Navaneethan Palanisamy

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7357355/publications.pdf

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25 papers 439 citations

933410 10 h-index ⁷⁵²⁶⁷⁹
20
g-index

27 all docs

27 docs citations

times ranked

27

848 citing authors

#	Article	IF	CITATIONS
1	Engineering AraC to make it responsive to light instead of arabinose. Nature Chemical Biology, 2021, 17, 817-827.	8.0	40
2	Expanding the SiMPl Plasmid Toolbox for Use with Spectinomycin/Streptomycin. ACS Omega, 2021, 6, 14148-14153.	3.5	1
3	Targeting the NS2B-NS3 protease of tick-borne encephalitis virus with pan-flaviviral protease inhibitors. Antiviral Research, 2021, 190, 105074.	4.1	12
4	Global Prevalence of Adaptive and Prolonged Infections' Mutations in the Receptor-Binding Domain of the SARS-CoV-2 Spike Protein. Viruses, 2021, 13, 1974.	3.3	9
5	Identification of a C2-symmetric diol based human immunodeficiency virus protease inhibitor targeting Zika virus NS2B-NS3 protease. Journal of Biomolecular Structure and Dynamics, 2020, 38, 5526-5536.	3.5	9
6	C-terminal eYFP fusion impairs <i>Escherichia coli</i> MinE function. Open Biology, 2020, 10, 200010.	3.6	7
7	Effect of the baseline Y93H resistance-associated substitution in HCV genotype 3 for direct-acting antiviral treatment: real-life experience from a multicenter study in Sweden and Norway. Scandinavian Journal of Gastroenterology, 2019, 54, 1042-1050.	1.5	10
8	Split intein-mediated selection of cells containing two plasmids using a single antibiotic. Nature Communications, 2019, 10, 4967.	12.8	20
9	A Multicolor Large Stokes Shift Fluorogenâ€Activating RNA Aptamer with Cationic Chromophores. Chemistry - A European Journal, 2019, 25, 1931-1935.	3.3	44
10	Biophysical Studies on HCV 1a NS3/4A Protease and Its Catalytic Triad in Wild Type and Mutants by the In Silico Approach. Interdisciplinary Sciences, Computational Life Sciences, 2018, 10, 143-156.	3.6	3
11	Protein backbone flexibility pattern is evolutionarily conserved in the Flaviviridae family: A case of NS3 protease in Flavivirus and Hepacivirus. Molecular Phylogenetics and Evolution, 2018, 118, 58-63.	2.7	8
12	Baseline dasabuvir resistance in Hepatitis C virus from the genotypes 1, 2 and 3 and modeling of the NS5B-dasabuvir complex by thein silicoapproach. Infection Ecology and Epidemiology, 2018, 8, 1528117.	0.8	5
13	Comparative genome analysis of Alkhumra hemorrhagic fever virus with Kyasanur forest disease and tick-borne encephalitis viruses by the in silico approach. Pathogens and Global Health, 2018, 112, 210-226.	2.3	5
14	Worldwide Prevalence of Baseline Resistance-Associated Polymorphisms and Resistance Mutations in HCV against Current Direct-Acting Antivirals. Antiviral Therapy, 2018, 23, 485-493.	1.0	15
15	Identification of putative drug targets and annotation of unknown proteins in Tropheryma whipplei. Computational Biology and Chemistry, 2018, 76, 130-138.	2.3	2
16	Deleterious single nucleotide polymorphisms of protein kinase R identified by the computational approach. Molecular Immunology, 2018, 101, 65-73.	2.2	9
17	Does antiretroviral treatment change HIV-1 codon usage patterns in its genes: a preliminary bioinformatics study. AIDS Research and Therapy, 2017, 14, 2.	1.7	8
18	Computational Prediction of Usutu Virus E Protein B Cell and T Cell Epitopes for Potential Vaccine Development. Scandinavian Journal of Immunology, 2017, 85, 350-364.	2.7	10

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19	Evaluation of Sofosbuvir (β-D-2′-deoxy-2′-α-fluoro-2′-β-C-methyluridine) as an inhibitor of Dengue virus replication #. Scientific Reports, 2017, 7, 6345.	3.3	58
20	SYBR Green II Dye-Based Real-Time Assay for Measuring Inhibitor Activity Against HIV-1 Reverse Transcriptase. Molecular Biotechnology, 2016, 58, 619-625.	2.4	7
21	Prevalence of polymorphisms with significant resistance to NS5A inhibitors in treatment-naive patients with hepatitis C virus genotypes 1a and 3a in Sweden. Infectious Diseases, 2015, 47, 555-562.	2.8	34
22	Influenza A Virus Encoding Secreted Gaussia Luciferase as Useful Tool to Analyze Viral Replication and Its Inhibition by Antiviral Compounds and Cellular Proteins. PLoS ONE, 2014, 9, e97695.	2.5	50
23	Transmission of hepatitis C virus among intravenous drug users in the Uppsala region of Sweden. Infection Ecology and Epidemiology, 2014, 4, 22251.	0.8	3
24	Implications of baseline polymorphisms for potential resistance to NS3 protease inhibitors in Hepatitis C virus genotypes 1a, 2b and 3a. Antiviral Research, 2013, 99, 12-17.	4.1	62
25	Nutritional status of children in rural India: a case study from Tamil Nadu, first in the world to initiate the Mid-Day Meal scheme. Health, 2011, 03, 647-655.	0.3	7