Miguel E Quiñones-Mateu

List of Publications by Year in descending order

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Version: 2024-02-01



#	Article	IF	CITATIONS
1	Ultraviolet-C Irradiation, Heat, and Storage as Potential Methods of Inactivating SARS-CoV-2 and Bacterial Pathogens on Filtering Facepiece Respirators. Pathogens, 2022, 11, 83.	2.8	6
2	Reduced and highly diverse peripheral HIV-1 reservoir in virally suppressed patients infected with non-B HIV-1 strains in Uganda. Retrovirology, 2022, 19, 1.	2.0	5
3	Chikungunya Virus' High Genomic Plasticity Enables Rapid Adaptation to Restrictive A549 Cells. Viruses, 2022, 14, 282.	3.3	2
4	Characterization of the First SARS-CoV-2 Isolates from Aotearoa New Zealand as Part of a Rapid Response to the COVID-19 Pandemic. Viruses, 2022, 14, 366.	3.3	7
5	Manipulation of Spray-Drying Conditions to Develop an Inhalable Ivermectin Dry Powder. Pharmaceutics, 2022, 14, 1432.	4.5	7
6	Chronic opioid use modulates human enteric microbiota and intestinal barrier integrity. Gut Microbes, 2021, 13, 1946368.	9.8	36
7	Perspective: the nose and the stomach play a critical role in the NZACE2-PÄŧari* (modified ACE2) drug treatment project of SARS-CoV-2 infection. Expert Review of Clinical Immunology, 2021, 17, 553-560.	3.0	10
8	High-level resistance to bictegravir and cabotegravir in subtype A- and D-infected HIV-1 patients failing raltegravir with multiple resistance mutations. Journal of Antimicrobial Chemotherapy, 2021, 76, 2965-2974.	3.0	13
9	Rapid Response to SARS-CoV-2 in Aotearoa New Zealand: Implementation of a Diagnostic Test and Characterization of the First COVID-19 Cases in the South Island. Viruses, 2021, 13, 2222.	3.3	4
10	Uncoupling Molecular Testing for SARS-CoV-2 From International Supply Chains. Frontiers in Public Health, 2021, 9, 808751.	2.7	2
11	Prior Case of Resistance on Dolutegravir Plus Lamivudine Dual Therapy. AIDS Research and Human Retroviruses, 2020, 36, 254-255.	1.1	2
12	Decreased Enteric Bacterial Composition and Diversity in South American Crohn's Disease Vary With the Choice of Treatment Strategy and Time Since Diagnosis. Journal of Crohn's and Colitis, 2020, 14, 791-800.	1.3	4
13	Nanoscale flow cytometry reveals interpatient variability in HIV protease activity that correlates with viral infectivity and identifies drug-resistant viruses. Scientific Reports, 2020, 10, 18101.	3.3	3
14	Accumulation of integrase strand transfer inhibitor resistance mutations confers high-level resistance to dolutegravir in non-B subtype HIV-1 strains from patients failing raltegravir in Uganda. Journal of Antimicrobial Chemotherapy, 2020, 75, 3525-3533.	3.0	12
15	<i>In Vivo</i> Emergence of a Novel Protease Inhibitor Resistance Signature in HIV-1 Matrix. MBio, 2020, 11, .	4.1	11
16	Predictors of first-line antiretroviral therapy failure among adults and adolescents living with HIV/AIDS in a large prevention and treatment program in Nigeria. AIDS Research and Therapy, 2020, 17, 64.	1.7	5
17	Performance comparison of next generation sequencing analysis pipelines for HIV-1 drug resistance testing. Scientific Reports, 2020, 10, 1634.	3.3	45
18	The case for New Zealand to have its own COVID-19 vaccine programme. New Zealand Medical Journal, 2020, 133, 112-115.	0.5	2

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19	Inhaled modified angiotensin converting enzyme 2 (ACE2) as a decoy to mitigate SARS-CoV-2 infection. New Zealand Medical Journal, 2020, 133, 112-118.	0.5	7
20	First-line HIV treatment failures in non-B subtypes and recombinants: a cross-sectional analysis of multiple populations in Uganda. AIDS Research and Therapy, 2019, 16, 3.	1.7	8
