

Ofer Isakov

List of Publications by Year in descending order

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Version: 2024-02-01

44
papers

1,517
citations

430754

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330025

37
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docs citations

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times ranked

3186
citing authors

#	ARTICLE	IF	CITATIONS
1	Rapid 8â€Zone Lung Ultrasound Protocol is Comparable to a Full 12â€Zone Protocol for Outcome Prediction in Hospitalized <sc>COVID</sc>â€19 Patients. Journal of Ultrasound in Medicine, 2022, 41, 1677-1687.	0.8	7
2	Efficacy of a second external cephalic version (ECV) after a successful first external cephalic version with subsequent spontaneous reinversion to breech presentation: a retrospective cohort study. Archives of Gynecology and Obstetrics, 2021, 303, 911-916.	0.8	4
3	Can uric acid blood levels in renal transplant recipients predict allograft outcome?. Renal Failure, 2021, 43, 1240-1249.	0.8	3
4	Assessing the involvement of the placental microbiome and virome in preeclampsia using non coding RNA sequencing. Journal of Perinatal Medicine, 2021, 49, 1071-1083.	0.6	0
5	Targeted therapies for immune thrombocytopenic purpura: a meta-analysis of randomized controlled trials. Annals of Hematology, 2021, 100, 2879-2887.	0.8	3
6	Treatment with Ixekizumab Following Secukinumab Failure in Patients with Psoriatic Arthritis: Real-Life Experience from a Resistant Population. Biologics: Targets and Therapy, 2021, Volume 15, 463-470.	3.0	5
7	Resection Versus Observation of Small Asymptomatic Nonfunctioning Pancreatic Neuroendocrine Tumors. Journal of Gastrointestinal Surgery, 2020, 24, 1366-1374.	0.9	26
8	Posttransplantation Hypomagnesemia as a Predictor of Better Graft Function after Transplantation. Kidney and Blood Pressure Research, 2020, 45, 982-995.	0.9	3
9	Spectrum of genes for inherited hearing loss in the Israeli Jewish population, including the novel human deafness gene <sc><i>ATOH1</i></sc>. Clinical Genetics, 2020, 98, 353-364.	1.0	15
10	Early persistent hyperparathyroidism postâ€renal transplantation as a predictor of worse graft function and mortality after transplantation. Clinical Transplantation, 2020, 34, e14085.	0.8	11
11	The effects of a postmiscarriage menstrual period prior to reconceiving. American Journal of Obstetrics and Gynecology, 2020, 223, 444.e1-444.e5.	0.7	1
12	Effect of Ticagrelor on Reducing the Risk of Gram-Positive Infections in Patients With Acute Coronary Syndrome. American Journal of Cardiology, 2020, 130, 56-63.	0.7	12
13	The dark age of single organ screening is over: CD24 is a novel universal simple blood test for early detection of cancer.. Journal of Clinical Oncology, 2020, 38, e15591-e15591.	0.8	0
14	Effect of liver fibrosis on survival in patients with intrahepatic cholangiocarcinoma: a SEER population-based study. Oncotarget, 2020, 11, 4438-4447.	0.8	4
15	HIV-1 infection increases microRNAs that inhibit Dicer1, HRB and HIV-EP2, thereby reducing viral replication. PLoS ONE, 2019, 14, e0211111.	1.1	22
16	Variant <i>PADI3</i> in Central Centrifugal Cicatricial Alopecia. New England Journal of Medicine, 2019, 380, 833-841.	13.9	102
17	Prediction of Success in External Cephalic Version for Breech Presentation at Term. Obstetrics and Gynecology, 2019, 133, 857-866.	1.2	30
18	Genotypeâ€Serotype Interactions Shed Light on Genetic Components of Inflammatory Bowel Diseases. Inflammatory Bowel Diseases, 2019, 25, 336-344.	0.9	2

