Igor A Sidorov

List of Publications by Year in descending order

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49 papers

3,530 citations

279798
23
h-index

214800 47 g-index

57 all docs

57 docs citations

57 times ranked

6473 citing authors

| # | Article | IF | Citations |
|----|--|------|-----------|
| 1 | SARS-coronavirus-2 replication in Vero E6 cells: replication kinetics, rapid adaptation and cytopathology. Journal of General Virology, 2020, 101, 925-940. | 2.9 | 465 |
| 2 | Potent cross-reactive neutralization of SARS coronavirus isolates by human monoclonal antibodies. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 12123-12128. | 7.1 | 276 |
| 3 | Identification of Dynamically Distinct Subpopulations of T Lymphocytes That Are Differentially Affected by HIV. Journal of Experimental Medicine, 2001, 194, 1731-1741. | 8.5 | 203 |
| 4 | Broadly cross-reactive HIV-1-neutralizing human monoclonal Fab selected for binding to gp120-CD4-CCR5 complexes. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 6913-6918. | 7.1 | 203 |
| 5 | Discovery of an essential nucleotidylating activity associated with a newly delineated conserved domain in the RNA polymerase-containing protein of all nidoviruses. Nucleic Acids Research, 2015, 43, 8416-8434. | 14.5 | 197 |
| 6 | Dissection of Human Immunodeficiency Virus Type 1 Entry with Neutralizing Antibodies to gp41 Fusion Intermediates. Journal of Virology, 2002, 76, 6780-6790. | 3.4 | 115 |
| 7 | Induction of prolonged survival of CD4+ T lymphocytes by intermittent IL-2 therapy in HIV-infected patients. Journal of Clinical Investigation, 2005, 115, 2139-2148. | 8.2 | 115 |
| 8 | Leukocyte telomere dynamics and human hematopoietic stem cell kinetics during somatic growth. Experimental Hematology, 2009, 37, 514-524. | 0.4 | 114 |
| 9 | Identification and Characterization of a New Cross-Reactive Human Immunodeficiency Virus Type 1-Neutralizing Human Monoclonal Antibody. Journal of Virology, 2004, 78, 9233-9242. | 3.4 | 80 |
| 10 | Nail ve T-Cell Dynamics in Human Immunodeficiency Virus Type 1 Infection: Effects of Highly Active Antiretroviral Therapy Provide Insights into the Mechanisms of Nail ve T-Cell Depletion. Journal of Virology, 2006, 80, 2665-2674. | 3.4 | 66 |
| 11 | Increased efficacy of HIV-1 neutralization by antibodies at low CCR5 surface concentration. Biochemical and Biophysical Research Communications, 2006, 348, 1107-1115. | 2.1 | 62 |
| 12 | Cross-reactive HIV-1 neutralizing monoclonal antibodies selected by screening of an immune human phage library against an envelope glycoprotein (gp140) isolated from a patient (R2) with broadly HIV-1 neutralizing antibodies. Virology, 2007, 363, 79-90. | 2.4 | 57 |
| 13 | Retrospective Validation of a Metagenomic Sequencing Protocol for Combined Detection of RNA and DNA Viruses Using Respiratory Samples from Pediatric Patients. Journal of Molecular Diagnostics, 2020, 22, 196-207. | 2.8 | 56 |
| 14 | The respiratory virome and exacerbations in patients with chronic obstructive pulmonary disease. PLoS ONE, 2019, 14, e0223952. | 2.5 | 51 |
| 15 | Energy cost of infection burden: An approach to understanding the dynamics of host–pathogen interactions. Journal of Theoretical Biology, 2006, 241, 1-13. | 1.7 | 43 |
| 16 | STROBE-metagenomics: a STROBE extension statement to guide the reporting of metagenomics studies. Lancet Infectious Diseases, The, 2020, 20, e251-e260. | 9.1 | 40 |
| 17 | Practical application of bioinformatics by the multidisciplinary VIZIER consortium. Antiviral Research, 2010, 87, 95-110. | 4.1 | 39 |
| 18 | Recommendations for the introduction of metagenomic next-generation sequencing in clinical virology, part II: bioinformatic analysis and reporting. Journal of Clinical Virology, 2021, 138, 104812. | 3.1 | 39 |

