Beata Kasztelewicz

List of Publications by Year
in descending order

27	249	933447 10	996975 15
papers	citations	h-index	g-index

28	28	28	463
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Prevalence of IgG antibodies against SARS-CoV-2 among healthcare workers in a tertiary pediatric hospital in Poland. PLoS ONE, 2021, 16, e0249550.	2.5	14
2	The Limitations of Cytomegalovirus DNA Detection in Cerebrospinal Fluid of Newborn Infants With Congenital CMV Infection: A Tertiary Care Neonatal Center Experience. Pediatric Infectious Disease Journal, 2021, 40, 838-845.	2.0	6
3	Single Nucleotide Polymorphisms of Interleukins and Toll-like Receptors and Neuroimaging Results in Newborns with Congenital HCMV Infection. Viruses, 2021, 13, 1783.	3.3	0
4	Association between single nucleotide polymorphisms and viral load in congenital cytomegalovirus infection. Medycyna Wieku Rozwojowego, 2021, 24, 9-17.	0.2	2
5	Association between single nucleotide polymorphisms (SNPs) of IL1, IL12, IL28 and TLR4 and symptoms of congenital cytomegalovirus infection. PLoS ONE, 2020, 15, e0233096.	2.5	5
6	Antiviral treatment in congenital HCMV infection: The six-year experience of a single neonatal center in Poland. Advances in Clinical and Experimental Medicine, 2020, 29, 1161-1167.	1.4	4
7	Title is missing!. , 2020, 15, e0233096.		0
8	Title is missing!. , 2020, 15, e0233096.		0
9	Title is missing!. , 2020, 15, e0233096.		0
10	Title is missing!. , 2020, 15, e0233096.		0
11	Title is missing!. , 2020, 15, e0233096.		0
12	Title is missing!. , 2020, 15, e0233096.		0
13	Distribution of the CMV glycoprotein gH/gL/gO and gH/gL/pUL128/pUL130/pUL131A complex variants and associated clinical manifestations in infants infected congenitally or postnatally. Scientific Reports, 2019, 9, 16352.	3.3	11
14	Cytokine gene polymorphism associations with congenital cytomegalovirus infection and sensorineural hearing loss. European Journal of Clinical Microbiology and Infectious Diseases, 2017, 36, 1811-1818.	2.9	18
15	Lyme Neuroborreliosis (LNB) â€“ Clinical and Diagnostic Difficulties. , 2017, 26, 19-24.		0
16	Assessment of interleukin-17A, C5a and RANTES for early diagnosis of neonatal sepsis â€“ a preliminary study. Central-European Journal of Immunology, 2016, 4, 376-382.	1.2	4
17	Human cytomegalovirus UL55, UL144, and US28 genotype distribution in infants infected congenitally or postnatally. Journal of Medical Virology, 2015, 87, 1737-1748.	5.0	18
18	Cytomegalovirus glycoprotein H genotype distribution and the relationship with hearing loss in children. Journal of Medical Virology, 2014, 86, 1421-1427.	5.0	23

#	Article	IF	Citations
19	Cytomegalovirus alpha-chemokine genotypes are associated with clinical manifestations in children with congenital or postnatal infections. Virology, 2014, 462-463, 207-217.	2.4	12
20	Distribution of cytomegalovirus gN variants and associated clinical sequelae in infants. Journal of Clinical Virology, 2013, 58, 271-275.	3.1	27
21	The impact of cytokine gene polymorphisms on Epsteinâ€“Barr virus infection outcome in pediatric liver transplant recipients. Journal of Clinical Virology, 2012, 55, 226-232.	3.1	10
22	Mannan-binding lectin-2 (MBL2) gene polymorphisms in prenatal and perinatal cytomegalovirus infections. Molecular Immunology, 2011, 48, 2203-2206.	2.2	10
23	Epsteinâ€“Barr virus DNA load in peripheral blood mononuclear cells and whole blood from pediatric transplant recipients. Transplant Infectious Disease, 2011, 13, 471-479.	1.7	12
24	Epsteinâ€“Barr virus gene expression and latent membrane protein 1 gene polymorphism in pediatric liver transplant recipients. Journal of Medical Virology, 2011, 83, 2182-2190.	5.0	6
25	Nijmegen breakage syndrome: Long-term monitoring of viral and immunological biomarkers in peripheral blood before development of malignancy. Clinical Immunology, 2010, 135, 440-447.	3.2	25
26	Long-term monitoring of Epstein-Barr virus DNA load and humoral parameter abnormalities in pediatric liver transplant recipients before development of malignancy. Pediatric Transplantation, 2010, 14, 629-635.	1.0	6
27	Analysis of eotaxin 1/CCL11, eotaxin 2/CCL24 and eotaxin 3/CCL26 expression in lesional and non-lesional skin of patients with atopic dermatitis. Cytokine, 2010, 50, 181-185.	3.2	36