

Zohreh Moradi

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

103
citations

1478505

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1372567

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times ranked

74
citing authors

#	ARTICLE	IF	CITATIONS
1	Occurrence and Evolutionary Analysis of Coat Protein Gene Sequences of Iranian Isolates of Sugarcane mosaic virus. <i>Plant Pathology Journal</i> , 2017, 33, 296-306.	1.7	25
2	The complete genome sequences of two naturally occurring recombinant isolates of Sugarcane mosaic virus from Iran. <i>Virus Genes</i> , 2016, 52, 270-280.	1.6	22
3	Iranian johnsongrass mosaic virus: the complete genome sequence, molecular and biological characterization, and comparison of coat protein gene sequences. <i>Virus Genes</i> , 2017, 53, 77-88.	1.6	13
4	Diagnosis and Molecular Variability of Watermelon mosaic virus Isolates from North, East, North-east and North-west Regions of Iran. <i>Asian Journal of Plant Pathology</i> , 2011, 5, 115-125.	0.3	11
5	Genetic variability and molecular evolution of Bean common mosaic virus populations in Iran: comparison with the populations in the world. <i>European Journal of Plant Pathology</i> , 2019, 154, 673-690.	1.7	10
6	Genetic diversity and biological characterization of sugarcane streak mosaic virus isolates from Iran. <i>VirusDisease</i> , 2018, 29, 316-323.	2.0	7
7	Whole-Genome Characterization of Alfalfa Mosaic Virus Obtained from Metagenomic Analysis of <i>Vinca minor</i> and <i>Wisteria sinensis</i> in Iran: with Implications for the Genetic Structure of the Virus. <i>Plant Pathology Journal</i> , 2021, 37, 619-631.	1.7	6
8	Meta-transcriptomic analysis reveals an isolate of aphid lethal paralysis virus from <i>Wisteria sinensis</i> in Iran. <i>Virus Research</i> , 2022, 315, 198770.	2.2	3
9	Molecular characterization of two highly divergent Iranian johnsongrass mosaic virus isolates from <i>Zea mays</i> . <i>VirusDisease</i> , 2021, 32, 155-160.	2.0	2
10	Whole genome characterization of wisteria vein mosaic virus from Iran and its relationship to other members of bean common mosaic virus group. <i>3 Biotech</i> , 2021, 11, 407.	2.2	2
11	Analysis of the complete genome sequence of cucumber mosaic virus from <i>Vinca minor</i> and <i>Wisteria sinensis</i> in Iran. <i>Journal of Plant Pathology</i> , 2020, 102, 1263-1268.	1.2	1
12	Complete Genomic Characterization of Two Beet Soil-Borne Virus Isolates from Turkey: Implications of Comparative Analysis of Genome Sequences. <i>Plant Pathology Journal</i> , 2021, 37, 152-161.	1.7	1