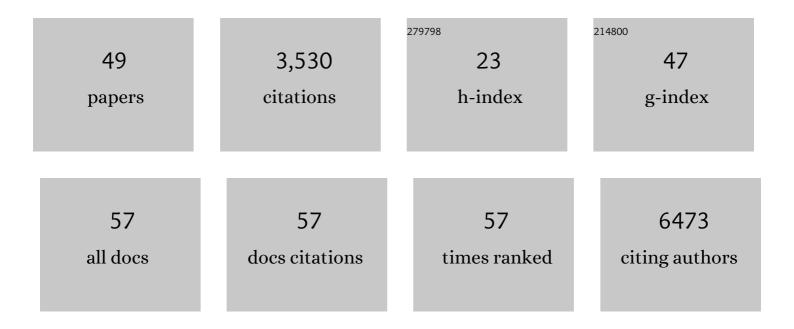
Igor A Sidorov

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Translating genomic exploration of the family Polyomaviridae into confident human polyomavirus detection. IScience, 2022, 25, 103613.	4.1	2
2	Genome-wide diversity of Zika virus: Exploring spatio-temporal dynamics to guide a new nomenclature proposal. Virus Evolution, 2022, 8, veac029.	4.9	5
3	Longitudinal Monitoring of DNA Viral Loads in Transplant Patients Using Quantitative Metagenomic Next-Generation Sequencing. Pathogens, 2022, 11, 236.	2.8	3
4	Performance of Five Metagenomic Classifiers for Virus Pathogen Detection Using Respiratory Samples from a Clinical Cohort. Pathogens, 2022, 11, 340.	2.8	4
5	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
6	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
7	Torque teno virus loads after kidney transplantation predict allograft rejection but not viral infection. Journal of Clinical Virology, 2021, 140, 104871.	3.1	28
8	Benchmark of thirteen bioinformatic pipelines for metagenomic virus diagnostics using datasets from clinical samples. Journal of Clinical Virology, 2021, 141, 104908.	3.1	28
9	Viral metagenomic sequencing in a cohort of international travellers returning with febrile illness. Journal of Clinical Virology, 2021, 143, 104940.	3.1	10
10	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
11	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
12	Coronavirus discovery by metagenomic sequencing: a tool for pandemic preparedness. Journal of Clinical Virology, 2020, 131, 104594.	3.1	31
13	Improved diagnosis of viral encephalitis in adult and pediatric hematological patients using viral metagenomics. Journal of Clinical Virology, 2020, 130, 104566.	3.1	35
14	STROBE-metagenomics: a STROBE extension statement to guide the reporting of metagenomics studies. Lancet Infectious Diseases, The, 2020, 20, e251-e260.	9.1	40
15	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
16	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
17	The respiratory virome and exacerbations in patients with chronic obstructive pulmonary disease. PLoS ONE, 2019, 14, e0223952.	2.5	51
18	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

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19	Practical application of bioinformatics by the multidisciplinary VIZIER consortium. Antiviral Research, 2010, 87, 95-110.	4.1	39
20	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
21	SNAD: sequence name annotation-based designer. BMC Bioinformatics, 2009, 10, 251.	2.6	6
22	Leukocyte telomere dynamics and human hematopoietic stem cell kinetics during somatic growth. Experimental Hematology, 2009, 37, 514-524.	0.4	114
23	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
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	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
26	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
27	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
28	A probability model predicting initiation efficiency of retroviral vectors with two primer-binding sites. Journal of Theoretical Biology, 2006, 242, 347-355.	1.7	0
29	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
30	Nail̀^ve T-Cell Dynamics in Human Immunodeficiency Virus Type 1 Infection: Effects of Highly Active Antiretroviral Therapy Provide Insights into the Mechanisms of Nail̂ve T-Cell Depletion. Journal of Virology, 2006, 80, 2665-2674.	3.4	66
31	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
32	Retinoic acid-induced downmodulation of telomerase activity in human cancer cells. Experimental and Molecular Pathology, 2005, 79, 108-117.	2.1	14
33	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
34	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
35	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
36	A kinetic model of telomere shortening in infants and adults. Journal of Theoretical Biology, 2004, 226, 169-175.	1.7	19

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37	Role of Ets/Id proteins for telomerase regulation in human cancer cells. Experimental and Molecular Pathology, 2003, 75, 238-247.	2.1	32
38	Cancer treatment by telomerase inhibitors: predictions by a kinetic model. Mathematical Biosciences, 2003, 181, 209-221.	1.9	15
39	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
40	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
41	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
42	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
43	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
44	Cancer Cell Dynamics in Presence of Telomerase Inhibitors: Analysis of In Vitro Data. Journal of Theoretical Biology, 2002, 219, 225-233.	1.7	9
45	Renal transplant patients show variations in their self-reactive repertoires: a serial study. International Immunology, 2001, 13, 747-755.	4.0	7
46	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
47	Mathematical modeling of citric acid production by repeated batch culture. Enzyme and Microbial Technology, 2000, 26, 826-833.	3.2	17
48	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
49	Purification and properties of polyphosphatase fromSaccharomyces cerevisiae cytosol. , 1998, 14, 383-390.		13