21	HIV/AIDS in Sierra Leone: Characterizing the Hidden Epidemic. AIDS Reviews, 2019, 20, 104-113.	1.0	16
22	Absence of HIV-1 Drug Resistance Mutations Supports the Use of Dolutegravir in Uganda. AIDS Research and Human Retroviruses, 2018, 34, 404-414.	1.1	23
23	Tapering Courses of Oral Vancomycin Induce Persistent Disruption of the Microbiota That Provide Colonization Resistance to Clostridium difficile and Vancomycin-Resistant Enterococci in Mice. Antimicrobial Agents and Chemotherapy, 2018, 62, .	3.2	23
24	Characterization of minority HIV-1 drug resistant variants in the United Kingdom following the verification of a deep sequencing-based HIV-1 genotyping and tropism assay. AIDS Research and Therapy, 2018, 15, 18.	1.7	17
25	Emergence of Resistance to Colistin During the Treatment of Bloodstream Infection Caused by Klebsiella pneumoniae Carbapenemase–Producing Klebsiella pneumoniae. Open Forum Infectious Diseases, 2018, 5, ofy054.	0.9	11
26	Increased replication capacity following evolution of PYxE insertion in Gagâ€p6 is associated with enhanced virulence in HIVâ€I subtype C from East Africa. Journal of Medical Virology, 2017, 89, 106-111.	5.0	12
27	HIV-1 strains belonging to large phylogenetic clusters show accelerated escape from integrase inhibitors in cell culture compared with viral isolates from singleton/small clusters. Journal of Antimicrobial Chemotherapy, 2017, 72, 2171-2183.	3.0	8
28	Prion seeding activity and infectivity in skin samples from patients with sporadic Creutzfeldt-Jakob disease. Science Translational Medicine, 2017, 9, .	12.4	103
29	Impaired human immunodeficiency virus type 1 replicative fitness in atypical viremic non-progressor individuals. AIDS Research and Therapy, 2017, 14, 15.	1.7	9
30	Sensitive detection of HIV-1 resistance to Zidovudine and impact on treatment outcomes in low- to middle-income countries. Infectious Diseases of Poverty, 2017, 6, 163.	3.7	11
31	Low-Frequency Drug Resistance in HIV-Infected Ugandans on Antiretroviral Treatment Is Associated with Regimen Failure. Antimicrobial Agents and Chemotherapy, 2016, 60, 3380-3397.	3.2	49
32	Novel high throughput pooled shRNA screening identifies NQO1 as a potential drug target for host directed therapy for tuberculosis. Scientific Reports, 2016, 6, 27566.	3.3	16
33	Identification of Variants in Primary and Recurrent Glioblastoma Using a Cancer-Specific Gene Panel and Whole Exome Sequencing. PLoS ONE, 2015, 10, e0124178.	2.5	16
34	Contribution of Human Immunodeficiency Virus Type 1 Minority Variants to Reduced Drug Susceptibility in Patients on an Integrase Strand Transfer Inhibitor-Based Therapy. PLoS ONE, 2014, 9, e104512.	2.5	12
35	HIV-1 and GBV-C co-infection in Venezuela. Journal of Infection in Developing Countries, 2014, 8, 863-868.	1.2	6
36	Next-Generation Sequencing to Help Monitor Patients Infected with HIV: Ready for Clinical Use?. Current Infectious Disease Reports, 2014, 16, 401.	3.0	28

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37	Sensitive Deep-Sequencing-Based HIV-1 Genotyping Assay To Simultaneously Determine Susceptibility to Protease, Reverse Transcriptase, Integrase, and Maturation Inhibitors, as Well as HIV-1 Coreceptor Tropism. Antimicrobial Agents and Chemotherapy, 2014, 58, 2167-2185.	3.2	61
38	Deep sequencing: Becoming a critical tool in clinical virology. Journal of Clinical Virology, 2014, 61, 9-19.	3.1	123
39	Sensitive Cell-Based Assay for Determination of Human Immunodeficiency Virus Type 1 Coreceptor Tropism. Journal of Clinical Microbiology, 2013, 51, 1517-1527.	3.9	18
