

Mohammad Arif

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

43 papers	455 citations	13 h-index	19 g-index
59 ext. papers	638 ext. citations	3.5 avg, IF	4.03 L-index

#	Paper	IF	Citations
43	Genomic Comparisons and Phenotypic Diversity of Strains Causing Bacterial Soft Rot of Banana in China.. <i>Frontiers in Plant Science</i> , 2022 , 13, 822829	6.2	0
42	Genome-informed loop-mediated isothermal amplification assay for specific detection of <i>Pectobacterium parmentieri</i> in infected potato tissues and soil. <i>Scientific Reports</i> , 2021 , 11, 21948	4.9	1
41	Multiple internal controls enhance reliability for PCR and real time PCR detection of <i>Rathayibacter toxicus</i> . <i>Scientific Reports</i> , 2021 , 11, 8365	4.9	1
40	Multiplex recombinase polymerase amplification assay developed using unique genomic regions for rapid on-site detection of genus <i>Clavibacter</i> and <i>C. nebraskensis</i> . <i>Scientific Reports</i> , 2021 , 11, 12017	4.9	4
39	Improved multiplex TaqMan qPCR assay with universal internal control offers reliable and accurate detection of <i>Clavibacter michiganensis</i> . <i>Journal of Applied Microbiology</i> , 2021 , 131, 1405-1416	4.7	1
38	Genomic and Phenotypic Biology of Novel Strains of Isolated From Pineapple and Taro in Hawaii: Insights Into Genome Plasticity, Pathogenicity, and Virulence Determinants. <i>Frontiers in Plant Science</i> , 2021 , 12, 663851	6.2	2
37	First Report of <i>Pectobacterium brasiliense</i> Causing Soft Rot on Brassica oleracea var. sabellica in Hawaii, United States. <i>Plant Disease</i> , 2020 , 104, 2721	1.5	2
36	First Report of <i>Pectobacterium brasiliense</i> Causing Bacterial Soft Rot and Blackleg Diseases of Potato in Hawaii. <i>Plant Disease</i> , 2020 , 104, 2515-2515	1.5	1
35	Comparative genomics reveals signature regions used to develop a robust and sensitive multiplex TaqMan real-time qPCR assay to detect the genus <i>Dickeya</i> and <i>Dickeya dianthicola</i> . <i>Journal of Applied Microbiology</i> , 2020 , 128, 1703-1719	4.7	6
34	Genome-Informed Recombinase Polymerase Amplification Assay Coupled with a Lateral Flow Device for In-Field Detection of Species. <i>Plant Disease</i> , 2020 , 104, 2217-2224	1.5	4
33	First Report of Bacterial Soft Rot and Blackleg on Potato Caused by <i>Pectobacterium parmentieri</i> in Hawaii. <i>Plant Disease</i> , 2020 , 104, 970	1.5	7
32	Genomic divergence between <i>Dickeya zeae</i> strain EC2 isolated from rice and previously identified strains, suggests a different rice foot rot strain. <i>PLoS ONE</i> , 2020 , 15, e0240908	3.7	2
31	Taxonomy and Phylogenetic Research on <i>Ralstonia solanacearum</i> Species Complex: A Complex Pathogen with Extraordinary Economic Consequences. <i>Pathogens</i> , 2020 , 9,	4.5	13
30	Exploring the Use of High-Resolution Melting Analysis and Helicase-Dependent Amplification for Discrimination of <i>Bemisia tabaci</i> (Hemiptera: Aleyrodidae) Cryptic Species and <i>Trialeurodes vaporariorum</i> . <i>Journal of Economic Entomology</i> , 2020 , 113, 2511-2520	2.2	1
29	First Report of <i>Dickeya dianthicola</i> as a Causal Agent of Bacterial Soft Rot of Potato in Hawaii. <i>Plant Disease</i> , 2019 , 103, 2943-2943	1.5	9
28	Synergetic effect of non-complementary 5' AT-rich sequences on the development of a multiplex TaqMan real-time PCR for specific and robust detection of <i>Clavibacter michiganensis</i> and <i>C. michiganensis</i> subsp. <i>nebraskensis</i> . <i>PLoS ONE</i> , 2019 , 14, e0218530	3.7	4
27	Development of a robust, field-deployable loop-mediated isothermal amplification (LAMP) assay for specific detection of potato pathogen <i>Dickeya dianthicola</i> targeting a unique genomic region. <i>PLoS ONE</i> , 2019 , 14, e0218868	3.7	22

