

# Jae Sun Moon

## List of Publications by Year in descending order

Source: [//exaly.com/author-pdf/7628564/publications.pdf](https://exaly.com/author-pdf/7628564/publications.pdf)

Version: 2024-02-01

116  
papers

2,329  
citations

214121

24  
h-index

215937

43  
g-index

118  
all docs

118  
docs citations

118  
times ranked

2991  
citing authors

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 1  | Pyrrroloquinoline Quinone Is a Plant Growth Promotion Factor Produced by <i>Pseudomonas fluorescens</i> B16. <i>Plant Physiology</i> , 2008, 146, 657-668.  | 5.1 | 211       |
| 2  | The Activated SA and JA Signaling Pathways Have an Influence on flg22-Triggered Oxidative Burst and Callose Deposition. <i>PLoS ONE</i> , 2014, 9, e88951.  | 2.4 | 137       |
| 3  | Regulation of polar flagellum genes is mediated by quorum sensing and FlhDC in <i>Burkholderia glumae</i> . <i>Molecular Microbiology</i> , 2007, 64, 165-179.  | 2.5 | 111       |
| 4  | Expression of an evolutionarily distinct novel BiP gene during the unfolded protein response in <i>Arabidopsis thaliana</i> . <i>Gene</i> , 2003, 311, 81-91.   | 2.3 | 105       |
| 5  | Small-molecule inhibitor binding to an <i>N</i> -acyl-homoserine lactone synthase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 12089-12094.   | 7.6 | 102       |
| 6  | Amyloidogenesis of Type III-dependent Harpins from Plant Pathogenic Bacteria. <i>Journal of Biological Chemistry</i> , 2007, 282, 13601-13609.  | 3.5 | 95        |
| 7  | Rice Mitogen-Activated Protein Kinase Interactome Analysis Using the Yeast Two-Hybrid System. <i>Plant Physiology</i> , 2012, 160, 477-487.   | 5.1 | 81        |
| 8  | Bases of biocontrol: Sequence predicts synthesis and mode of action of agrocin 84, the Trojan Horse antibiotic that controls crown gall. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 8846-8851. | 7.6 | 79        |
| 9  | Synthesis of 13-(substituted benzyl) berberine and berberrubine derivatives as antifungal agents. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2006, 16, 3913-3916.  | 2.4 | 71        |
| 10 | Mutational Analysis of <i>Xanthomonas</i> Harpin HpaG Identifies a Key Functional Region That Elicits the Hypersensitive Response in Nonhost Plants. <i>Journal of Bacteriology</i> , 2004, 186, 6239-6247.   | 2.4 | 65        |
| 11 | Comparative genome analysis of rice-pathogenic <i>Burkholderia</i> provides insight into capacity to adapt to different environments and hosts. <i>BMC Genomics</i> , 2015, 16, 349.  | 2.9 | 49        |
| 12 | Complete Genome Sequence of <i>Burkholderia gladioli</i> BSR3. <i>Journal of Bacteriology</i> , 2011, 193, 3149-3149.   | 2.4 | 48        |
| 13 | Alcohol and Club Drug Use Among Same-Sex Attracted Young People: Associations With Frequenting the Lesbian and Gay Scene and Other Bars and Nightclubs. <i>Substance Use and Misuse</i> , 2013, 48, 129-136.  | 1.4 | 46        |
| 14 | Proteomic analysis of the proteins regulated by HrpB from the plant pathogenic bacterium <i>Burkholderia glumae</i> . <i>Proteomics</i> , 2008, 8, 106-121.   | 2.7 | 44        |
| 15 | <i>Xanthomonas oryzae</i> pv. <i>oryzae</i> Type III Effector XopN Targets OsVOZ2 and a Putative Thiamine Synthase as a Virulence Factor in Rice. <i>PLoS ONE</i> , 2013, 8, e73346.  | 2.4 | 40        |
| 16 | Monocyte Chemotactic Protein 1 (MCP-1) Modulates Pro-Survival Signaling to Promote Progression of Head and Neck Squamous Cell Carcinoma. <i>PLoS ONE</i> , 2014, 9, e88952.   | 2.4 | 38        |
| 17 | Complete genome sequence of a novel endornavirus isolated from hot pepper. <i>Archives of Virology</i> , 2015, 160, 3153-3156.  | 1.9 | 32        |
| 18 | Nucleotide sequence and genomic organization of a newly identified member of the genus Carmovirus, soybean yellow mottle mosaic virus, from soybean. <i>Archives of Virology</i> , 2009, 154, 1679-1684.  | 1.9 | 31        |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | Development of tobacco ringspot virus-based vectors for foreign gene expression and virus-induced gene silencing in a variety of plants. <i>Virology</i> , 2016, 492, 166-178.                        | 2.5 | 30        |
| 20 | The complete genomic sequence of a tentative new polerovirus identified in barley in South Korea. <i>Archives of Virology</i> , 2016, 161, 2047-2050.   | 1.9 | 29        |
| 21 | Retrospective surveillance of porcine circovirus 4 in pigs in Inner Mongolia, China, from 2016 to 2018. <i>Archives of Virology</i> , 2021, 166, 1951-1959.   | 1.9 | 29        |
| 22 | Construction of SARS-CoV-2 virus-like particles in plant. <i>Scientific Reports</i> , 2022, 12, 1005.   | 3.5 | 28        |
