GermÃ;n Añez

List of Publications by Year in descending order

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516710 580821 25 701 16 25 citations g-index h-index papers 27 27 27 1413 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Epidemiological Scenario of Dengue in Brazil. BioMed Research International, 2015, 2015, 1-13.	1.9	87
2	Circulation of Different Lineages of Dengue Virus Type 2 in Central America, Their Evolutionary Time-Scale and Selection Pressure Analysis. PLoS ONE, 2011, 6, e27459.	2.5	59
3	Evolutionary Dynamics of West Nile Virus in the United States, 1999–2011: Phylogeny, Selection Pressure and Evolutionary Time-Scale Analysis. PLoS Neglected Tropical Diseases, 2013, 7, e2245.	3.0	59
4	Increment of interleukin 6, tumour necrosis factor alpha, nitric oxide, C-reactive protein and apoptosis in dengue. Transactions of the Royal Society of Tropical Medicine and Hygiene, 2010, 104, 16-23.	1.8	58
5	Dengue in the United States of America: A Worsening Scenario?. BioMed Research International, 2013, 2013, 1-13.	1.9	46
6	Short report: increased level of serum nitric oxide in patients with dengue American Journal of Tropical Medicine and Hygiene, 2002, 66, 762-764.	1.4	45
7	Passage of Dengue Virus Type 4 Vaccine Candidates in Fetal Rhesus Lung Cells Selects Heparin-Sensitive Variants That Result in Loss of Infectivity and Immunogenicity in Rhesus Macaques. Journal of Virology, 2009, 83, 10384-10394.	3.4	41
8	Ultrastructural studies on dengue virus type 2 infection of cultured human monocytes. Virology Journal, 2005, 2, 26.	3.4	30
9	Differential Induction of Cytokines by Human Neonatal, Adult, and Elderly Monocyte/Macrophages Infected with Dengue Virus. Viral Immunology, 2014, 27, 151-159.	1.3	29
10	Phylogenetic Analysis of Dengue Virus Types 1 and 4 Circulating in Puerto Rico and Key West, Florida, during 2010 Epidemics. American Journal of Tropical Medicine and Hygiene, 2012, 87, 548-553.	1.4	26
11	Differential Oxidative Stress Induced by Dengue Virus in Monocytes from Human Neonates, Adult and Elderly Individuals. PLoS ONE, 2013, 8, e73221.	2.5	24
12	Immunogenicity and safety of a booster dose of a quadrivalent meningococcal tetanus toxoid-conjugate vaccine (MenACYW-TT) in adolescents and adults: a Phase III randomized study. Human Vaccines and Immunotherapeutics, 2020, 16, 1292-1298.	3.3	23
13	Complete Genome Sequences of Dengue Virus Type 1 to 4 Strains Used for the Development of CBER/FDA RNA Reference Reagents and WHO International Standard Candidates for Nucleic Acid Testing. Genome Announcements, 2016 , 4 , .	0.8	20
14	Collaborative study to establish World Health Organization international reference reagents for dengue virus Types 1 to 4 RNA for use in nucleic acid testing. Transfusion, 2017, 57, 1977-1987.	1.6	17
15	Differential Pattern of Soluble Immune Markers in Asymptomatic Dengue, West Nile and Zika Virus Infections. Scientific Reports, 2019, 9, 17172.	3.3	16
16	Dengue virus and other arboviruses: a global view of risks. ISBT Science Series, 2012, 7, 274-282.	1.1	15
17	Distribution of Dengue Virus Types 1 and 4 in Blood Components from Infected Blood Donors from Puerto Rico. PLoS Neglected Tropical Diseases, 2016, 10, e0004445.	3.0	14
18	Standardized methods to generate mock (spiked) clinical specimens by spiking blood or plasma with cultured pathogens. Journal of Applied Microbiology, 2016, 120, 1119-1129.	3.1	13

#	Article	IF	CITATIONS
19	Genetic Variability of West Nile Virus in U.S. Blood Donors from the 2012 Epidemic Season. PLoS Neglected Tropical Diseases, 2016, 10, e0004717.	3.0	12
20	Highly Multiplex Real-Time PCR–Based Screening for Blood-Borne Pathogens on an OpenArray Platform. Journal of Molecular Diagnostics, 2017, 19, 549-560.	2.8	8
21	Collaborative study for the characterization of a chikungunya virus <scp>RNA</scp> reference reagent for use in nucleicÂacid testing. Vox Sanguinis, 2015, 109, 312-318.	1.5	7
22	Complete Coding Region Sequence of a Chikungunya Virus Strain Used for Formulation of CBER/FDA RNA Reference Reagents for Nucleic Acid Testing. Genome Announcements, 2014, 2, .	0.8	5
23	Genetic Analysis of West Nile Virus Isolates from an Outbreak in Idaho, United States, 2006–2007. International Journal of Environmental Research and Public Health, 2013, 10, 4486-4506.	2.6	3
24	Complete Genome Sequence of West Nile Virus Strains Used for the Formulation of CBER/FDA RNA Reference Reagents and Lot Release Panels for Nucleic Acid Testing. Genome Announcements, 2014, 2, .	0.8	2
25	Flying pain. Medical Humanities, 2015, 41, e12-e12.	1.2	0