Miguel E Quiñones-Mateu

List of Publications by Year in descending order

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105 papers 4,029 citations

35 h-index 59 g-index

108 all docs

108 docs citations

108 times ranked 4339 citing authors

#	Article	IF	Citations
1	Human epithelial \hat{I}^2 -defensins 2 and 3 inhibit HIV-1 replication. Aids, 2003, 17, F39-F48.	2.2	388
2	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
3	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
4	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
5	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
6	Deep sequencing: Becoming a critical tool in clinical virology. Journal of Clinical Virology, 2014, 61, 9-19.	3.1	123
7	Prion seeding activity and infectivity in skin samples from patients with sporadic Creutzfeldt-Jakob disease. Science Translational Medicine, 2017, 9, .	12.4	103
8	Role of Human \hat{I}^2 -defensins in HIV Infection. Advances in Dental Research, 2006, 19, 42-48.	3.6	84
9	Fitness of drug resistant HIV-1: methodology and clinical implications. Drug Resistance Updates, 2002, 5, 224-233.	14.4	82
10	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
11	Impact of tuberculosis on HIV-1 replication, diversity, and disease progression. AIDS Reviews, 2002, 4, 165-76.	1.0	81
12	Use of Four Next-Generation Sequencing Platforms to Determine HIV-1 Coreceptor Tropism. PLoS ONE, 2012, 7, e49602.	2.5	78
13	Role of the Human Immunodeficiency Virus Type 1 Envelope Gene in Viral Fitness. Journal of Virology, 2003, 77, 9069-9073.	3.4	77
14	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
15	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
16	HIV tropism: diagnostic tools and implications for disease progression and treatment with entry inhibitors. Aids, 2006, 20, 1359-1367.	2.2	71
17	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
18	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

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19	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
20	Characterization of the Reverse Transcriptase of a Human Immunodeficiency Virus Type 1 Group O Isolate. Virology, 1997, 236, 364-373.	2.4	60
21	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
22	Reduced Fitness of HIV-1 Resistant to Cxcr4 Antagonists. Antiviral Therapy, 2003, 8, 1-8.	1.0	51
23	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
24	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
25	Mucosal Transmission of Human Immunodeficiency Virus. Current HIV Research, 2012, 10, 3-8.	0.5	46
26	Performance comparison of next generation sequencing analysis pipelines for HIV-1 drug resistance testing. Scientific Reports, 2020, 10, 1634.	3.3	45
27	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
28	Drug Resistance, Virus Fitness and HIV-1 Mutagenesis. Current Pharmaceutical Design, 2004, 10, 4065-4070.	1.9	41
29	Mechanisms Involved in Stimulation of Human Immunodeficiency Virus Type 1 Replication by Aminooxypentane RANTES. Journal of Virology, 2001, 75, 8624-8638.	3.4	40
30	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
31	Current tests to evaluate HIV-1 coreceptor tropism. Current Opinion in HIV and AIDS, 2009, 4, 136-142.	3.8	40
32	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
33	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
34	Fitness Variations and their Impact on the Evolution of Antiretroviral Drug Resistance. Current Drug Targets Infectious Disorders, 2003, 3, 355-371.	2.1	37
35	Chronic opioid use modulates human enteric microbiota and intestinal barrier integrity. Gut Microbes, 2021, 13, 1946368.	9.8	36
36	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

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37	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
38	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
39	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
40	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
41	Viral Drug Resistance and Fitness. Advances in Pharmacology, 2008, 56, 257-296.	2.0	30
42	HIV type 1 tropism and inhibitors of viral entry: clinical implications. AIDS Reviews, 2006, 8, 60-77.	1.0	30
43	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
44	Next-Generation Sequencing to Help Monitor Patients Infected with HIV: Ready for Clinical Use?. Current Infectious Disease Reports, 2014, 16, 401.	3.0	28
45	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
46	Role of Baseline pol Genotype in HIV-1 Fitness Evolution. Journal of Acquired Immune Deficiency Syndromes (1999), 2003, 33, 448-460.	2.1	25
47	Molecular Characterization of Human Immunodeficiency Virus Type 1 Isolates from Venezuela. AIDS Research and Human Retroviruses, 1995, 11, 605-616.	1.1	23
48	Novel Method for Simultaneous Quantification of Phenotypic Resistance to Maturation, Protease, Reverse Transcriptase, and Integrase HIV Inhibitors Based on $3\hat{a}\in^2$ Gag(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
49	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
50	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
51	HIV type 1 integrase inhibitors: from basic research to clinical implications. AIDS Reviews, 2008, 10, 172-89.	1.0	23
52	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
53	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
54	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