| 23 | Antifungal Activity of CHE-23C, a Dimeric Sesquiterpene from <i>Chloranthus henryi</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2009, 57, 5750-5755.                                     | 5.4 | 27        |
| 24 | Genome sequence of a recombinant brassica yellows virus infecting Chinese cabbage. <i>Archives of Virology</i> , 2015, 160, 597-600.  | 1.9 | 27        |
| 25 | Sesquiterpene Furan Compound CJ-01, a Novel Chitin Synthase 2 Inhibitor from <i>Chloranthus japonicus</i> SIEB.. <i>Biological and Pharmaceutical Bulletin</i> , 2008, 31, 1041-1044.                 | 1.5 | 26        |
| 26 | Synthesis and antifungal activity of a novel series of 13-(4-isopropylbenzyl)berberine derivatives. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2010, 20, 6551-6554.                          | 2.4 | 26        |
| 27 | Rapid and Specific Detection of Apple stem grooving virus by Reverse Transcription-recombinase Polymerase Amplification. <i>Plant Pathology Journal</i> , 2018, 34, 575-579.                          | 1.7 | 26        |
| 28 | Development of Multiplex RT-PCR for Simultaneous Detection of Garlic Viruses and the Incidence of Garlic Viral Disease in Garlic Genetic Resources. <i>Plant Pathology Journal</i> , 2015, 31, 90-96. | 1.7 | 24        |
| 29 | A novel light-dependent selection marker system in plants. <i>Plant Biotechnology Journal</i> , 2011, 9, 348-358.   | 8.5 | 22        |
| 30 | Comparative analysis of three indigenous plasmids from <i>Xanthomonas axonopodis</i> pv. <i>glycines</i> . <i>Plasmid</i> , 2006, 56, 79-87.  | 1.4 | 21        |
| 31 | Biochemical Evidence for ToxR and ToxJ Binding to the <i>tox</i> Operons of <i>Burkholderia glumae</i> and Mutational Analysis of ToxR. <i>Journal of Bacteriology</i> , 2009, 191, 4870-4878.        | 2.4 | 20        |
| 32 | Isolation and characterization of a high iturin yielding <i>Bacillus velezensis</i> UV mutant with improved antifungal activity. <i>PLoS ONE</i> , 2020, 15, e0234177.                                | 2.4 | 20        |
| 33 | Deep Sequencing Analysis of Apple Infecting Viruses in Korea. <i>Plant Pathology Journal</i> , 2016, 32, 441-451.   | 1.7 | 19        |
| 34 | Inhibition of Chitin Synthase 2 and Antifungal Activity of Lignans from the Stem Bark of <i>Lindera erythrocarpa</i> . <i>Planta Medica</i> , 2007, 73, 679-682.                                      | 1.8 | 18        |
| 35 | Complete nucleotide sequence and genome organization of peach virus D, a putative new member of the genus Marafivirus. <i>Archives of Virology</i> , 2017, 162, 1769-1772.                            | 1.9 | 18        |
| 36 | An HrpB-dependent but type III-independent extracellular aspartic protease is a virulence factor of <i>Ralstonia solanacearum</i> . <i>Molecular Plant Pathology</i> , 2011, 12, 373-380.             | 4.3 | 17        |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 37 | Arabidopsis TTR1 Causes LRR-Dependent Lethal Systemic Necrosis, rather than Systemic Acquired Resistance, to Tobacco Ringspot Virus. <i>Molecules and Cells</i> , 2011, 32, 421-430.   | 2.6 | 17        |
| 38 | Development of the Large-Scale Oligonucleotide Chip for the Diagnosis of Plant Viruses and its Practical Use. <i>Plant Pathology Journal</i> , 2014, 30, 51-57.  | 1.7 | 17        |
| 39 | Development of a new vector using Soybean yellow common mosaic virus for gene function study or heterologous protein expression in soybeans. <i>Journal of Virological Methods</i> , 2016, 228, 1-9.                           | 2.2 | 17        |
| 40 | Organization and characterization of genetic regions in <i>Bacillus subtilis</i> subsp. <i>krietiensis</i> ATCC55079 associated with the biosynthesis of iturin and surfactin compounds. <i>PLoS ONE</i> , 2017, 12, e0188179. | 2.4 | 17        |
| 41 | Genomic detection and characterization of a Korean isolate of Little cherry virus 1 sampled from a peach tree. <i>Virus Genes</i> , 2015, 51, 260-266.   | 1.9 | 16        |
| 42 | The complete genome sequences of two isolates of cnidium vein yellowing virus, a tentative new member of the family Secoviridae. <i>Archives of Virology</i> , 2015, 160, 2911-2914.   | 1.9 | 15        |
| 43 | Optimization of a Virus-Induced Gene Silencing System with Soybean yellow common mosaic virus for Gene Function Studies in Soybeans. <i>Plant Pathology Journal</i> , 2016, 32, 112-122.                                       | 1.7 | 15        |
| 44 | Protein interactome analysis of 12 mitogen-activated protein kinase kinase kinase in rice using a yeast two-hybrid system. <i>Proteomics</i> , 2014, 14, 105-115.  | 2.7 | 14        |
