Olivier Moncorge

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7425961/publications.pdf

Version: 2024-02-01

24 papers 1,775 citations

16 h-index 25 g-index

29 all docs 29 docs citations

times ranked

29

3177 citing authors

| # | Article | IF | CITATIONS |
|----|---|-------------|-----------|
| 1 | Mitochondrial morphodynamics alteration induced by influenza virus infection as a new antiviral strategy. PLoS Pathogens, 2021, 17, e1009340. | 4.7 | 19 |
| 2 | SARS-CoV-2 Triggers an MDA-5-Dependent Interferon Response Which Is Unable To Control Replication in Lung Epithelial Cells. Journal of Virology, 2021, 95, . | 3.4 | 168 |
| 3 | Mammalian and Avian Host Cell Influenza A Restriction Factors. Viruses, 2021, 13, 522. | 3. 3 | 16 |
| 4 | Alarmin S100A9 restricts retroviral infection by limiting reverse transcription in human dendritic cells. EMBO Journal, 2021, 40, e106540. | 7.8 | 12 |
| 5 | Crystal structure of the TLDc domain of human NCOA7-AS. Acta Crystallographica Section F, Structural Biology Communications, 2021, 77, 230-237. | 0.8 | 3 |
| 6 | Regulation of influenza A virus mRNA splicing by CLK1. Antiviral Research, 2019, 168, 187-196. | 4.1 | 21 |
| 7 | TRIM8 is required for virus-induced IFN response in human plasmacytoid dendritic cells. Science Advances, 2019, 5, eaax3511. | 10.3 | 40 |
| 8 | The interferon-inducible isoform of NCOA7 inhibits endosome-mediated viral entry. Nature Microbiology, 2018, 3, 1369-1376. | 13.3 | 54 |
| 9 | Species difference in ANP32A underlies influenza A virus polymerase host restriction. Nature, 2016, 529, 101-104. | 27.8 | 228 |
| 10 | Involvement of an Arginine Triplet in M1 Matrix Protein Interaction with Membranes and in M1 Recruitment into Virus-Like Particles of the Influenza A(H1N1)pdm09 Virus. PLoS ONE, 2016, 11, e0165421. | 2.5 | 20 |
| 11 | Viral determinants of influenza A virus host range. Journal of General Virology, 2014, 95, 1193-1210. | 2.9 | 132 |
| 12 | Transfer of the Amino-Terminal Nuclear Envelope Targeting Domain of Human MX2 Converts MX1 into an HIV-1 Resistance Factor. Journal of Virology, 2014, 88, 9017-9026. | 3.4 | 87 |
| 13 | The Effect of the PB2 Mutation 627K on Highly Pathogenic H5N1 Avian Influenza Virus Is Dependent on the Virus Lineage. Journal of Virology, 2013, 87, 9983-9996. | 3.4 | 56 |
| 14 | Human MX2 is an interferon-induced post-entry inhibitor of HIV-1 infection. Nature, 2013, 502, 559-562. | 27.8 | 505 |
| 15 | Investigation of Influenza Virus Polymerase Activity in Pig Cells. Journal of Virology, 2013, 87, 384-394. | 3.4 | 46 |
| 16 | Unstable Polymerase-Nucleoprotein Interaction Is Not Responsible for Avian Influenza Virus Polymerase Restriction in Human Cells. Journal of Virology, 2013, 87, 1278-1284. | 3.4 | 41 |
| 17 | Different effects of the TAR structure on HIV-1 and HIV-2 genomic RNA translation. Nucleic Acids Research, 2012, 40, 2653-2667. | 14.5 | 38 |
| 18 | Characterization of Peptide Aptamers Targeting Bfl-1 Anti-Apoptotic Protein. Biochemistry, 2011, 50, 5120-5129. | 2.5 | 12 |

| # | Article | IF | CITATION |
|----|---|------|----------|
| 19 | Evidence for Avian and Human Host Cell Factors That Affect the Activity of Influenza Virus Polymerase. Journal of Virology, 2010, 84, 9978-9986. | 3.4 | 88 |
| 20 | Rapid generation of a well-matched vaccine seed from a modern influenza A virus primary isolate without recourse to eggs. Vaccine, 2010, 28, 2973-2979. | 3.8 | 9 |
| 21 | A RasGAP SH3 Peptide Aptamer Inhibits RasGAP-Aurora Interaction and Induces Caspase-Independent Tumor Cell Death. PLoS ONE, 2008, 3, e2902. | 2.5 | 14 |
| 22 | A Comparative Analysis of Perturbations Caused by a Gene Knock-out, a Dominant Negative Allele, and a Set of Peptide Aptamers. Molecular and Cellular Proteomics, 2007, 6, 2110-2121. | 3.8 | 19 |
| 23 | Back to basics: the untreated rabbit reticulocyte lysate as a competitive system to recapitulate cap/poly(A) synergy and the selective advantage of IRES-driven translation. Nucleic Acids Research, 2007, 35, e121-e121. | 14.5 | 60 |
| 24 | Selection and characterization of large collections of peptide aptamers through optimized yeast two-hybrid procedures. Nature Protocols, 2006, 1, 1066-1091. | 12.0 | 50 |