanerochaete chrysosporium: cloning, sequence analysis, and regulation of differential expression. Fungal Genetics and Biology, 2003, 38, 10-21.	0.9	32
41	DNA-Based Methodologies for Rapid Detection, Quantification, and Species- or Strain-Level Identification of Respiratory Pathogens (Mycobacteria and Pseudomonads) in Metalworking Fluids. Journal of Occupational and Environmental Hygiene, 2003, 18, 966-975.	0.5	30
42	A new in vitro model using small intestinal epithelial cells to enhance infection of Cryptosporidium parvum. Journal of Microbiological Methods, 2014, 106, 47-54.	0.7	30
43	Aquaporins in lung health and disease: Emerging roles, regulation, and clinical implications. Respiratory Medicine, 2020, 174, 106193.	1.3	30
44	Method for Rapid Identification and Differentiation of the Species of the Mycobacterium chelonae Complex Based on 16S-23S rRNA Gene Internal Transcribed Spacer PCR-Restriction Analysis. Journal of Clinical Microbiology, 2005, 43, 4466-4472.	1.8	29
45	Regulation and heterologous expression of P450 enzyme system components of the white rot fungus Phanerochaete chrysosporium. Enzyme and Microbial Technology, 2008, 43, 205-213.	1.6	25
46	DNA damage, redox changes, and associated stress-inducible signaling events underlying the apoptosis and cytotoxicity in murine alveolar macrophage cell line MH-S by methanol-extracted Stachybotrys chartarum toxins. Toxicology and Applied Pharmacology, 2006, 214, 297-308.	1.3	24
47	Rational engineering of the fungal P450 monooxygenase CYP5136A3 to improve its oxidizing activity toward polycyclic aromatic hydrocarbons. Protein Engineering, Design and Selection, 2013, 26, 553-557.	1.0	23
48	MyD88 mediates in vivo effector functions of alveolar macrophages in acute lung inflammatory responses to carbon nanotube exposure. Toxicology and Applied Pharmacology, 2015, 288, 322-329.	1.3	23
49	The protecting-group directed diastereoselective Nozaki–Hiyama–Kishi (NHK) reaction: total synthesis and biological evaluation of zeaenol, 7-epi-zeaenol and its analogues. Organic and Biomolecular Chemistry, 2014, 12, 9683-9695.	1.5	20
50	Cloning and Characterization of the Cytochrome P450 Oxidoreductase Gene from the Zygomycete Fungus Cunninghamella. Biochemical and Biophysical Research Communications, 2000, 268, 345-353.	1.0	19
51	Development of a Rapid ATP Bioluminescence Assay for Biocidal Susceptibility Testing of Rapidly Growing Mycobacteria. Journal of Clinical Microbiology, 2010, 48, 3725-3728.	1.8	19
52	Draft genome sequence of a monokaryotic model brown-rot fungus Postia (Rhodonia) placenta SB12. Genomics Data, 2017, 14, 21-23.	1.3	19
53	Exposure to perfluorooctanoic acid (PFOA) decreases neutrophil migration response to injury in zebrafish embryos. BMC Research Notes, 2020, 13, 408.	0.6	18
54	Molecular Detection, Quantification, and Toxigenicity Profiling of Aeromonas spp. in Source- and Drinking-Water. Open Microbiology Journal, 2014, 8, 32-39.	0.2	16

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55	Microbial P450 Enzymes in Bioremediation and Drug Discovery: Emerging Potentials and Challenges. Current Protein and Peptide Science, 2017, 19, 75-86.	0.7	16
56	Peptide nucleic acid-fluorescence in situ hybridization (PNA-FISH) assay for specific detection of Mycobacterium immunogenum and DNA-FISH assay for analysis of pseudomonads in metalworking fluids and sputum. Molecular and Cellular Probes, 2008, 22, 273-280.	0.9	15
57	Expanding the mycobacterial diversity of metalworking fluids (MWFs): evidence showing MWF colonization by Mycobacterium abscessus. FEMS Microbiology Ecology, 2012, 79, 392-399.	1.3	15
58	Multifaceted Supramolecular Interactions from Câ€Methylresorcin[4]arene Lead to an Enhancement in In Vitro Antibacterial Activity of Gatifloxacin. Chemistry - A European Journal, 2017, 23, 18171-18179.	1.7	15
59	Electrically heatable carbon nanotube point-of-use filters for effective separation and in-situ inactivation of Legionella pneumophila. Chemical Engineering Journal, 2019, 366, 21-26.	6.6	15
60	Multigenic Control and Sex Bias in Host Susceptibility to Spore-Induced Pulmonary Anthrax in Mice. Infection and Immunity, 2011, 79, 3204-3215.	1.0	14
61	Carbon Nanotube and Asbestos Exposures Induce Overlapping but Distinct Profiles of Lung Pathology in Non-Swiss Albino CF-1 Mice. Toxicologic Pathology, 2016, 44, 211-225.	0.9	14
62	Differential biocide susceptibility of the multiple genotypes of Mycobacterium immunogenum. Journal of Industrial Microbiology and Biotechnology, 2008, 35, 197-203.	1.4	13