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55	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
56	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
57	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
58	Can HIV-1 superinfection compromise antiretroviral therapy?. Aids, 2004, 18, 131-134.	2.2	16
59	Editorial: [Hot Topic: Use of Microbicides to Prevent HIV Sexual Transmission (Guest Editor: Miguel E.) Tj ETQq $1\ 1$	0,784314 0.5	rgBT /Over
60	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
61	HIV/AIDS in Sierra Leone: Characterizing the Hidden Epidemic. AIDS Reviews, 2019, 20, 104-113.	1.0	16
62	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
63	Methods to Determine HIV-1 Ex Vivo Fitness. , 2005, 304, 355-368.		15
64	Sorting out the complexities of HIV-1 fitness. Aids, 2003, 17, 780-781.	2.2	15
65	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
66	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
67	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
68	Increased replication capacity following evolution of PYxE insertion in Gagâ€p6 is associated with enhanced virulence in HIVâ€1 subtype C from East Africa. Journal of Medical Virology, 2017, 89, 106-111.	5.0	12
69	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
70	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
71	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
72	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

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7 3	Is HIV-1 evolving to a less virulent (pathogenic) virus?. Aids, 2005, 19, 1689-1690.	2.2	11
74	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
7 5	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
76	<i>In Vivo</i> Emergence of a Novel Protease Inhibitor Resistance Signature in HIV-1 Matrix. MBio, 2020, 11, .	4.1	11
77	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
78	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
79	Reduced fitness of HIV-1 resistant to CXCR4 antagonists. Antiviral Therapy, 2003, 8, 1-8.	1.0	10
80	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
81	Impaired human immunodeficiency virus type 1 replicative fitness in atypical viremic non-progressor individuals. AIDS Research and Therapy, 2017, 14, 15.	1.7	9
82	Sequence Note: env Gene Diversity of HIV Type 1 Isolates from Spain. AIDS Research and Human Retroviruses, 1996 , 12 , 955 - 957 .	1.1	8
83	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
84	HIV-1 mutagenesis during antiretroviral therapy: implications for successful drug treatment. Frontiers in Bioscience - Landmark, 2005, 10, 743.	3.0	8
85	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
86	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
87	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
88	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
89	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
90	Manipulation of Spray-Drying Conditions to Develop an Inhalable Ivermectin Dry Powder. Pharmaceutics, 2022, 14, 1432.	4.5	7

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91	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
92	Virus-inhibitory peptide. Aids, 2011, 25, 1663-1664.	2.2	6
93	HIV-1 and GBV-C co-infection in Venezuela. Journal of Infection in Developing Countries, 2014, 8, 863-868.	1.2	6
94	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
95	The impact of viral and host elements on HIV fitness and disease progression. Current HIV/AIDS Reports, 2007, 4, 36-41.	3.1	5
96	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
97	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
98	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
99	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
100	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
101	Prior Case of Resistance on Dolutegravir Plus Lamivudine Dual Therapy. AIDS Research and Human Retroviruses, 2020, 36, 254-255.	1.1	2
102	Uncoupling Molecular Testing for SARS-CoV-2 From International Supply Chains. Frontiers in Public Health, 2021, 9, 808751.	2.7	2
103	Chikungunya Virus' High Genomic Plasticity Enables Rapid Adaptation to Restrictive A549 Cells. Viruses, 2022, 14, 282.	3.3	2
104	The case for New Zealand to have its own COVID-19 vaccine programme. New Zealand Medical Journal, 2020, 133, 112-115.	0.5	2
105	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	O