40	Resistance Mutations outside the Integrase Coding Region Have an Effect on Human Immunodeficiency Virus Replicative Fitness but Do Not Affect Its Susceptibility to Integrase Strand Transfer Inhibitors. PLoS ONE, 2013, 8, e65631.	2.5	10
41	Mucosal Transmission of Human Immunodeficiency Virus. Current HIV Research, 2012, 10, 3-8.	0.5	46
42	Editorial: [Hot Topic: Use of Microbicides to Prevent HIV Sexual Transmission (Guest Editor: Miguel E.) Tj ETQqO	0 0 rgBT /0	Overlock 10 T
43	Use of Four Next-Generation Sequencing Platforms to Determine HIV-1 Coreceptor Tropism. PLoS ONE, 2012, 7, e49602.	2.5	78
44	Virus-inhibitory peptide. Aids, 2011, 25, 1663-1664.	2.2	6
45	Identification of low-molecular weight inhibitors of HIV-1 reverse transcriptase using a cell-based high-throughput screening system. Antiviral Research, 2011, 91, 94-98.	4.1	9
46	Novel Method for Simultaneous Quantification of Phenotypic Resistance to Maturation, Protease, Reverse Transcriptase, and Integrase HIV Inhibitors Based on 3â€2Gag(p2/p7/p1/p6)/PR/RT/INT-Recombinant Viruses: a Useful Tool in the Multitarget Era of Antiretroviral Therapy. Antimicrobial Agents and Chemotherapy, 2011, 55, 3729-3742.	3.2	23
47	Human Immunodeficiency Virus Type 1 Resistance or Cross-Resistance to Nonnucleoside Reverse Transcriptase Inhibitors Currently under Development as Microbicides. Antimicrobial Agents and Chemotherapy, 2011, 55, 3645-3645.	3.2	0
48	Human Immunodeficiency Virus Type 1 Resistance or Cross-Resistance to Nonnucleoside Reverse Transcriptase Inhibitors Currently under Development as Microbicides. Antimicrobial Agents and Chemotherapy, 2011, 55, 1403-1413.	3.2	29
49	Novel Recombinant Virus Assay for Measuring Susceptibility of Human Immunodeficiency Virus Type 1 Group M Subtypes To Clinically Approved Drugs. Journal of Clinical Microbiology, 2009, 47, 2232-2242.	3.9	13
50	Current tests to evaluate HIV-1 coreceptor tropism. Current Opinion in HIV and AIDS, 2009, 4, 136-142.	3.8	40
51	Viral Drug Resistance and Fitness. Advances in Pharmacology, 2008, 56, 257-296.	2.0	30
52	Impact on Replicative Fitness of the G48E Substitution in the Protease of HIV-1. Journal of Acquired Immune Deficiency Syndromes (1999), 2008, 48, 255-262.	2.1	6
53	Increased Levels of Human Beta-Defensins mRNA in Sexually HIV-1 Exposed But Uninfected Individuals. Current HIV Research, 2008, 6, 531-538.	0.5	74
54	HIV type 1 integrase inhibitors: from basic research to clinical implications. AIDS Reviews, 2008, 10, 172-89.	1.0	23

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55	Viral fitness: relation to drug resistance mutations and mechanisms involved: nucleoside reverse transcriptase inhibitor mutations. Current Opinion in HIV and AIDS, 2007, 2, 81-87.	3.8	7
56	The impact of viral and host elements on HIV fitness and disease progression. Current HIV/AIDS Reports, 2007, 4, 36-41.	3.1	5
57	Cystic Fibrosis and Normal Human Airway Epithelial Cell Response to Influenza A Viral Infection Journal of Interferon and Cytokine Research, 2006, 26, 609-627.	1.2	35
58	Role of Human $\hat{1}^2$ -defensins in HIV Infection. Advances in Dental Research, 2006, 19, 42-48.	3.6	84
59	HIV tropism: diagnostic tools and implications for disease progression and treatment with entry inhibitors. Aids, 2006, 20, 1359-1367.	2.2	71
60	Use of a novel assay based on intact recombinant viruses expressing green (EGFP) or red (DsRed2) fluorescent proteins to examine the contribution of pol and env genes to overall HIV-1 replicative fitness. Journal of Virological Methods, 2006, 136, 102-117.	2.1	47
61	HIV type 1 tropism and inhibitors of viral entry: clinical implications. AIDS Reviews, 2006, 8, 60-77.	1.0	30
