

Tina Ganzenmueller

List of Publications by Citations

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47
papers

978
citations

19
h-index

30
g-index

51
ext. papers

1,275
ext. citations

6
avg, IF

4.03
L-index

#	Paper	IF	Citations
47	Chronic hepatitis e in heart transplant recipients. <i>American Journal of Transplantation</i> , 2012 , 12, 3128-338.	7	120
46	Patient, virus, and treatment-related risk factors in pediatric adenovirus infection after stem cell transplantation: results of a routine monitoring program. <i>Biology of Blood and Marrow Transplantation</i> , 2014 , 20, 250-6	4.7	62
45	Human mastadenovirus type 70: a novel, multiple recombinant species D mastadenovirus isolated from diarrhoeal faeces of a haematopoietic stem cell transplantation recipient. <i>Journal of General Virology</i> , 2015 , 96, 2734-2742	4.9	61
44	Prevalence of SARS-CoV-2 Infection in Children and Their Parents in Southwest Germany. <i>JAMA Pediatrics</i> , 2021 , 175, 586-593	8.3	61
43	Characterization of Human Cytomegalovirus Genome Diversity in Immunocompromised Hosts by Whole-Genome Sequencing Directly From Clinical Specimens. <i>Journal of Infectious Diseases</i> , 2017 , 215, 1673-1683	7	58
42	Molecular Evolution of Human Adenovirus (HAdV) Species C. <i>Scientific Reports</i> , 2019 , 9, 1039	4.9	57
41	High lethality of human adenovirus disease in adult allogeneic stem cell transplant recipients with high adenoviral blood load. <i>Journal of Clinical Virology</i> , 2011 , 52, 55-9	14.5	44
40	Comparison of the performance of direct fluorescent antibody staining, a point-of-care rapid antigen test and virus isolation with that of RT-PCR for the detection of novel 2009 influenza A (H1N1) virus in respiratory specimens. <i>Journal of Medical Microbiology</i> , 2010 , 59, 713-717	3.2	44
39	Human Cytomegalovirus Genomes Sequenced Directly From Clinical Material: Variation, Multiple-Strain Infection, Recombination, and Gene Loss. <i>Journal of Infectious Diseases</i> , 2019 , 220, 781-791	7.1	43
38	Quantification of cytomegalovirus DNA levels in intestinal biopsies as a diagnostic tool for CMV intestinal disease. <i>Journal of Clinical Virology</i> , 2009 , 46, 254-8	14.5	28
37	Next-generation sequencing fails to identify human virus sequences in cutaneous squamous cell carcinoma. <i>International Journal of Cancer</i> , 2012 , 131, E1173-9	7.5	27
36	Varicella zoster virus infections in neurological patients: a clinical study. <i>BMC Infectious Diseases</i> , 2018 , 18, 238	4	24
35	Guillain-Barré syndrome associated with autochthonous infection by hepatitis E virus subgenotype 3c. <i>Infection</i> , 2014 , 42, 171-3	5.8	24
34	Improved quantitative PCR protocols for adenovirus and CMV with an internal inhibition control system and automated nucleic acid isolation. <i>Journal of Medical Virology</i> , 2012 , 84, 890-6	19.7	24
33	Generation of high-titre virus stocks using BrK.219, a B-cell line infected stably with recombinant Kaposi's sarcoma-associated herpesvirus. <i>Journal of Virological Methods</i> , 2015 , 217, 79-86	2.6	22
32	A human adenovirus species B subtype 21a associated with severe pneumonia. <i>Journal of Infection</i> , 2014 , 69, 490-9	18.9	22
31	Detection of cytomegalovirus (CMV) by real-time PCR in fecal samples for the non-invasive diagnosis of CMV intestinal disease. <i>Journal of Clinical Virology</i> , 2014 , 61, 517-22	14.5	22