| 45 | Complete genome sequence of a tentative new caulimovirus from the medicinal plant <i>Atractylodes macrocephala</i> . <i>Archives of Virology</i> , 2015, 160, 3127-3131.   | 1.9 | 14        |
| 46 | The complete nucleotide sequence and genome organization of lychnis mottle virus. <i>Archives of Virology</i> , 2015, 160, 2891-2894.  | 1.9 | 14        |
| 47 | Complete nucleotide sequence of a highly divergent cherry-associated luteovirus (ChALV) isolate from peach in South Korea. <i>Archives of Virology</i> , 2017, 162, 2893-2896.   | 1.9 | 14        |
| 48 | Gibberellin Promotes Bolting and Flowering via the Floral Integrators RsFT and RsSOC1-1 under Marginal Vernalization in Radish. <i>Plants</i> , 2020, 9, 594.  | 3.6 | 14        |
| 49 | Biological and molecular characterization of Soybean yellow common mosaic virus, a new species in the genus Sobemovirus. <i>Virus Research</i> , 2012, 163, 363-367.   | 2.3 | 13        |
| 50 | Complete genome sequence and construction of infectious full-length cDNA clones of tobacco ringspot Nepovirus, a viral pathogen causing bud blight in soybean. <i>Virus Genes</i> , 2015, 51, 163-166.                         | 1.9 | 13        |
| 51 | First report of citrus leaf blotch virus in Satsuma mandarin in Korea. <i>Journal of Plant Pathology</i> , 2019, 101, 1229-1229.   | 1.2 | 12        |
| 52 | Comparative proteomic analysis of host responses to <i>Plasmodiophora brassicae</i> infection in susceptible and resistant <i>Brassica oleracea</i> . <i>Plant Biotechnology Reports</i> , 2020, 14, 263-274.                  | 1.6 | 12        |
| 53 | Lymphatic Involvement in the Disappearance of Steroidogenic Cells from the Corpus Luteum during Luteolysis. <i>PLoS ONE</i> , 2014, 9, e88953.   | 2.4 | 11        |
| 54 | Molecular detection and characterization of a divergent isolate of <i>Plantago asiatica</i> mosaic virus in <i>Plantago asiatica</i> . <i>VirusDisease</i> , 2016, 27, 307-310.  | 2.1 | 10        |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 55 | Genome-wide identification of flowering time genes associated with vernalization and the regulatory flowering networks in Chinese cabbage. <i>Plant Biotechnology Reports</i> , 2018, 12, 347-363.  | 1.6 | 10        |
| 56 | The complete genome sequence of apple rootstock virus A, a novel nucleorhabdovirus identified in apple rootstocks. <i>Archives of Virology</i> , 2019, 164, 2641-2644.  | 1.9 | 10        |
| 57 | Suppression of pepper SGT1 and SKP1 causes severe retardation of plant growth and compromises basal resistance. <i>Physiologia Plantarum</i> , 2006, 126, 060217072449001-???   | 5.3 | 9         |
| 58 | Genomic detection and molecular characterization of two distinct isolates of cycas necrotic stunt virus from <i>Paonia suffruticosa</i> and <i>Daphne odora</i> . <i>Virus Genes</i> , 2019, 55, 734-737.   | 1.9 | 9         |
| 59 | A Phenylpropanoid Glycoside as a Calcineurin Inhibitor Isolated from <i>Magnolia obovata</i> Thunb.. <i>Journal of Microbiology and Biotechnology</i> , 2015, 25, 1429-1432.  | 2.1 | 9         |
| 60 | Actin Cytoskeleton and Golgi Involvement in Barley stripe mosaic virus Movement and Cell Wall Localization of Triple Gene Block Proteins. <i>Plant Pathology Journal</i> , 2013, 29, 17-30.   | 1.7 | 9         |
| 61 | Molecular and Biological Characterization of a Trackable Illinois Isolate of Barley yellow dwarf virus-PAV. <i>Plant Disease</i> , 2000, 84, 483-486.   | 1.5 | 8         |
| 62 | Complete genome sequence of cnidium virus 1, a novel betanucleorhabdovirus infecting <i>Cnidium officinale</i> . <i>Archives of Virology</i> , 2022, 167, 973-977.  | 1.9 | 8         |
| 63 | Complete genome sequence of keunjorong mosaic virus, a potyvirus from <i>Cynanchum wilfordii</i> . <i>Archives of Virology</i> , 2013, 158, 1817-1820.  | 1.9 | 7         |
| 64 | Complete Genome Sequence of Ornithogalum Mosaic Virus Infecting <i>Gladiolus</i> spp. in South Korea. <i>Genome Announcements</i> , 2016, 4, .  | 0.8 | 7         |
| 65 | Occurrence of Three Major Soybean Viruses, Soybean mosaic virus, Soybean yellow mottle mosaic virus and Soybean yellow common mosaic virus Revealed by a Nationwide Survey of Subsistence Farming Soybean Fields. <i>Research in Plant Disease</i> , 2013, 19, 319-325. | 0.8 | 7         |
| 66 | Complete genome sequence of Codonopsis torradovirus A, a novel torradovirus infecting <i>Codonopsis lanceolata</i> in South Korea. <i>Archives of Virology</i> , 2021, 166, 3473-3476.  | 1.9 | 7         |