62	Is HIV-1 evolving to a less virulent (pathogenic) virus?. Aids, 2005, 19, 1689-1690.	2.2	11
63	Combination of a mutagenic agent with a reverse transcriptase inhibitor results in systematic inhibition of HIV-1 infection. Virology, 2005, 338, 1-8.	2.4	41
64	HIV-1 mutagenesis during antiretroviral therapy: implications for successful drug treatment. Frontiers in Bioscience - Landmark, 2005, 10, 743.	3.0	8
65	The Replicative Fitness of Primary Human Immunodeficiency Virus Type 1 (HIV-1) Group M, HIV-1 Group O, and HIV-2 Isolates. Journal of Virology, 2005, 79, 8979-8990.	3.4	179
66	Persistent Replication of Human Immunodeficiency Virus Type 1 despite Treatment of Pulmonary Tuberculosis in Dually Infected Subjects. Vaccine Journal, 2005, 12, 1298-1304.	3.1	20
67	Methods to Determine HIV-1 Ex Vivo Fitness. , 2005, 304, 355-368.		15
68	Diminished Replicative Fitness of Primary Human Immunodeficiency Virus Type 1 Isolates Harboring the K65R Mutation. Journal of Clinical Microbiology, 2005, 43, 1395-1400.	3.9	76
69	Failure to Detect Human Immunodeficiency Virus Type 1 Superinfection in 28 HIV-Seroconcordant Individuals with High Risk of Reexposure to the Virus. AIDS Research and Human Retroviruses, 2004, 20, 1026-1031.	1.1	32
70	Can HIV-1 superinfection compromise antiretroviral therapy?. Aids, 2004, 18, 131-134.	2.2	16
71	Drug Resistance, Virus Fitness and HIV-1 Mutagenesis. Current Pharmaceutical Design, 2004, 10, 4065-4070.	1.9	41
72	A novel TaqMan real-time PCR assay to estimate ex vivo human immunodeficiency virus type 1 fitness in the era of multi-target (pol and env) antiretroviral therapy. Journal of General Virology, 2003, 84, 2217-2228	2.9	37

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73	Role of the Human Immunodeficiency Virus Type 1 Envelope Gene in Viral Fitness. Journal of Virology, 2003, 77, 9069-9073.	3.4	77
74	Comparing the Ex Vivo Fitness of CCR5-Tropic Human Immunodeficiency Virus Type 1 Isolates of Subtypes B and C. Journal of Virology, 2003, 77, 1021-1038.	3.4	189
75	Human epithelial β-defensins 2 and 3 inhibit HIV-1 replication. Aids, 2003, 17, F39-F48.	2.2	388
76	Role of Baseline pol Genotype in HIV-1 Fitness Evolution. Journal of Acquired Immune Deficiency Syndromes (1999), 2003, 33, 448-460.	2.1	25
77	Sorting out the complexities of HIV-1 fitness. Aids, 2003, 17, 780-781.	2.2	15
78	Fitness Variations and their Impact on the Evolution of Antiretroviral Drug Resistance. Current Drug Targets Infectious Disorders, 2003, 3, 355-371.	2.1	37
79	Reduced fitness of HIV-1 resistant to CXCR4 antagonists. Antiviral Therapy, 2003, 8, 1-8.	1.0	10
80	Reduced Fitness of HIV-1 Resistant to Cxcr4 Antagonists. Antiviral Therapy, 2003, 8, 1-8.	1.0	51
81	Human Immunodeficiency Virus Type 1 (HIV-1)Quasispecies at the Sites of Mycobacterium tuberculosis InfectionContribute to Systemic HIV-1 Heterogeneity. Journal of Virology, 2002, 76, 1697-1706.	3.4	66
82	In Vitro Intersubtype Recombinants of Human Immunodeficiency Virus Type 1: Comparison to Recent and Circulating In Vivo Recombinant Forms. Journal of Virology, 2002, 76, 9600-9613.	3.4	51
83	Insertions in the Reverse Transcriptase Increase both Drug Resistance and Viral Fitness in a Human Immunodeficiency Virus Type 1 Isolate Harboring the Multi-Nucleoside Reverse Transcriptase Inhibitor Resistance 69 Insertion Complex Mutation. Journal of Virology, 2002, 76, 10546-10552.	3.4	40
84	Fitness of drug resistant HIV-1: methodology and clinical implications. Drug Resistance Updates, 2002, 5, 224-233.	14.4	82
85	Impact of tuberculosis on HIV-1 replication, diversity, and disease progression. AIDS Reviews, 2002, 4, 165-76.	1.0	81
