Agnieszka PaweÅ,czyk

List of Publications by Year in descending order

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Version: 2024-02-01

46 papers 888 citations

567281 15 h-index 28 g-index

52 all docs 52 docs citations

times ranked

52

1287 citing authors

#	Article	IF	CITATIONS
1	Seronegative Infection with Toxoplasma gondii in Asymptomatic Human Immunodeficiency Virus Type 1 (HIV-1)-Infected Patients and in Blood Donors. Journal of Clinical Medicine, 2022, 11, 638.	2.4	6
2	Long-term study of Borrelia and Babesia prevalence and co-infection in Ixodes ricinus and Dermacentor recticulatus ticks removed from humans in Poland, 2016–2019. Parasites and Vectors, 2021, 14, 348.	2.5	10
3	Search for Viral Infections in Cerebrospinal Fluid From Patients With Autoimmune Encephalitis. Open Forum Infectious Diseases, 2020, 7, ofaa468.	0.9	6
4	Next-generation sequencing in the diagnosis of viral encephalitis: sensitivity and clinical limitations. Scientific Reports, 2020, 10, 16173.	3.3	23
5	Search for viral agents in cerebrospinal fluid in patients with multiple sclerosis using real-time PCR and metagenomics. PLoS ONE, 2020, 15, e0240601.	2.5	5
6	Hepatitis C virus (HCV) genotype 1b displays higher genetic variability of hypervariable region 1 (HVR1) than genotype 3. Scientific Reports, 2019, 9, 12846.	3.3	4
7	Expression of programmed cell death protein 1 and Tâ€cell immunoglobulinâ€and mucinâ€domainâ€containing moleculeâ€3 on peripheral blood CD4+CD8+ double positive T cells in patients with chronic hepatitis C virus infection and in subjects who spontaneously cleared the virus. Journal of Viral Hepatitis, 2019, 26. 942-950.	2.0	14
8	Seroprevalence of six pathogens transmitted by the Ixodes ricinus ticks in asymptomatic individuals with HIV infection and in blood donors. Scientific Reports, 2019, 9, 2117.	3.3	13
9	TICK-BORNE PATHOGENS IN INDIVIDUALS WITH HUMAN IMMUNODEFICIENCY VIRUS TYPE 1 (HIV-1) INFECTION. Postepy Mikrobiologii, 2019, 57, 251-259.	0.1	O
10	Human Pegivirus in Patients with Encephalitis of Unclear Etiology, Poland. Emerging Infectious Diseases, 2018, 24, 1785-1794.	4.3	15
11	Molecular identification of tick-borne pathogens in asymptomatic individuals with human immunodeficiency virus type 1 (HIV-1) infection: a retrospective study. BMC Infectious Diseases, 2018, 18, 227.	2.9	5
12	Next-generation sequencing analysis of a cluster of hepatitis C virus infections in a haematology and oncology center. PLoS ONE, 2018, 13, e0194816.	2.5	8
13	Selected aspects of helminth infections Schistosoma sp., Ascaris lumbricoides, Strongyloides stercoralis in individuals diagnosed with human immunodeficiency virus (HIV) infection. Przeglad Epidemiologiczny, 2018, 72, 349-361.	0.2	2
14	A Cluster of Fatal Tick-borne Encephalitis Virus Infection in Organ Transplant Setting. Journal of Infectious Diseases, 2017, 215, 896-901.	4.0	67
15	Sensitivity of Next-Generation Sequencing Metagenomic Analysis for Detection of RNA and DNA Viruses in Cerebrospinal Fluid: The Confounding Effect of Background Contamination. Advances in Experimental Medicine and Biology, 2017, , 53-62.	1.6	10
16	Viral etiologies in adult patients with encephalitis in Poland: A prospective single center study. PLoS ONE, 2017, 12, e0178481.	2.5	15
17	Sensitivity of Next-Generation Sequencing Metagenomic Analysis for Detection of RNA and DNA Viruses in Cerebrospinal Fluid: The Confounding Effect of Background Contamination. Advances in Experimental Medicine and Biology, 2016, , 53-62.	1.6	49
18	Spouse-to-Spouse Transmission and Evolution of Hypervariable Region 1 and 5' Untranslated Region of Hepatitis C Virus Analyzed by Next-Generation Sequencing. PLoS ONE, 2016, 11, e0150311.	2.5	4

