

Elena Knops

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

466
citations

840776

11
h-index

713466

21
g-index

27
all docs

27
docs citations

27
times ranked

1128
citing authors

#	ARTICLE	IF	CITATIONS
1	Analysis of antibodies from HCV elite neutralizers identifies genetic determinants of broad neutralization. <i>Immunity</i> , 2022, 55, 341-354.e7.	14.3	21
2	SARS-CoV-2 neutralizing antibody treatment in patients with COVID-19 and immunodeficiency due to B-cell non-Hodgkin lymphoma. <i>Blood Advances</i> , 2022, 6, 1580-1584.	5.2	8
3	CXCR3 Expression Pattern on CD4+ T Cells and IP-10 Levels with Regard to the HIV-1 Reservoir in the Gut-Associated Lymphatic Tissue. <i>Pathogens</i> , 2022, 11, 483.	2.8	4
4	Detection of SARS-CoV-2 viremia before onset of COVID-19 symptoms in an allo-transplanted patient with acute leukemia. <i>Bone Marrow Transplantation</i> , 2021, 56, 716-719.	2.4	20
5	Combined Therapy with Intravenous Immunoglobulins, Letermovir and (Val-)Ganciclovir in Complicated Courses of CMV-Infection in Transplant Recipients. <i>Microorganisms</i> , 2021, 9, 1666.	3.6	1
6	Characterization of HIV-1 drug resistance among patients with failure of second-line combined antiretroviral therapy in central Ethiopia. <i>HIV Medicine</i> , 2021, , .	2.2	1
7	Safe and effective pool testing for SARS-CoV-2 detection. <i>Journal of Clinical Virology</i> , 2021, 145, 105018.	3.1	9
8	RNAemia Corresponds to Disease Severity and Antibody Response in Hospitalized COVID-19 Patients. <i>Viruses</i> , 2020, 12, 1045.	3.3	53
9	Vulnerability to reservoir reseeded due to high immune activation after allogeneic hematopoietic stem cell transplantation in individuals with HIV-1. <i>Science Translational Medicine</i> , 2020, 12, .	12.4	17
10	Molecular Characterization and Clinical Description of Non-Polio Enteroviruses Detected in Stool Samples from HIV-Positive and HIV-Negative Adults in Ghana. <i>Viruses</i> , 2020, 12, 221.	3.3	8
11	HIV-1 Sub-Subtype A6: Settings for Normalised Identification and Molecular Epidemiology in the Southern Federal District, Russia. <i>Viruses</i> , 2020, 12, 475.	3.3	20
12	Lower prevalence of <i>Blastocystis</i> sp. infections in HIV positive compared to HIV negative adults in Ghana. <i>PLoS ONE</i> , 2019, 14, e0221968.	2.5	11
13	1269. Cohort Profile: The Translational Platform HIV (TP-HIV), a Multicenter Cohort Project in Germany. <i>Open Forum Infectious Diseases</i> , 2019, 6, S456-S457.	0.9	0
14	Hotspots of Transmission Driving the Local Human Immunodeficiency Virus Epidemic in the Cologne-Bonn Region, Germany. <i>Clinical Infectious Diseases</i> , 2019, 68, 1539-1546.	5.8	11
15	geno2pheno[ngs-freq]: a genotypic interpretation system for identifying viral drug resistance using next-generation sequencing data. <i>Nucleic Acids Research</i> , 2018, 46, W271-W277.	14.5	37
16	Epistatic Interactions in NS5A of Hepatitis C Virus Suggest Drug Resistance Mechanisms. <i>Genes</i> , 2018, 9, 343.	2.4	12
17	Dynamics of BKP γ reactivation and risk of hemorrhagic cystitis after allogeneic hematopoietic stem cell transplantation. <i>European Journal of Haematology</i> , 2017, 99, 133-140.	2.2	11
18	Results of the first international HIV-1 coreceptor proficiency panel test. <i>Journal of Clinical Virology</i> , 2017, 93, 53-56.	3.1	4

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19	Dynamics of Torque Teno virus viremia could predict risk of complications after allogeneic hematopoietic stem cell transplantation. <i>Medical Microbiology and Immunology</i> , 2017, 206, 355-362.	4.8	39
20	HCV Resistance Profile Evolution in a GT1b, DAA-Naive Patient Before, On, and After Failing Triple DAA Therapy. <i>Clinical Gastroenterology and Hepatology</i> , 2017, 15, 307-309.	4.4	1
21	Geno2pheno[HCV] – A Web-based Interpretation System to Support Hepatitis C Treatment Decisions in the Era of Direct-Acting Antiviral Agents. <i>PLoS ONE</i> , 2016, 11, e0155869.	2.5	101
22	Development of a phenotypic susceptibility assay for HIV-1 integrase inhibitors. <i>Journal of Virological Methods</i> , 2016, 238, 29-37.	2.1	6
23	Detection and characterization of enteroviruses and parechoviruses in healthy people living in the South of Côte d'Ivoire. <i>Journal of Clinical Virology</i> , 2015, 71, 40-43.	3.1	37
24	Mutational patterns in the frameshift-regulating site of HIV-1 selected by protease inhibitors. <i>Medical Microbiology and Immunology</i> , 2012, 201, 213-218.	4.8	15
25	Evolution of protease inhibitor resistance in the gag and pol genes of HIV subtype G isolates. <i>Journal of Antimicrobial Chemotherapy</i> , 2010, 65, 1472-1476.	3.0	9
26	The evolution of protease mutation 76V is associated with protease mutation 46I and gag mutation 431V. <i>Aids</i> , 2010, 24, 779-781.	2.2	9