| 67 | Toxoflavin Lyase Enzyme as a Marker for Selecting Potato Plant Transformants. <i>Bioscience, Biotechnology and Biochemistry</i> , 2012, 76, 2354-2356.  | 1.3 | 6         |
| 68 | Nucleotide sequence and genome organization of a new proposed crinivirus, tetterwort vein chlorosis virus. <i>Archives of Virology</i> , 2015, 160, 2899-2902.  | 1.9 | 6         |
| 69 | Complete Genome Sequence of <i>Rehmannia Mosaic Virus</i> Infecting <i>Rehmannia glutinosa</i> in South Korea. <i>Genome Announcements</i> , 2016, 4, .   | 0.8 | 6         |
| 70 | Complete genome analysis of a novel umbravirus-polerovirus combination isolated from <i>Ixeridium dentatum</i> . <i>Archives of Virology</i> , 2017, 162, 3893-3897.  | 1.9 | 6         |
| 71 | Complete genome sequence of cnidium closterovirus 1, a novel member of the genus <i>Closterovirus</i> infecting <i>Cnidium officinale</i> . <i>Archives of Virology</i> , 2022, 167, 1491-1494.   | 1.9 | 6         |
| 72 | Complete genome sequence of daphne virus 1, a novel cytorhabdovirus infecting <i>Daphne odora</i> . <i>Archives of Virology</i> , 2023, 168, .  | 1.9 | 6         |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 73 | Complete Genome Sequences of Grapevine Yellow Speckle Viroid 1 and Hop Stunt Viroid Assembled from the Transcriptome of <i>Ixeridium dentatum</i> Plants. <i>Genome Announcements</i> , 2015, 3, . | 0.8 | 5         |
| 74 | Complete genome sequence of a novel potyvirus, callistephus mottle virus, identified in <i>Callistephus chinensis</i> . <i>Archives of Virology</i> , 2016, 161, 3281-3283.                        | 1.9 | 5         |
| 75 | Complete genome sequence of a putative new caulimovirus which exists as endogenous pararetroviral sequences in <i>Angelica dahurica</i> . <i>Archives of Virology</i> , 2017, 162, 3837-3842.      | 1.9 | 5         |
| 76 | Complete Genome Sequence of Nectarine stem pitting-associated virus , Isolated from <i>Prunus persica</i> in Cheongdo County, South Korea. <i>Genome Announcements</i> , 2017, 5, .                | 0.8 | 5         |
| 77 | Characteristics of a Lettuce mosaic virus Isolate Infecting Lettuce in Korea. <i>Plant Pathology Journal</i> , 2014, 30, 183-187.  | 1.7 | 5         |
| 78 | Complete genome sequence of pueraria virus A, a new member of the genus Caulimovirus. <i>Archives of Virology</i> , 2022, 167, 1481-1485.  | 1.9 | 5         |
| 79 | Complete genome sequence of a South Korean isolate of <i>Brugmansia</i> mosaic virus. <i>Archives of Virology</i> , 2013, 158, 2019-2022.  | 1.9 | 4         |
| 80 | Complete genome sequence of artemisia virus B, a new polerovirus infecting <i>Artemisia princeps</i> in South Korea. <i>Archives of Virology</i> , 2021, 166, 1495-1499.                           | 1.9 | 4         |
| 81 | Inhibition of the Calcineurin Pathway by Two Flavonoids Isolated from <i>Milvusa sinensis</i> Finet & Gagnep.. <i>Journal of Microbiology and Biotechnology</i> , 2016, 26, 1696-1700.             | 2.1 | 4         |
| 82 | Beet western yellows virus (BWYV): Aspect of Outbreak and Survey, and First Complete Genome Sequence of a Korea Isolate of BWYV. <i>Research in Plant Disease</i> , 2018, 24, 276-284.             | 0.8 | 4         |
| 83 | First Report of Cotton Leafroll Dwarf Virus Infecting <i>Hibiscus syriacus</i> in South Korea. <i>Plant Disease</i> , 2022, 106, 3003.   | 1.5 | 4         |
| 84 | Production of <i>Barley yellow dwarf virus</i> antisera by DNA immunization. <i>Canadian Journal of Plant Pathology</i> , 2000, 22, 410-415.   | 1.6 | 3         |
| 85 | The first complete sequence and genome structure of daphne virus Y. <i>Archives of Virology</i> , 2016, 161, 2905-2908.  | 1.9 | 3         |
| 86 | The complete sequence and genome organization of <i>ligustrum virus A</i> , a novel carlavirus. <i>Archives of Virology</i> , 2016, 161, 3593-3596.  | 1.9 | 3         |
| 87 | First report of cherry virus A infecting <i>Prunus mume</i> in South Korea. <i>VirusDisease</i> , 2017, 28, 220-221.   | 2.1 | 3         |
| 88 | Complete genome sequence of rice virus A, a new member of the family Tombusviridae. <i>Archives of Virology</i> , 2017, 162, 3247-3250.  | 1.9 | 3         |
| 89 | Complete genome sequence of a tentative new member of the genus Badnavirus identified in <i>Codonopsis lanceolata</i> . <i>Archives of Virology</i> , 2019, 164, 1733-1737.                        | 1.9 | 3         |
| 90 | Complete genome sequence and genome organization of <i>achyranthes virus A</i> , a novel member of the genus Potyvirus. <i>Archives of Virology</i> , 2020, 165, 2695-2698.                        | 1.9 | 3         |

