Aleksandr Ianevski

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6118740/publications.pdf

Version: 2024-02-01

318942 299063 2,802 46 23 42 citations h-index g-index papers 59 59 59 5331 docs citations times ranked citing authors all docs

| # | Article | IF | Citations |
|----|--|-----|-----------|
| 1 | Integrated root phenotypes for improved rice performance under low nitrogen availability. Plant, Cell and Environment, 2022, 45, 805-822. | 2.8 | 23 |
| 2 | Targeting Apoptosis Pathways With BCL2 and MDM2 Inhibitors in Adult B-cell Acute Lymphoblastic Leukemia. HemaSphere, 2022, 6, e701. | 1.2 | 4 |
| 3 | Fully-automated and ultra-fast cell-type identification using specific marker combinations from single-cell transcriptomic data. Nature Communications, 2022, 13, 1246. | 5.8 | 163 |
| 4 | Mono- and combinational drug therapies for global viral pandemic preparedness. IScience, 2022, 25, 104112. | 1.9 | 19 |
| 5 | SynergyFinder 3.0: an interactive analysis and consensus interpretation of multi-drug synergies across multiple samples. Nucleic Acids Research, 2022, 50, W739-W743. | 6.5 | 139 |
| 6 | DrugVirus.info 2.0: an integrative data portal for broad-spectrum antivirals (BSA) and BSA-containing drug combinations (BCCs). Nucleic Acids Research, 2022, 50, W272-W275. | 6.5 | 15 |
| 7 | Computational Pipeline for Rational Drug Combination Screening in Patient-Derived Cells. Methods in Molecular Biology, 2022, 2449, 327-348. | 0.4 | 4 |
| 8 | RUNX1 mutations in blast-phase chronic myeloid leukemia associate with distinct phenotypes, transcriptional profiles, and drug responses. Leukemia, 2021, 35, 1087-1099. | 3.3 | 32 |
| 9 | Safe-in-Man Broad Spectrum Antiviral Agents. Advances in Experimental Medicine and Biology, 2021, 1322, 313-337. | 0.8 | 1 |
| 10 | Neural signaling modulates metabolism of gastric cancer. IScience, 2021, 24, 102091. | 1.9 | 14 |
| 11 | Patient-tailored design for selective co-inhibition of leukemic cell subpopulations. Science Advances, 2021, 7, . | 4.7 | 28 |
| 12 | Machine Learning of Bone Marrow Histopathology Identifies Genetic and Clinical Determinants in Patients with MDS. Blood Cancer Discovery, 2021, 2, 238-249. | 2.6 | 25 |
| 13 | Computational Drug Repositioning and Experimental Validation of Ivermectin in Treatment of Gastric Cancer. Frontiers in Pharmacology, 2021, 12, 625991. | 1.6 | 7 |
| 14 | Inhibition of Arenaviruses by Combinations of Orally Available Approved Drugs. Antimicrobial Agents and Chemotherapy, 2021, 65, . | 1.4 | 27 |
| 15 | Screening of FDA-Approved Drugs Using a MERS-CoV Clinical Isolate from South Korea Identifies Potential Therapeutic Options for COVID-19. Viruses, 2021, 13, 651. | 1.5 | 50 |
| 16 | Development of HDAC Inhibitors Exhibiting Therapeutic Potential in T-Cell Prolymphocytic Leukemia. Journal of Medicinal Chemistry, 2021, 64, 8486-8509. | 2.9 | 28 |
| 17 | High-throughput compound screening identifies navitoclax combined with irradiation as a candidate therapy for HPV-negative head and neck squamous cell carcinoma. Scientific Reports, 2021, 11, 14755. | 1.6 | 7 |
| 18 | Active Components of Commonly Prescribed Medicines Affect Influenza A Virus–Host Cell Interaction: A Pilot Study. Viruses, 2021, 13, 1537. | 1.5 | 3 |