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19	The correlation between pretreatment cytokine expression patterns in peripheral blood mononuclear cells with chronic hepatitis c outcome. BMC Infectious Diseases, 2015, 15, 556.	2.9	13
20	Next-Generation Sequencing of 5′ Untranslated Region of Hepatitis C Virus in Search of Minor Viral Variant in a Patient Who Revealed New Genotype While on Antiviral Treatment. Advances in Experimental Medicine and Biology, 2015, 885, 11-23.	1.6	3
21	Pathogens vectored by the tick, Dermacentor reticulatus, in endemic regions and zones of expansion in Poland. Parasites and Vectors, 2015, 8, 490.	2.5	62
22	Genetic Variability of Hepatitis C Virus (HCV) 5' Untranslated Region in HIV/HCV Coinfected Patients Treated with Pegylated Interferon and Ribavirin. PLoS ONE, 2015, 10, e0125604.	2.5	5
23	First report of two asymptomatic cases of human infection with <i>Babesia microti </i> (Franca, 1910) in Poland. Annals of Agricultural and Environmental Medicine, 2015, 22, 51-54.	1.0	32
24	Analysis of Genotype 1b Hepatitis C Virus IRES in Serum and Peripheral Blood Mononuclear Cells in Patients Treated with Interferon and Ribavirin. BioMed Research International, 2014, 2014, 1-7.	1.9	1
25	Evidence for immune activation in patients with residual hepatitis C virus RNA long after successful treatment with IFN and ribavirin. Journal of General Virology, 2014, 95, 2004-2009.	2.9	14
26	Seronegative hepatitis C virus infection in patients with lymphoproliferative disorders. Journal of Viral Hepatitis, 2014, 21, 424-429.	2.0	7
27	Seronegative Hepatitis C Virus Infection. Archivum Immunologiae Et Therapiae Experimentalis, 2014, 62, 145-151.	2.3	17
28	Deep sequencing of hepatitis C virus hypervariable region 1 reveals no correlation between genetic heterogeneity and antiviral treatment outcome. BMC Infectious Diseases, 2014, 14, 389.	2.9	9
29	Hepatitis C virus $5\hat{a} \in \mathbb{R}^2$ untranslated region variability correlates with treatment outcome. Journal of Viral Hepatitis, 2014, 21, 551-559.	2.0	4
30	Detection of hepatitis C virus (HCV) negative strand RNA and NS3 protein in peripheral blood mononuclear cells (PBMC): CD3+, CD14+ and CD19+. Virology Journal, 2013, 10, 346.	3.4	39
31	Hepatitis C virus (HCV) infection of peripheral blood mononuclear cells in patients with type II cryoglobulinemia. Human Immunology, 2013, 74, 1559-1562.	2.4	10
32	Hepatitis G virus/GBV-C in serum, peripheral blood mononuclear cells and bone marrow in patients with hematological malignancies. Infection, Genetics and Evolution, 2013, 19, 195-199.	2.3	13
33	Ultradeep Pyrosequencing of Hepatitis C Virus Hypervariable Region 1 in Quasispecies Analysis. BioMed Research International, 2013, 2013, 1-10.	1.9	18
34	Variability of hepatitis C virus hypervariable region 1 (HVR-1) during the early phase of pegylated interferon and ribavirin therapy. Advances in Medical Sciences, 2012, 57, 370-374.	2.1	3
35	Progress in the detection of productive HCV infection – the presence of the non-structural NS3 protein in peripheral blood mononuclear cell (PBMC). Experimental and Clinical Hepatology, 2011, 7, 16-19.	0.3	0
36	Diversity of Thalassemia Variants in Poland – Screening by Real-Time PCR. Acta Haematologica, 2008, 120, 153-157.	1.4	4

#	Article	IF	Citations
37	Babesia microti: Prevalence in wild rodents and Ixodes ricinus ticks from the Mazury Lakes District of north-eastern Poland. International Journal of Medical Microbiology, 2006, 296, 137-143.	3.6	65
38	Differential display analysis of gene expression in brains from hepatitis C-infected patients. Aids, 2005, 19, S145-S150.	2.2	14
39	Medium-term temporal stability of the helminth component community structure in bank voles (Clethrionomys glareolus) from the Mazury Lake District region of Poland. Parasitology, 2005, 130, 213-228.	1.5	44
40	Evidence for Viral Persistence in Patients Who Test Positive for Anti–Hepatitis C Virus Antibodies and Have Normal Alanine Aminotransferase Levels. Journal of Infectious Diseases, 2005, 191, 1730-1733.	4.0	42
41	Factors affecting the component community structure of haemoparasites in common voles () Tj ETQq1 1 0.7843 270-284.	14 rgBT /0 1.6	Overlock 10 55
42	The zoonotic reservoir of Borrelia burgdorferi sensu lato in the Mazury Lakes district of North-Eastern Poland. International Journal of Medical Microbiology Supplements, 2004, 293, 167-171.	0.4	4
43	Parasites of chaffinch (Fringilla coelebs) population. Part II. Blood parasites. Annals of Parasitology, 2003, 49, 31-8.	0.1	1
44	Prevalence and abundance of Cryptosporidium parvum and Giardia spp. in wild rural rodents from the Mazury Lake District region of Poland. Parasitology, 2002, 125, 21-34.	1.5	68
45	Factors affecting the component community structure of haemoparasites in bank voles (Clethrionomys glareolus) from the Mazury Lake District region of Poland. Parasitology, 2001, 122, 43-54.	1.5	70
46	Detection of reservoirs for Lyme borreliosis in the Mazury Lakes District, Poland. Zentralblatt Fur Bakteriologie: International Journal of Medical Microbiology, 1999, 289, 698-703.	0.5	4