| #   | ARTICLE  | IF  | CITATIONS |
|-----|--|-----|-----------|
| 91  | Complete genome sequence of platycodon closterovirus 1, a novel putative member of the genus Closterovirus. Archives of Virology, 2021, 166, 2051-2054.  | 1.9 | 3         |
| 92  | Complete genome sequence of Plantago asiatica virus A, a novel putative member of the genus Ploverovirus. Archives of Virology, 2022, 167, 219-222.  | 1.9 | 3         |
| 93  | Molecular characterization of a novel umbra-like virus from Thuja orientalis (arborvitae) in South Korea. Archives of Virology, 2023, 168, .   | 1.9 | 3         |
| 94  | Qualitative and Quantitative Detection of Agricultural Microorganisms Expressing Iturin and Mop Cyclase in Soils. Journal of Agricultural and Food Chemistry, 2010, 58, 12657-12663.                       | 5.4 | 2         |
| 95  | Generation of an Infectious Clone of a New Korean Isolate of Apple chlorotic leaf spot virus Driven by Dual 35S and T7 Promoters in a Versatile Binary Vector. Plant Pathology Journal, 2017, 33, 608-613. | 1.7 | 2         |
| 96  | Characterization of Brugmansia mosaic virus Isolated from Brugmansia spp. in Korea. Research in Plant Disease, 2014, 20, 307-313.  | 0.8 | 2         |
| 97  | Construction of full-length infectious cDNA clones of two Korean isolates of turnip mosaic virus breaking resistance in Brassica napus. Archives of Virology, 2022, 167, 1157-1162.                        | 1.9 | 2         |
| 98  | Genome sequence and characterization of a novel fabavirus infecting Cirsium setidens (gondre) in South Korea. Archives of Virology, 2023, 168, .   | 1.9 | 2         |
| 99  | Nucleotide sequence and genome organization of atractylodes mottle virus, a new member of the genus Carlavirus. Archives of Virology, 2015, 160, 2895-2898.  | 1.9 | 1         |
| 100 | Complete genome sequence of a tentative new umbravirus isolated from Patrinia scabiosaefolia. Archives of Virology, 2019, 164, 2375-2378.  | 1.9 | 1         |
| 101 | Complete genome sequence of a putative novel potyvirus isolated from Platycodon grandiflorum. Archives of Virology, 2019, 164, 621-624.  | 1.9 | 1         |
| 102 | Identification and molecular characterization of a novel kudzu-infecting virus of the family Betaflexiviridae. Archives of Virology, 0, , .  | 1.9 | 1         |
| 103 | Complete genome sequence of a new virus, a tentative member of the family Geminiviridae, infecting hedge bindweed (Calystegia sepium) in South Korea. Archives of Virology, 2024, 169, .                   | 1.9 | 1         |
| 104 | Highly stereoselective metal-mediated domino aldol reactions of propiophenone enolates with heteroaromatic, aliphatic, and unsaturated aldehydes. Monatshefte für Chemie, 2016, 147, 1925-1932.            | 1.9 | 0         |
| 105 | Complete Genome Sequence of a South Korean Isolate of Habenaria mosaic virus. Genome Announcements, 2016, 4, .   | 0.8 | 0         |
| 106 | Complete Genome Sequence of a Papaya ringspot virus Isolate from South Korea That Infects Cucurbita pepo. Genome Announcements, 2017, 5, .   | 0.8 | 0         |
| 107 | Psychiatrische Behandlung eines alkoholabhÄngigen Patienten mit Lebererkrankung im Endstadium. Psychopraxis Neuropraxis, 2019, 22, 108-113.  | 0.2 | 0         |
| 108 | Complete genome sequence and genome organization of scorzonera virus A (SCoVA), a novel member of the genus Potyvirus. Archives of Virology, 2021, 166, 2901-2904.   | 1.9 | 0         |