30	Hepatitis E: an emerging infectious disease in Germany?. <i>Zeitschrift Fur Gastroenterologie</i> , 2011 , 49, 1255-6	22
29	Adenoviral load diagnostics by quantitative polymerase chain reaction: techniques and application. <i>Reviews in Medical Virology</i> , 2012 , 22, 194-208	11.7 20
28	No human virus sequences detected by next-generation sequencing in benign verrucous skin tumors occurring in BRAF-inhibitor-treated patients. <i>Experimental Dermatology</i> , 2013 , 22, 725-9	4 18
27	Cerebrospinal fluid features in adults with enteroviral nervous system infection. <i>International Journal of Infectious Diseases</i> , 2018 , 68, 94-101	10.5 17
26	The E7 protein of the cottontail rabbit papillomavirus immortalizes normal rabbit keratinocytes and reduces pRb levels, while E6 cooperates in immortalization but neither degrades p53 nor binds E6AP. <i>Virology</i> , 2008 , 372, 313-24	3.6 16
25	The Influence of Blood Contamination on Cerebrospinal Fluid Diagnostics. <i>Frontiers in Neurology</i> , 2019 , 10, 584	4.1 15
24	Clinical significance of quantitative cytomegalovirus detection in bronchoalveolar lavage fluid in lung transplant recipients. <i>Transplant Infectious Disease</i> , 2013 , 15, 60-9	2.7 15
23	Three novel, multiple recombinant types of species of human mastadenovirus D (HAdV-D 73, 74 & 75) isolated from diarrhoeal faeces of immunocompromised patients. <i>Journal of General Virology</i> , 2017 , 98, 3037-3045	4.9 14
22	Molecular phylogeny of a novel human adenovirus type 8 strain causing a prolonged, multi-state keratoconjunctivitis epidemic in Germany. <i>Scientific Reports</i> , 2017 , 7, 40680	4.9 13
21	Expansion of recipient-derived antiviral T cells may influence donor chimerism after allogeneic stem cell transplantation. <i>Transplant Infectious Disease</i> , 2013 , 15, 627-33	2.7 12
20	Varicella Zoster Virus Meningitis in a Young Immunocompetent Adult without Rash: A Misleading Clinical Presentation. <i>Case Reports in Neurological Medicine</i> , 2014 , 2014, 686218	0.7 11
19	A case series of children and young people admitted to a tertiary care hospital in Germany with COVID-19. <i>BMC Infectious Diseases</i> , 2021 , 21, 133	4 7
18	Evaluation of the analytical performance and specificity of a SARS-CoV-2 transcription-mediated amplification assay. <i>Journal of Virological Methods</i> , 2021 , 294, 114182	2.6 7
17	Prevalence of SARS-CoV-2 Infection in Children and Their Parents in Southwest Germany. <i>SSRN Electronic Journal</i> ,	1 6
16	Whole-Genome Approach to Assessing Human Cytomegalovirus Dynamics in Transplant Patients Undergoing Antiviral Therapy. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020 , 10, 267	5.9 5
15	Comparison of the performance of the Panther Fusion respiratory virus panel to R-Gene and laboratory developed tests for diagnostic and hygiene screening specimens from the upper and lower respiratory tract. <i>Journal of Medical Microbiology</i> , 2020 , 69, 427-435	3.2 5
14	Outcome of pregnancies with recent primary cytomegalovirus infection in first trimester treated with hyperimmunoglobulin: observational study. <i>Ultrasound in Obstetrics and Gynecology</i> , 2021 , 57, 560-567	5.8 5
13	Haploflow: strain-resolved de novo assembly of viral genomes. <i>Genome Biology</i> , 2021 , 22, 212	18.3 5

12	Influenza and respiratory syncytial virus screening for the detection of asymptotically infected patients in hematology and oncology. <i>GMS Hygiene and Infection Control</i> , 2018 , 13, Doc08	2.4	4
11	A rolling circle amplification screen for polyomaviruses other than BKPyV in renal transplant recipients confirms high prevalence of urinary JCPyV shedding. <i>Intervirology</i> , 2015 , 58, 88-94	2.5	3
10	Prolonged detection of herpes simplex virus type 2 (HSV-2) DNA in cerebrospinal fluid despite antiviral therapy in a patient with HSV-2-associated radiculitis. <i>Antiviral Therapy</i> , 2012 , 17, 125-8	1.6	3
9	Transmission of Severe Acute Respiratory Syndrome Coronavirus 2 in Households with Children, Southwest Germany, May-August 2020. <i>Emerging Infectious Diseases</i> , 2021 , 27, 3009-3019	10.2	3
8	Human Cytomegalovirus Genomes Sequenced Directly from Clinical Material: Variation, Multiple-Strain Infection, Recombination and Mutation		3
7	Novel Virus Related to Kaposi's Sarcoma-Associated Herpesvirus from Colobus Monkey. <i>Emerging Infectious Diseases</i> , 2019 , 25, 1548-1551	10.2	2
6	Human cytomegalovirus multiple-strain infections and viral population diversity in haematopoietic stem cell transplant recipients analysed by high-throughput sequencing. <i>Medical Microbiology and Immunology</i> , 2021 , 210, 291-304	4	1
5	Reverse genetics systems for contemporary isolates of respiratory syncytial virus enable rapid evaluation of antibody escape mutants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	1
4	Human Cytomegalovirus Genome Diversity in Longitudinally Collected Breast Milk Samples. <i>Frontiers in Cellular and Infection Microbiology</i> , 2021 , 11, 664247	5.9	1
3	The probe target mutation G18913A of adenovirus type 5 is not associated with underquantification in a generic adenovirus real-time PCR. <i>Diagnostic Microbiology and Infectious Disease</i> , 2018 , 90, 71-72	2.9	
2	Virus-Associated Hemophagocytic Syndrome Contributes to Fatal Outcome In Critical Ill Patients with 2009 Influenza A (H1N1) Infection.. <i>Blood</i> , 2010 , 116, 3795-3795	2.2	
1	Persistence of Recipient-Derived Antiviral T cells Severely Influence Donor Chimerism Post Allogeneic HSCT,. <i>Blood</i> , 2011 , 118, 3240-3240	2.2	