Mohsen Mehrvar

List of Publications by Year
in descending order

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26	195	1163117 8	1125743 13
papers	citations	h-index	g-index

27	27	27	203
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Diversity of Beet curly top Iran virus isolated from different hosts in Iran. Virus Genes, 2013, 46, 571-575.	1.6	29
2	Occurrence and Evolutionary Analysis of Coat Protein Gene Sequences of Iranian Isolates of Sugarcane mosaic virus. Plant Pathology Journal, 2017, 33, 296-306.	1.7	25
3	The complete genome sequences of two naturally occurring recombinant isolates of Sugarcane mosaic virus from Iran. Virus Genes, 2016, 52, 270-280.	1.6	22
4	Iranian beet necrotic yellow vein virus (BNYVV): pronounced diversity of the p25 coding region in A-type BNYVV and identification of P-type BNYVV lacking a fifth RNA species. Archives of Virology, 2009, 154, 501-506.	2.1	18
5	Genetic structure and molecular variability of potato virus M populations. Archives of Virology, 2014, 159, 2081-2090.	2.1	13
6	Iranian johnsongrass mosaic virus: the complete genome sequence, molecular and biological characterization, and comparison of coat protein gene sequences. Virus Genes, 2017, 53, 77-88.	1.6	13
7	A full-length infectious clone of beet soil-borne virus indicates the dispensability of the RNA-2 for virus survival in planta and symptom expression on Chenopodium quinoa leaves. Journal of General Virology, 2009, 90, 3051-3056.	2.9	11
8	Genetic variability and molecular evolution of Bean common mosaic virus populations in Iran: comparison with the populations in the world. European Journal of Plant Pathology, 2019, 154, 673-690.	1.7	10
9	Nucleotide sequence analyses of coat protein gene of peanut stunt virus isolates from alfalfa and different hosts show a new tentative subgroup from Iran. VirusDisease, 2017, 28, 295-302.	2.0	7
10	Identification and characterization of a phytoplasma associated with black locust yellow disease in two provinces of Iran. Crop Protection, 2018, 110, 261-268.	2.1	7
11	Genetic diversity and biological characterization of sugarcane streak mosaic virus isolates from Iran. VirusDisease, 2018, 29, 316-323.	2.0	7
12	Whole-Genome Characterization of Alfalfa Mosaic Virus Obtained from Metagenomic Analysis of Vinca minor and Wisteria sinensis in Iran: with Implications for the Genetic Structure of the Virus. Plant Pathology Journal, 2021, 37, 619-631.	1.7	6
13	Growth analysis of rhizomania infected and healthy sugar beet. Journal of Crop Science and Biotechnology, 2014, 17, 59-69.	1.5	4
14	First report of Wisteria vein mosaic virus on Wisteria sinensis in Iran. New Disease Reports, 2018, 38, 18-18.	0.8	4
15	Molecular variability in the cysteine rich protein of potato virus M. VirusDisease, 2015, 26, 117-122.	2.0	3
16	Comparative genetic diversity of potato virus Y populations based on coat protein gene. Acta Virologica, 2017, 61, 161-174.	0.8	3
17	Molecular characterization of two sugarcane streak mosaic virus isolates from Iran with emphasis on its population structure. Acta Virologica, 2017, 61, 428-437.	0.8	3
18	Dynamics of the population structure and genetic variability within Iranian isolates of grapevine fanleaf virus: evidence for polyphyletic origin. Acta Virologica, 2017, 61, 324-335.	0.8	2

#	Article	IF	Citations
19	Molecular characterization of two highly divergent Iranian johnsongrass mosaic virus isolates from Zea mays. VirusDisease, 2021, 32, 155-160.	2.0	2
20	Whole genome characterization of wisteria vein mosaic virus from Iran and its relationship to other members of bean common mosaic virus group. 3 Biotech, 2021, 11, 407.	2.2	2
21	Analysis of the complete genome sequence of cucumber mosaic virus from Vinca minor and Wisteria sinensis in Iran. Journal of Plant Pathology, 2020, 102, 1263-1268.	1.2	1
22	Complete Genomic Characterization of Two Beet Soil-Borne Virus Isolates from Turkey: Implications of Comparative Analysis of Genome Sequences. Plant Pathology Journal, 2021, 37, 152-161.	1.7	1
23	Analysis of the molecular and biological variability of Zucchini yellow mosaic virus isolates from Iran and Iraq. Gene, 2021, 788, 145674.	2.2	1
24	Identification of garlic-infecting leek yellow stripe virus through deep-sequencing analyses from Iran. VirusDisease, 2021, 32, 595-600.	2.0	1
25	Distribution and phylogenetic analysis of the 3â€²UTR and coat protein gene of Iranian Beet black scorch virus. Journal of Plant Diseases and Protection, 2019, 126, 535-542.	2.9	0
26	Identification of the experimental herbaceous host range of the Apscaviroids infecting citrus species. Acta Virologica, 2019, 63, 415-422.	0.8	0