| #   | ARTICLE   | IF  | CITATIONS |
|-----|---|-----|-----------|
| 109 | Complete genome sequence of a putative novel alphaendornavirus isolated from Fagopyrum esculentum in South Korea. Archives of Virology, 2022, , 1.                | 1.9 | 0         |
| 110 | Complete nucleotide sequence of hemisteptia virus A, a polero-like virus. Archives of Virology, 0, , .  | 1.9 | 0         |
| 111 | Complete genome sequence of Stellaria aquatica virus B, a novel polerovirus that infects Stellaria aquatica. Archives of Virology, 2023, 168, .                   | 1.9 | 0         |
| 112 | Complete genome sequence and genome characterization of a novel potyvirus from Lamprocapnos spectabilis. Archives of Virology, 2023, 168, .                       | 1.9 | 0         |
| 113 | The complete genome sequence of Stellaria aquatica virus A, a new member of the genus Alphacarmovirus, family Tombusviridae. Archives of Virology, 2023, 168, .   | 1.9 | 0         |
| 114 | Metatranscriptome-Based Analysis of Viral Incidence in Jujube (Ziziphus jujuba) in Korea. Research in Plant Disease, 2023, 29, 276-285.                           | 0.8 | 0         |
| 115 | Complete genome sequence of a new putative member of the genus Pelarspovirus that infects Quercus aliena Blume in South Korea. Archives of Virology, 2024, 169, . | 1.9 | 0         |
| 116 | Complete nucleotide sequence of chrysanthemum virus D, a polero-like virus. Archives of Virology, 2024, 169, .  | 1.9 | 0         |