

Maria Teresa Cuevas

List of Publications by Year in descending order

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42
papers

1,028
citations

430874

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454955

30
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43
times ranked

1225
citing authors

#	ARTICLE	IF	CITATIONS
1	Diverse Large HIV-1 Non-subtype B Clusters Are Spreading Among Men Who Have Sex With Men in Spain. <i>Frontiers in Microbiology</i> , 2019, 10, 655.	3.5	31
2	HIV-1 Genetic Diversity in Recently Diagnosed Infections in Moscow: Predominance of A_{FSU}, Frequent Branching in Clusters, and Circulation of the Iberian Subtype G Variant. <i>AIDS Research and Human Retroviruses</i> , 2018, 34, 629-634.	1.1	16
3	Genome-scale analysis of evolutionary rate and selection in a fast-expanding Spanish cluster of HIV-1 subtype F1. <i>Infection, Genetics and Evolution</i> , 2018, 66, 43-47.	2.3	4
4	Genetic Diversity of HIV-1 in Tunisia. <i>AIDS Research and Human Retroviruses</i> , 2017, 33, 77-81.	1.1	11
5	Respiratory Infections by Enterovirus D68 in Outpatients and Inpatients Spanish Children. <i>Pediatric Infectious Disease Journal</i> , 2016, 35, 45-49.	2.0	16
6	Transmission dynamics of HIV-1 subtype B in the Basque Country, Spain. <i>Infection, Genetics and Evolution</i> , 2016, 40, 91-97.	2.3	11
7	Identification of an HIV-1 BG Intersubtype Recombinant Form (CRF73_BG), Partially Related to CRF14_BG, Which Is Circulating in Portugal and Spain. <i>PLoS ONE</i> , 2016, 11, e0148549.	2.5	14
8	CCR5 deficiency predisposes to fatal outcome in influenza virus infection. <i>Journal of General Virology</i> , 2015, 96, 2074-2078.	2.9	55
9	Epidemiological Surveillance of HIV-1 Transmitted Drug Resistance in Spain in 2004-2012: Relevance of Transmission Clusters in the Propagation of Resistance Mutations. <i>PLoS ONE</i> , 2015, 10, e0125699.	2.5	37
10	Phylogeny and Phylogeography of a Recent HIV-1 Subtype F Outbreak among Men Who Have Sex with Men in Spain Deriving from a Cluster with a Wide Geographic Circulation in Western Europe. <i>PLoS ONE</i> , 2015, 10, e0143325.	2.5	29
11	Characterization of an enhanced antigenic change in the pandemic 2009 H1N1 influenza virus haemagglutinin. <i>Journal of General Virology</i> , 2014, 95, 1033-1042.	2.9	10
12	Higher vaccine effectiveness in seasons with predominant circulation of seasonal influenza A(H1N1) than in A(H3N2) seasons: Test-negative case-control studies using surveillance data, Spain, 2003-2011. <i>Vaccine</i> , 2014, 32, 4404-4411.	3.8	16
13	Genetic diversity of HA1 domain of haemagglutinin gene of influenza A(H1N1)pdm09 in Tunisia. <i>Virology Journal</i> , 2013, 10, 150.	3.4	13
14	Frequency of D222G haemagglutinin mutant of pandemic (H1N1) pdm09 influenza virus in Tunisia between 2009 and 2011. <i>Diagnostic Pathology</i> , 2013, 8, 124.	2.0	6
15	Genetic diversity of Influenza A virus in 2009-2010 and 2010-2011 in Tunisia. <i>Médecine Et Maladies Infectieuses</i> , 2013, 43, 337-344.	5.0	3
16	Spread of different rhinovirus B genotypes in hospitalized children in Spain. <i>Influenza and Other Respiratory Viruses</i> , 2013, 7, 623-628.	3.4	7
17	Improvement of HIV-1 coreceptor tropism prediction by employing selected nucleotide positions of the env gene in a Bayesian network classifier. <i>Journal of Antimicrobial Chemotherapy</i> , 2013, 68, 1471-1485.	3.0	12
18	Identification of New and Unusual <i>rev</i> and <i>nef</i> Transcripts Expressed by an HIV Type 1 Primary Isolate. <i>AIDS Research and Human Retroviruses</i> , 2013, 29, 1075-1078.	1.1	3

