

# Peter Kojo Quashie

## List of Publications by Year in descending order

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Version: 2024-02-01

34  
papers

1,187  
citations

566801

15  
h-index

414034

32  
g-index

39  
all docs

39  
docs citations

39  
times ranked

1319  
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of the R263K Mutation in HIV-1 Integrase That Confers Low-Level Resistance to the Second-Generation Integrase Strand Transfer Inhibitor Dolutegravir. <i>Journal of Virology</i> , 2012, 86, 2696-2705.	1.5	212
2	A year of genomic surveillance reveals how the SARS-CoV-2 pandemic unfolded in Africa. <i>Science</i> , 2021, 374, 423-431.	6.0	144
3	Viral fitness cost prevents HIV-1 from evading dolutegravir drug pressure. <i>Retrovirology</i> , 2013, 10, 22.	0.9	114
4	The M50I polymorphic substitution in association with the R263K mutation in HIV-1 subtype B integrase increases drug resistance but does not restore viral replicative fitness. <i>Retrovirology</i> , 2014, 11, 7.	0.9	74
5	Differential Effects of the G118R, H51Y, and E138K Resistance Substitutions in Different Subtypes of HIV Integrase. <i>Journal of Virology</i> , 2015, 89, 3163-3175.	1.5	66
6	Evolution of HIV integrase resistance mutations. <i>Current Opinion in Infectious Diseases</i> , 2013, 26, 43-49.	1.3	63
7	Biochemical Analysis of the Role of G118R-Linked Dolutegravir Drug Resistance Substitutions in HIV-1 Integrase. <i>Antimicrobial Agents and Chemotherapy</i> , 2013, 57, 6223-6235.	1.4	62
8	Addition of E138K to R263K in HIV integrase increases resistance to dolutegravir, but fails to restore activity of the HIV integrase enzyme and viral replication capacity. <i>Journal of Antimicrobial Chemotherapy</i> , 2014, 69, 2733-2740.	1.3	47
9	Integrase strand transfer inhibitors in the management of HIV-positive individuals. <i>Annals of Medicine</i> , 2014, 46, 123-129.	1.5	43
10	Evolution of a novel pathway leading to dolutegravir resistance in a patient harbouring N155H and multiclass drug resistance. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 405-411.	1.3	35
11	Identification of a Pyridoxine-Derived Small-Molecule Inhibitor Targeting Dengue Virus RNA-Dependent RNA Polymerase. <i>Antimicrobial Agents and Chemotherapy</i> , 2016, 60, 600-608.	1.4	33
12	HIV Drug Resistance and the Advent of Integrase Inhibitors. <i>Current Infectious Disease Reports</i> , 2013, 15, 85-100.	1.3	29
13	Effect of HIV-1 Integrase Resistance Mutations When Introduced into SIVmac239 on Susceptibility to Integrase Strand Transfer Inhibitors. <i>Journal of Virology</i> , 2014, 88, 9683-9692.	1.5	22
14	Genetic diversity of SARS-CoV-2 infections in Ghana from 2020-2021. <i>Nature Communications</i> , 2022, 13, 2494.	5.8	22
15	Detection of SARS-CoV-2 intra-host recombination during superinfection with Alpha and Epsilon variants in New York City. <i>Nature Communications</i> , 2022, 13, .	5.8	22
16	Genomic analysis of SARS-CoV-2 reveals local viral evolution in Ghana. <i>Experimental Biology and Medicine</i> , 2021, 246, 960-970.	1.1	20
17	Development of a fluorescence-based HIV-1 integrase DNA binding assay for identification of novel HIV-1 integrase inhibitors. <i>Antiviral Research</i> , 2013, 98, 441-448.	1.9	17
18	Trends of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) antibody prevalence in selected regions across Ghana. <i>Wellcome Open Research</i> , 0, 6, 173.	0.9	16

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19	The R263K substitution in HIV-1 subtype C is more deleterious for integrase enzymatic function and viral replication than in subtype B. <i>Aids</i> , 2015, 29, 1459-1466.	1.0	15
20	Recommendations for empowering early career researchers to improve research culture and practice. <i>PLoS Biology</i> , 2022, 20, e3001680.	2.6	15
21	A resveratrol analog termed 3,3,4,4,5,5-hexahydroxy- <i>trans</i> -stilbene is a potent HIV-1 inhibitor. <i>Journal of Medical Virology</i> , 2015, 87, 2054-2060.	2.5	14
22	Structural Studies of the HIV-1 Integrase Protein: Compound Screening and Characterization of a DNA-Binding Inhibitor. <i>PLoS ONE</i> , 2015, 10, e0128310.	1.1	14
23	Rapid, Cheap, and Effective COVID-19 Diagnostics for Africa. <i>Diagnostics</i> , 2021, 11, 2105.	1.3	11
24	The R263K mutation in HIV integrase that is selected by dolutegravir may actually prevent clinically relevant resistance to this compound. <i>Journal of the International AIDS Society</i> , 2014, 17, 19518.	1.2	10
25	HIV-1 Group O Integrase Displays Lower Enzymatic Efficiency and Higher Susceptibility to Raltegravir than HIV-1 Group M Subtype B Integrase. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 7141-7150.	1.4	8
26	Progressive emergence of an S153F plus R263K combination of integrase mutations in the proviral DNA of one individual successfully treated with dolutegravir. <i>Journal of Antimicrobial Chemotherapy</i> , 2021, 76, 639-647.	1.3	8
27	Dolutegravir maintains a durable effect against HIV replication in tissue culture even after drug washout. <i>Journal of Antimicrobial Chemotherapy</i> , 2015, 70, 2810-2815.	1.3	7
28	A SARS-CoV-2 nucleocapsid ELISA represents a low-cost alternative to lateral flow testing for community screening in LMI countries. <i>Journal of Infection</i> , 2022, 84, 48-55.	1.7	7
29	Characterization of the Drug Resistance Profiles of Integrase Strand Transfer Inhibitors in Simian Immunodeficiency Virus SIVmac239. <i>Journal of Virology</i> , 2015, 89, 12002-12013.	1.5	6
30	Subtype-Specific Analysis of the K65R Substitution in HIV-1 That Confers Hypersusceptibility to a Novel Nucleotide-Competing Reverse Transcriptase Inhibitor. <i>Antimicrobial Agents and Chemotherapy</i> , 2015, 59, 3189-3196.	1.4	5
31	HIV-1 group O integrase displays lower susceptibility to raltegravir and has a different mutational pathway for resistance than HIV-1 group M. <i>Journal of the International AIDS Society</i> , 2014, 17, 19738.	1.2	4
32	Low COVID-19 impact in Africa: The multifactorial Nexus. <i>AAS Open Research</i> , 0, 4, 47.	1.5	4
33	Biochemical Analysis of the Role of G118R-Linked Dolutegravir Drug Resistance Substitutions in HIV-1 Integrase. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 3580-3580.	1.4	3
34	Explaining the unexpected COVID-19 trends and potential impact across Africa.. <i>F1000Research</i> , 0, 10, 1177.	0.8	0