86	Functional Characterization of Chimeric Reverse Transcriptases with Polypeptide Subunits of Highly Divergent HIV-1 Group M and O Strains. Journal of Biological Chemistry, 2001, 276, 27470-27479.	3.4	32
87	Mechanisms Involved in Stimulation of Human Immunodeficiency Virus Type 1 Replication by Aminooxypentane RANTES. Journal of Virology, 2001, 75, 8624-8638.	3.4	40
88	Greater Diversity of HIV-1 Quasispecies in HIV-Infected Individuals With Active Tuberculosis. Journal of Acquired Immune Deficiency Syndromes (1999), 2000, 24, 408-417.	2.1	16
89	A Dual Infection/Competition Assay Shows a Correlation between Ex Vivo Human Immunodeficiency Virus Type 1 Fitness and Disease Progression. Journal of Virology, 2000, 74, 9222-9233.	3.4	224
90	Greater Diversity of HIV-1 Quasispecies in HIV-Infected Individuals With Active Tuberculosis. Journal of Acquired Immune Deficiency Syndromes (1999), 2000, 24, 408-417.	2.1	34

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91	Variable Sensitivity of CCR5-Tropic Human Immunodeficiency Virus Type 1 Isolates to Inhibition by RANTES Analogs. Journal of Virology, 2000, 74, 4868-4876.	3.4	81
92	Molecular Epidemiology of HIV Type 1 Isolates from the Czech Republic: Identification of an env E Subtype Case. AIDS Research and Human Retroviruses, 1999, 15, 85-89.	1.1	11
93	Nucleotide Diversity in Three Different Genomic Regions of Venezuelan HIV Type 1 Isolates: A Subtyping Update. AIDS Research and Human Retroviruses, 1999, 15, 73-79.	1.1	8
94	Phylogeny of HIV Type 1 Group O Isolates Based on env Gene Sequences. AIDS Research and Human Retroviruses, 1999, 15, 769-773.	1.1	11
95	LTR and tat variability of HIV-1 isolates from patients with divergent rates of disease progression. Virus Research, 1998, 57, 11-20.	2.2	27
96	Mechanisms of clinical resistance by HIV-I variants to zidovudine and the paradox of reverse transcriptase sensitivity. Drug Resistance Updates, 1998, 1, 21-28.	14.4	11
97	Analysis of pol Gene Heterogeneity, Viral Quasispecies, and Drug Resistance in Individuals Infected with Group O Strains of Human Immunodeficiency Virus Type 1. Journal of Virology, 1998, 72, 9002-9015.	3.4	64
98	3′-Azido-3′-Deoxythymidine (AZT) Mediates Cross-Resistance to Nucleoside Analogs in the Case of AZT-Resistant Human Immunodeficiency Virus Type 1 Variants. Journal of Virology, 1998, 72, 4858-4865.	3.4	21
99	Characterization of the Reverse Transcriptase of a Human Immunodeficiency Virus Type 1 Group O Isolate. Virology, 1997, 236, 364-373.	2.4	60
100	Sequence Note:envGene Characterization of the First HIV Type 1 Group O Spanish Isolate. AIDS Research and Human Retroviruses, 1996, 12, 1647-1649.	1.1	21
101	Sequence Note: env Gene Diversity of HIV Type 1 Isolates from Spain. AIDS Research and Human Retroviruses, 1996, 12, 955-957.	1.1	8
102	Point Mutant Frequencies in the <i>pol</i> Gene of Human Immunodeficiency Virus Type 1 Are Two- to Threefold Lower Than Those of <i>env</i> . AIDS Research and Human Retroviruses, 1996, 12, 1117-1128.	1.1	35
103	Activation of antigen-induced lymphocyte proliferation by interleukin-15 without the mitogenic effect of interleukin-2 that may induce human immunodeficiency virus-1 expression Journal of Clinical Investigation, 1996, 98, 616-621.	8.2	38
104	Molecular Characterization of Human Immunodeficiency Virus Type 1 Isolates from Venezuela. AIDS Research and Human Retroviruses, 1995, 11, 605-616.	1.1	23
105	Pol gene quasispecies of human immunodeficiency virus: mutations associated with drug resistance in virus from patients undergoing no drug therapy. Journal of Virology, 1995, 69, 23-31.	3.4	240