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|----|--|-----|-----------|
| 19 | Nafamostat–Interferon-α Combination Suppresses SARS-CoV-2 Infection In Vitro and In Vivo by Cooperatively Targeting Host TMPRSS2. Viruses, 2021, 13, 1768. | 1.5 | 15 |
| 20 | Drug Combinations as a First Line of Defense against Coronaviruses and Other Emerging Viruses. MBio, 2021, 12, e0334721. | 1.8 | 45 |
| 21 | Synergistic Interferon-Alpha-Based Combinations for Treatment of SARS-CoV-2 and Other Viral Infections. Viruses, 2021, 13, 2489. | 1.5 | 20 |
| 22 | Identification and Tracking of Antiviral Drug Combinations. Viruses, 2020, 12, 1178. | 1.5 | 48 |
| 23 | Potential Antiviral Options against SARS-CoV-2 Infection. Viruses, 2020, 12, 642. | 1.5 | 92 |
| 24 | Chemical, Physical and Biological Triggers of Evolutionary Conserved Bcl-xL-Mediated Apoptosis. Cancers, 2020, 12, 1694. | 1.7 | 13 |
| 25 | SynToxProfiler: An interactive analysis of drug combination synergy, toxicity and efficacy. PLoS Computational Biology, 2020, 16, e1007604. | 1.5 | 43 |
| 26 | Discovery and development of safe-in-man broad-spectrum antiviral agents. International Journal of Infectious Diseases, 2020, 93, 268-276. | 1.5 | 169 |
| 27 | Breeze: an integrated quality control and data analysis application for high-throughput drug screening. Bioinformatics, 2020, 36, 3602-3604. | 1.8 | 68 |
| 28 | SynergyFinder 2.0: visual analytics of multi-drug combination synergies. Nucleic Acids Research, 2020, 48, W488-W493. | 6.5 | 545 |
| 29 | Integrated drug profiling and CRISPR screening identify essential pathways for CAR T-cell cytotoxicity. Blood, 2020, 135, 597-609. | 0.6 | 134 |
| 30 | SynToxProfiler: An interactive analysis of drug combination synergy, toxicity and efficacy., 2020, 16, e1007604. | | 0 |
| 31 | SynToxProfiler: An interactive analysis of drug combination synergy, toxicity and efficacy. , 2020, 16, e1007604. | | 0 |
| 32 | SynToxProfiler: An interactive analysis of drug combination synergy, toxicity and efficacy., 2020, 16, e1007604. | | 0 |
| 33 | SynToxProfiler: An interactive analysis of drug combination synergy, toxicity and efficacy. , 2020, 16, e1007604. | | 0 |
| 34 | Common Nodes of Virus–Host Interaction Revealed Through an Integrated Network Analysis. Frontiers in Immunology, 2019, 10, 2186. | 2.2 | 67 |
| 35 | Novel Antiviral Activities of Obatoclax, Emetine, Niclosamide, Brequinar, and Homoharringtonine. Viruses, 2019, 11, 964. | 1.5 | 68 |
| 36 | Genome-wide off-targets of drugs: risks and opportunities. Cell Biology and Toxicology, 2019, 35, 485-487. | 2.4 | 16 |

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|----|---|-----|-----------|
| 37 | Low Temperature and Low UV Indexes Correlated with Peaks of Influenza Virus Activity in Northern Europe during 2010–2018. Viruses, 2019, 11, 207. | 1.5 | 81 |
| 38 | Expanding the activity spectrum of antiviral agents. Drug Discovery Today, 2019, 24, 1224-1228. | 3.2 | 31 |
| 39 | Prediction of drug combination effects with a minimal set of experiments. Nature Machine Intelligence, 2019, 1, 568-577. | 8.3 | 99 |
| 40 | Novel activities of safe-in-human broad-spectrum antiviral agents. Antiviral Research, 2018, 154, 174-182. | 1.9 | 64 |
| 41 | Interactive visual analysis of drug–target interaction networks using Drug Target Profiler, with applications to precision medicine and drug repurposing. Briefings in Bioinformatics, 2018, , . | 3.2 | 25 |
| 42 | A Systems Approach to Study Immuno- and Neuro-Modulatory Properties of Antiviral Agents. Viruses, 2018, 10, 423. | 1.5 | 10 |
| 43 | Single Passage of Human Metapneumovirus in LLC-MK2 Cells Does Not Affect Viral Protein-Coding Capacity. Genome Announcements, 2018, 6, . | 0.8 | 0 |
| 44 | SynergyFinder: a web application for analyzing drug combination dose–response matrix data. Bioinformatics, 2017, 33, 2413-2415. | 1.8 | 403 |
| 45 | Obatoclax, saliphenylhalamide and gemcitabine inhibit Zika virus infection inÂvitro and differentially affect cellular signaling, transcription and metabolism. Antiviral Research, 2017, 139, 117-128. | 1.9 | 88 |
| 46 | Antiviral Properties of Chemical Inhibitors of Cellular Anti-Apoptotic Bcl-2 Proteins. Viruses, 2017, 9, 271. | 1.5 | 39 |