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19	Data from an integrated cancer prevention center screening for multiple cancer types.. Journal of Clinical Oncology, 2019, 37, e13069-e13069.	0.8	2
20	Exhaustive non-synonymous variants functionality prediction enables high resolution characterization of the neurofibromin architecture. EBioMedicine, 2018, 36, 508-516.	2.7	1
21	Seasonal Genetic Drift of Human Influenza A Virus Quasispecies Revealed by Deep Sequencing. Frontiers in Microbiology, 2018, 9, 2596.	1.5	27
22	Neoadjuvant FOLFIRINOX for locally advanced and borderline resectable pancreatic cancer: An intention to treat analysis. European Journal of Surgical Oncology, 2018, 44, 1619-1623.	0.5	82
23	X-linked elliptocytosis with impaired growth is related to mutated AMMECR1. Gene, 2017, 606, 47-52.	1.0	6
24	Machine Learning-Based Gene Prioritization Identifies Novel Candidate Risk Genes for Inflammatory Bowel Disease. Inflammatory Bowel Diseases, 2017, 23, 1516-1523.	0.9	49
25	Calpain 12 Function Revealed through the Study of an Atypical Case of Autosomal Recessive Congenital Ichthyosis. Journal of Investigative Dermatology, 2017, 137, 385-393.	0.3	19
26	Influence of population diversity on neurovirulence potential of plaque purified L-Zagreb variants. Vaccine, 2016, 34, 2383-2389.	1.7	2
27	Whole-Genome Sequencing Analysis from the Chikungunya Virus Caribbean Outbreak Reveals Novel Evolutionary Genomic Elements. PLoS Neglected Tropical Diseases, 2016, 10, e0004402.	1.3	96
28	MicroRNA regulation of p21 and TASK1 cellular restriction-factors enhances HIV-1 infection. Journal of Cell Science, 2015, 128, 1607-16.	1.2	51
29	Actionable clinical decisions based on comprehensive genomic evaluation in asymptomatic adults. Molecular Genetics & Genomic Medicine, 2015, 3, 433-439.	0.6	7
30	Rare genetic variants in Tunisian Jewish patients suffering from age-related macular degeneration. Journal of Medical Genetics, 2015, 52, 484-492.	1.5	19
31	Crowdfunding Effort Identifies the Causative Mutation in a Patient with Nystagmus, Microcephaly, Dystonia and Hypomyelination. Journal of Genetics and Genomics, 2015, 42, 79-81.	1.7	14
32	Group Selection and Contribution of Minority Variants during Virus Adaptation Determines Virus Fitness and Phenotype. PLoS Pathogens, 2015, 11, e1004838.	2.1	106
33	Mutations in Coronavirus Nonstructural Protein 10 Decrease Virus Replication Fidelity. Journal of Virology, 2015, 89, 6418-6426.	1.5	56
34	Deep sequencing analysis of viral infection and evolution allows rapid and detailed characterization of viral mutant spectrum. Bioinformatics, 2015, 31, 2141-2150.	1.8	62
35	Non-syndromic retinitis pigmentosa due to mutations in the mucopolysaccharidosis type IIIC gene, heparan-alpha-glucosaminide N-acetyltransferase (HGSNAT). Human Molecular Genetics, 2015, 24, 3742-51.	1.4	47
36	Phylogenetic and Genome-Wide Deep-Sequencing Analyses of Canine Parvovirus Reveal Co-Infection with Field Variants and Emergence of a Recent Recombinant Strain. PLoS ONE, 2014, 9, e111779.	1.1	73

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37	Sequencing your genome: your future is here, but are you sure you want to know it?. Genetical Research, 2014, 96, e006.	0.3	2
38	MicroRNA Editing Facilitates Immune Elimination of HCMV Infected Cells. PLoS Pathogens, 2014, 10, e1003963.	2.1	40
39	Emergence and Transmission of Arbovirus Evolutionary Intermediates with Epidemic Potential. Cell Host and Microbe, 2014, 15, 706-716.	5.1	107
40	Exome Sequencing Analysis: A Guide to Disease Variant Detection. Methods in Molecular Biology, 2013, 1038, 137-158.	0.4	18
41	Desmoglein 1 deficiency results in severe dermatitis, multiple allergies and metabolic wasting. Nature Genetics, 2013, 45, 1244-1248.	9.4	289
42	Missense mutation in the MEN1 gene discovered through whole exome sequencing co-segregates with familial hyperparathyroidism. Genetical Research, 2013, 95, 114-120.	0.3	11
43	Novel insight into the non-coding repertoire through deep sequencing analysis. Nucleic Acids Research, 2012, 40, e86-e86.	6.5	17
44	Pathogen detection using short-RNA deep sequencing subtraction and assembly. Bioinformatics, 2011, 27, 2027-2030.	1.8	63