63	Global gene expression changes underlying Stachybotrys chartarum toxin-induced apoptosis in murine alveolar macrophages: Evidence of multiple signal transduction pathways. Apoptosis: an International Journal on Programmed Cell Death, 2007, 12, 535-548.	2.2	12
64	Specific detection and quantification of culturable and non-culturable mycobacteria in metalworking fluids by fluorescence-based methods. Letters in Applied Microbiology, 2008, 47, 451-456.	1.0	12
65	Susceptibility of Mycobacterium immunogenum and Pseudomonas fluorescens to Formaldehyde and Non-Formaldehyde Biocides in Semi-Synthetic Metalworking Fluids. International Journal of Molecular Sciences, 2011, 12, 725-741.	1.8	11
66	Alveolar Macrophage Innate Response to Mycobacterium immunogenum, the Etiological Agent of Hypersensitivity Pneumonitis: Role of JNK and p38 MAPK Pathways. PLoS ONE, 2013, 8, e83172.	1.1	11
67	Crosstalk between gut microbiota and lung inflammation in murine toxicity models of respiratory exposure or co-exposure to carbon nanotube particles and cigarette smoke extract. Toxicology and Applied Pharmacology, 2022, 447, 116066.	1.3	11
68	CYPome of the conifer pathogen Heterobasidion irregulare: Inventory, phylogeny, and transcriptional analysis of the response to biocontrol. Fungal Biology, 2017, 121, 158-171.	1.1	9
69	Towards on-site detection of cadmium in human urine. Journal of Electroanalytical Chemistry, 2020, 859, 113808.	1.9	9
70	Extended tracking of the microbial community structure and dynamics in an industrial synthetic metalworking fluid system. FEMS Microbiology Ecology, 2014, 87, 664-677.	1.3	8
71	Early immunopathological events in acute model of mycobacterial hypersensitivity pneumonitis in mice. Journal of Immunotoxicology, 2017, 14, 77-88.	0.9	8
72	Genetic susceptibility to toxicologic lung responses among inbred mouse strains following exposure to carbon nanotubes and profiling of underlying gene networks. Toxicology and Applied Pharmacology, 2017, 327, 59-70.	1.3	8

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73	Mineralization of Trichloroethylene (TCE) by the White Rot Fungus Phanerochaete chrysosporium. Bulletin of Environmental Contamination and Toxicology, 2000, 65, 28-34.	1.3	7
74	Development of a species-specific colorimetric-PCR assay for detection and species differentiation of Mycobacterium immunogenum and Mycobacterium chelonae and its comparison with quantitative real-time PCR for field metalworking fluids. Molecular and Cellular Probes, 2009, 23, 75-82.	0.9	7
75	P450 Redox Enzymes in the White Rot Fungus Phanerochaete chrysosporium: Gene Transcription, Heterologous Expression, and Activity Analysis on the Purified Proteins. Current Microbiology, 2010, 61, 306-314.	1.0	7
76	Novel antigens of Mycobacterium immunogenum relevant inÂserodiagnosis of occupational hypersensitivity pneumonitis inÂmachinists. Annals of Allergy, Asthma and Immunology, 2015, 114, 525-526.e4.	0.5	6
77	T-cell antigens of Mycobacterium immunogenum, an etiological agent of occupational hypersensitivity pneumonitis. Molecular Immunology, 2016, 75, 168-177.	1.0	6
78	Secretome differences between the taxonomically related but clinically differing mycobacterial species Mycobacterium abscessus and M. chelonae. Journal of Integrated OMICS, 2012, 2, .	0.5	6
79	Omics analyses and biochemical study of Phlebiopsis gigantea elucidate its degradation strategy of wood extractives. Scientific Reports, 2021, 11, 12528.	1.6	5
80	Antimicrobial activity of selected natural products against Gram-positive, Gram-negative and Acid-fast bacterial pathogens. Alternative Medicine Studies, 2012, 2, 13.	0.2	4
81	SHP-2 Mediates Cryptosporidium parvum Infectivity in Human Intestinal Epithelial Cells. PLoS ONE, 2015, 10, e0142219.	1.1	4
82	Association of Streptomyces community composition determined by PCR-denaturing gradient gel electrophoresis with indoor mold status. Environmental Monitoring and Assessment, 2014, 186, 8773-8783.	1.3	3
83	Human leukocyte antigen (HLA)-binding epitopes dataset for the newly identified T-cell antigens of Mycobacterium immunogenum. Data in Brief, 2016, 8, 1069-1071.	0.5	3
84	Frontispiece: Multifaceted Supramolecular Interactions from Câ€Methylresorcin[4]arene Lead to an Enhancement in In Vitro Antibacterial Activity of Gatifloxacin. Chemistry - A European Journal, 2017, 23,	1.7	2
85	Capture of Magnetic Microspheres in Electrokinetic Flow for Application in Lab-on-Chip Devices. , 2012, , .		1
86	Modeling and simulation of colonization of water-based metalworking fluid by Mycobacterium immunogenum. Journal of Environmental Chemical Engineering, 2018, 6, 4953-4960.	3.3	1
87	Factors Alleviating Cadmium Toxicity in White Rot Fungus. Journal of Bioremediation & Biodegradation, 2014, 05, .	0.5	0