26	Phylogenetic Analyses of Xanthomonads Causing Bacterial Leaf Spot of Tomato and Pepper: Revealed Homologous Populations Despite Distant Geographical Distribution. <i>Microorganisms</i> , 2019 , 7,	4.9	4
25	Genome-Wide Analyses Revealed Remarkable Heterogeneity in Pathogenicity Determinants, Antimicrobial Compounds, and CRISPR-Cas Systems of Complex Phytopathogenic Genus. <i>Pathogens</i> , 2019 , 8,	4.5	9
24	Development of a loop-mediated isothermal amplification assay for specific detection of all known subspecies of <i>Clavibacter michiganensis</i> . <i>Journal of Applied Microbiology</i> , 2019 , 126, 388-401	4.7	12
23	Development of a genome-informed loop-mediated isothermal amplification assay for rapid and specific detection of <i>Xanthomonas euvesicatoria</i> . <i>Scientific Reports</i> , 2018 , 8, 14298	4.9	29
22	Genome-informed diagnostics for specific and rapid detection of <i>Pectobacterium</i> species using recombinase polymerase amplification coupled with a lateral flow device. <i>Scientific Reports</i> , 2018 , 8, 15972	4.9	24
21	Single-Target and Multiplex Discrimination of Whiteflies (Hemiptera: Aleyrodidae) <i>Bemisia tabaci</i> and <i>Trialeurodes vaporariorum</i> With Modified Priming Oligonucleotide Thermodynamics. <i>Journal of Economic Entomology</i> , 2017 , 110, 1821-1830	2.2	7
20	Development of a Loop-Mediated Isothermal Amplification Assay for the Detection of <i>Dickeya</i> spp. <i>Phytopathology</i> , 2017 , 107, 1339-1345	3.8	11
19	Antibacterial Effect of Potassium Tetraborate Tetrahydrate against Soft Rot Disease Agent in Tomato. <i>Frontiers in Microbiology</i> , 2017 , 8, 1728	5.7	10
18	A simplified strategy for sensitive detection of Rose rosette virus compatible with three RT-PCR chemistries. <i>Journal of Virological Methods</i> , 2016 , 232, 47-56	2.6	22
17	Emergence of a New Population of <i>Rathayibacter toxicus</i> : An Ecologically Complex, Geographically Isolated Bacterium. <i>PLoS ONE</i> , 2016 , 11, e0156182	3.7	13
16	Array of Synthetic Oligonucleotides to Generate Unique Multi-Target Artificial Positive Controls and Molecular Probe-Based Discrimination of <i>Liposcelis</i> Species. <i>PLoS ONE</i> , 2015 , 10, e0129810	3.7	14
15	Sensitive detection and discrimination method for studying multiple infections of five major plant viruses infecting ornamental plants in nursery environments. <i>Annals of Applied Biology</i> , 2015 , 166, 286-296	3.6	7
14	Morphological and comparative genomic analyses of pathogenic and non-pathogenic <i>Fusarium solani</i> isolated from <i>Dalbergia sissoo</i> . <i>Molecular Biology Reports</i> , 2015 , 42, 1107-22	2.8	9
13	Primer modification improves rapid and sensitive in vitro and field-deployable assays for detection of high plains virus variants. <i>Applied and Environmental Microbiology</i> , 2014 , 80, 320-7	4.8	27
12	Highly Sensitive End-Point PCR and SYBR Green qPCR Detection of <i>Phymatotrichopsis omnivora</i> , Causal Fungus of Cotton Root Rot. <i>Plant Disease</i> , 2014 , 98, 1205-1212	1.5	7
11	A multi-target, non-infectious and clonable artificial positive control for routine PCR-based assays. <i>Journal of Microbiological Methods</i> , 2013 , 95, 229-34	2.8	6
10	Comparative assessment of 5' A/T-rich overhang sequences with optimal and sub-optimal primers to increase PCR yields and sensitivity. <i>Molecular Biotechnology</i> , 2013 , 55, 17-26	3	31
9	Development of a rapid, sensitive, and field-deployable razor ex BioDetection system and quantitative PCR assay for detection of <i>Phymatotrichopsis omnivora</i> using multiple gene targets. <i>Applied and Environmental Microbiology</i> , 2013 , 79, 2312-20	4.8	26

8	Molecular phylogeny and pathotyping of <i>Fusarium solani</i> : a causal agent of <i>Dalbergia sissoo</i> decline. <i>Forest Pathology</i> , 2013 , 43, 478-487	1.2	4
7	Enhanced reliability and accuracy for field deployable bioforensic detection and discrimination of <i>Xylella fastidiosa</i> subsp. <i>pauca</i> , causal agent of citrus variegated chlorosis using razor ex technology and TaqMan quantitative PCR. <i>PLoS ONE</i> , 2013 , 8, e81647	3.7	29
6	PCR and isothermal-based molecular identification of the stored-product psocid pest <i>Lepinotus reticulatus</i> (Psocoptera: Trogiidae). <i>Journal of Stored Products Research</i> , 2012 , 49, 184-188	2.5	16
5	Development of specific primers for genus <i>Fusarium</i> and <i>F. solani</i> using rDNA sub-unit and transcription elongation factor (TEF-1) gene. <i>African Journal of Biotechnology</i> , 2011 , 11,	0.6	7
4	PCR-Based Identification and Characterization of <i>Fusarium</i> sp. Associated with Mango Malformation. <i>Biotechnology Research International</i> , 2011 , 2011, 141649		11
3	A Comparative Analysis of ISSR and RAPD Markers for Study of Genetic Diversity in Shisham (<i>Dalbergia sissoo</i>). <i>Plant Molecular Biology Reporter</i> , 2009 , 27, 488-495	1.7	33
2	New plant hosts for group 16SrII, <i>Candidatus Phytoplasma aurantifolia</i> in India. <i>Plant Pathology</i> , 2009 , 58, 391-391	2.8	5
1	Comparative genomics reveals signature regions used to develop a robust and sensitive multiplex TaqMan real-time qPCR assay to detect the genus <i>Dickeya</i> and <i>Dickeya dianthicola</i>		1