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|----|--|------|-----------|
| 19 | Improved diagnosis of viral encephalitis in adult and pediatric hematological patients using viral metagenomics. Journal of Clinical Virology, 2020, 130, 104566. | 3.1 | 35 |
| 20 | Haem is crucial for medium-dependent metronidazole resistance in clinical isolates of <i>Clostridioides difficile</i> Journal of Antimicrobial Chemotherapy, 2021, 76, 1731-1740. | 3.0 | 34 |
| 21 | Role of Ets/Id proteins for telomerase regulation in human cancer cells. Experimental and Molecular Pathology, 2003, 75, 238-247. | 2.1 | 32 |
| 22 | Selection of a novel gp41-specific HIV-1 neutralizing human antibody by competitive antigen panning. Journal of Immunological Methods, 2006, 317, 21-30. | 1.4 | 32 |
| 23 | Coronavirus discovery by metagenomic sequencing: a tool for pandemic preparedness. Journal of Clinical Virology, 2020, 131, 104594. | 3.1 | 31 |
| 24 | Correlation between binding rate constants and individual information of E. coli Fis binding sites. Nucleic Acids Research, 2007, 35, 5275-5283. | 14.5 | 30 |
| 25 | Torque teno virus loads after kidney transplantation predict allograft rejection but not viral infection. Journal of Clinical Virology, 2021, 140, 104871. | 3.1 | 28 |
| 26 | Benchmark of thirteen bioinformatic pipelines for metagenomic virus diagnostics using datasets from clinical samples. Journal of Clinical Virology, 2021, 141, 104908. | 3.1 | 28 |
| 27 | Identification of a novel CD4i human monoclonal antibody Fab that neutralizes HIV-1 primary isolates from different clades. Antiviral Research, 2004, 61, 161-164. | 4.1 | 26 |
| 28 | Concordant Modulation of Neutralization Resistance and High Infectivity of the Primary Human Immunodeficiency Virus Type 1 MN Strain and Definition of a Potential gp41 Binding Site in gp120. Journal of Virology, 2003, 77, 560-570. | 3.4 | 23 |
| 29 | Quantitative cortical synapse proteomics of a transgenic migraine mouse model with mutated Ca _V 2.1 calcium channels. Proteomics, 2010, 10, 2531-2535. | 2.2 | 21 |
| 30 | Conserved Structures Exposed in HIV-1 Envelope Glycoproteins Stabilized by Flexible Linkers as Potent Entry Inhibitors and Potential Immunogensâ€. Biochemistry, 2002, 41, 7176-7182. | 2.5 | 19 |
| 31 | A kinetic model of telomere shortening in infants and adults. Journal of Theoretical Biology, 2004, 226, 169-175. | 1.7 | 19 |
| 32 | Multiple Interactions across the Surface of the gp120 Core Structure Determine the Global Neutralization Resistance Phenotype of Human Immunodeficiency Virus Type 1. Journal of Virology, 2003, 77, 8061-8071. | 3.4 | 18 |
| 33 | Mathematical modeling of citric acid production by repeated batch culture. Enzyme and Microbial Technology, 2000, 26, 826-833. | 3.2 | 17 |
| 34 | Cancer treatment by telomerase inhibitors: predictions by a kinetic model. Mathematical Biosciences, 2003, 181, 209-221. | 1.9 | 15 |
| 35 | Retinoic acid-induced downmodulation of telomerase activity in human cancer cells. Experimental and Molecular Pathology, 2005, 79, 108-117. | 2.1 | 14 |
| 36 | Purification and properties of polyphosphatase from Saccharomyces cerevisiae cytosol., 1998, 14, 383-390. | | 13 |

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|----|--|-----|-----------|
| 37 | Viral metagenomic sequencing in a cohort of international travellers returning with febrile illness. Journal of Clinical Virology, 2021, 143, 104940. | 3.1 | 10 |
| 38 | Cancer Cell Dynamics in Presence of Telomerase Inhibitors: Analysis of In Vitro Data. Journal of Theoretical Biology, 2002, 219, 225-233. | 1.7 | 9 |
| 39 | Rapid Elimination of Broadly Neutralizing Antibodies Correlates with Treatment Failure in the Acute Phase of Simian-Human Immunodeficiency Virus Infection. Journal of Virology, 2019, 93, . | 3.4 | 8 |
| 40 | Renal transplant patients show variations in their self-reactive repertoires: a serial study. International Immunology, 2001, 13, 747-755. | 4.0 | 7 |
| 41 | SNAD: sequence name annotation-based designer. BMC Bioinformatics, 2009, 10, 251. | 2.6 | 6 |
| 42 | Inhibition analysis of hydroxyquinol-cleaving dioxygenases from the chlorophenol-degradingAzotobacter sp. GP1 andStreptomyces rochei 303. Journal of Basic Microbiology, 1999, 39, 61-73. | 3.3 | 5 |
| 43 | Limiting Dilution Analysis of Interleukin-2 Producing Helper T-cell Frequencies as a Tool in Allogeneic Hematopoietic Cell Transplantation. Transplantation, 2005, 80, 573-581. | 1.0 | 5 |
| 44 | Genome-wide diversity of Zika virus: Exploring spatio-temporal dynamics to guide a new nomenclature proposal. Virus Evolution, 2022, 8, veac029. | 4.9 | 5 |
| 45 | Performance of Five Metagenomic Classifiers for Virus Pathogen Detection Using Respiratory Samples from a Clinical Cohort. Pathogens, 2022, 11, 340. | 2.8 | 4 |
| 46 | Longitudinal Monitoring of DNA Viral Loads in Transplant Patients Using Quantitative Metagenomic Next-Generation Sequencing. Pathogens, 2022, 11, 236. | 2.8 | 3 |
| 47 | Rhinovirus Detection in the Nasopharynx of Children Undergoing Cardiac Surgery Is Not Associated With Longer PICU Length of Stay: Results of the Impact of Rhinovirus Infection After Cardiac Surgery in Kids (RISK) Study. Pediatric Critical Care Medicine, 2021, 22, e79-e90. | 0.5 | 2 |
| 48 | Translating genomic exploration of the family Polyomaviridae into confident human polyomavirus detection. IScience, 2022, 25, 103613. | 4.1 | 2 |
| 49 | A probability model predicting initiation efficiency of retroviral vectors with two primer-binding sites. Journal of Theoretical Biology, 2006, 242, 347-355. | 1.7 | O |