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19	Characterization In Vitro and In Vivo of a Pandemic H1N1 Influenza Virus from a Fatal Case. PLoS ONE, 2013, 8, e53515.	2.5	29
20	Virological Surveillance of Influenza Viruses during the 2008â€™09, 2009â€™10 and 2010â€™11 Seasons in Tunisia. PLoS ONE, 2013, 8, e74064.	2.5	20
21	Haemagglutinin D222G mutation found in a fatal case of pandemic (H1N1) flu in Tunisia. Archives of Virology, 2012, 157, 1813-1814.	2.1	7
22	Genetic diversity of influenza A(H1N1)2009 virus circulating during the season 2010â€™2011 in Spain. Journal of Clinical Virology, 2012, 53, 16-21.	3.1	18
23	Substitutions in position 222 of haemagglutinin of pandemic influenza A (H1N1) 2009 viruses in Spain. Journal of Clinical Virology, 2011, 51, 75-78.	3.1	30
24	Oseltamivir-resistant pandemic influenza a (H1N1) 2009 viruses in Spain. Journal of Clinical Virology, 2011, 51, 205-208.	3.1	7
25	Short Communication: Biological and Genetic Characterization of HIV Type 1 Subtype B and Nonsubtype B Transmitted Viruses: Usefulness for Vaccine Candidate Assessment. AIDS Research and Human Retroviruses, 2010, 26, 1019-1025.	1.1	23
26	Identification of a New HIV Type 1 Circulating BF Intersubtype Recombinant Form (CRF47_BF) in Spain. AIDS Research and Human Retroviruses, 2010, 26, 827-832.	1.1	37
27	Near Full-Length Genome Characterization of a Newly Identified HIV Type 1 Subtype F Variant Circulating in St. Petersburg, Russia. AIDS Research and Human Retroviruses, 2009, 25, 1187-1191.	1.1	10
28	HIV-1 Transmission Cluster With T215D Revertant Mutation Among Newly Diagnosed Patients From the Basque Country, Spain. Journal of Acquired Immune Deficiency Syndromes (1999), 2009, 51, 99-103.	2.1	35
29	Development of a Panel of Well-Characterized Human Immunodeficiency Virus Type 1 Isolates from Newly Diagnosed Patients Including Acute and Recent Infections. AIDS Research and Human Retroviruses, 2009, 25, 93-102.	1.1	19
30	Incidence of non-B subtypes of HIV-1 in Galicia, Spain: high frequency and diversity of HIV-1 among men who have sex with men. Eurosurveillance, 2009, 14, .	7.0	20
31	Reaction-diffusion model for pattern formation in E. coli swarming colonies with slime. Physical Review E, 2005, 71, 031908.	2.1	22
32	High HIV-1 genetic diversity in Cuba. Aids, 2002, 16, 1643-1653.	2.2	46
33	Diversity of mosaic structures and common ancestry of human immunodeficiency virus type 1 BF intersubtype recombinant viruses from Argentina revealed by analysis of near full-length genome sequences. Journal of General Virology, 2002, 83, 107-119.	2.9	85
34	Identification of a Newly Characterized HIV-1 BG Intersubtype Circulating Recombinant Form in Galicia, Spain, Which Exhibits a Pseudotype-Like Virion Structure. Journal of Acquired Immune Deficiency Syndromes (1999), 2002, 29, 536-543.	2.1	92
35	Inhibitory effect against polymerase and ribonuclease activities of HIV-reverse transcriptase of the aqueous leaf extract of Terminalia triflora. Phytotherapy Research, 2002, 16, 778-780.	5.8	10
36	Biological characteristics of newly described HIV-1 BG recombinants in Spanish individuals. Aids, 2002, 16, 669-672.	2.2	11

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37	Analysis of HIV Type 1 Protease and Reverse Transcriptase Sequences from Venezuela for Drug Resistance-Associated Mutations and Subtype Classification: A UNAIDS Study. <i>AIDS Research and Human Retroviruses</i> , 2001, 17, 753-758.	1.1	33
38	HIV-1 genetic diversity in Galicia Spain: BG intersubtype recombinant viruses circulating among injecting drug users. <i>Aids</i> , 2001, 15, 509-516.	2.2	76
39	HIV-1 subtype G and BG recombinant viruses in Spanish natives: evidence of characteristic mutations in reverse transcriptase and protease. <i>Aids</i> , 2001, 15, 1907-1910.	2.2	10
40	Genotypic resistance mutations to antiretroviral drugs in HIV-1 B and non-B subtypes from Cuba. <i>Revista Panamericana De Salud Publica/Pan American Journal of Public Health</i> , 2001, 10, 174-180.	1.1	11
41	Widespread circulation of a B/F intersubtype recombinant form among HIV-1-infected individuals in Buenos Aires, Argentina. <i>Aids</i> , 2000, 14, 897.	2.2	64
42	Argentine plant extracts active against polymerase and ribonuclease H activities of HIV-1 reverse transcriptase. , 1999, 13, 206-209.		18