

Kevin Galinsky

List of Publications by Year in descending order

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14
papers

1,204
citations

932766

10
h-index

1125271

13
g-index

15
all docs

15
docs citations

15
times ranked

2814
citing authors

#	ARTICLE	IF	CITATIONS
1	Sapanisertib plus Fulvestrant in Postmenopausal Women with Estrogen Receptorâ€“Positive/HER2-Negative Advanced Breast Cancer after Progression on Aromatase Inhibitor. <i>Clinical Cancer Research</i> , 2022, 28, 1107-1116.	3.2	7
2	Mucosal Eosinophil Abundance in Non-Inflamed Colonic Tissue Is Associated with Response to Vedolizumab Induction Therapy in Inflammatory Bowel Disease. <i>Journal of Clinical Medicine</i> , 2022, 11, 4141.	1.0	1
3	Evaluating Responses to Gluten Challenge: A Randomized, Double-Blind, 2-Dose Gluten Challenge Trial. <i>Gastroenterology</i> , 2021, 160, 720-733.e8.	0.6	53
4	Sapanisertib Plus Exemestane or Fulvestrant in Women with Hormone Receptorâ€“Positive/HER2-Negative Advanced or Metastatic Breast Cancer. <i>Clinical Cancer Research</i> , 2021, 27, 3329-3338.	3.2	8
5	Fast Principal-Component Analysis Reveals Convergent Evolution of ADH1B in Europe and East Asia. <i>American Journal of Human Genetics</i> , 2016, 98, 456-472.	2.6	335
6	Diversity-Oriented Synthesis Probe Targets <i>Plasmodium falciparum</i> Cytochrome b Ubiquinone Reduction Site and Synergizes With Oxidation Site Inhibitors. <i>Journal of Infectious Diseases</i> , 2015, 211, 1097-1103.	1.9	29
7	COIL: a methodology for evaluating malarial complexity of infection using likelihood from single nucleotide polymorphism data. <i>Malaria Journal</i> , 2015, 14, 4.	0.8	71
8	The cytoplasmic prolyl-tRNA synthetase of the malaria parasite is a dual-stage target of febrifugine and its analogs. <i>Science Translational Medicine</i> , 2015, 7, 288ra77.	5.8	82
9	Development of a Single Nucleotide Polymorphism Barcode to Genotype <i>Plasmodium vivax</i> Infections. <i>PLoS Neglected Tropical Diseases</i> , 2015, 9, e0003539.	1.3	90
10	An Epigenetic Antimalarial Resistance Mechanism Involving Parasite Genes Linked to Nutrient Uptake. <i>Journal of Biological Chemistry</i> , 2013, 288, 19429-19440.	1.6	71
11	Sequence-based association and selection scans identify drug resistance loci in the <i>Plasmodium falciparum</i> malaria parasite. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 13052-13057.	3.3	99
12	The malaria parasite <i>Plasmodium vivax</i> exhibits greater genetic diversity than <i>Plasmodium falciparum</i> . <i>Nature Genetics</i> , 2012, 44, 1046-1050.	9.4	256
13	Hybrid selection for sequencing pathogen genomes from clinical samples. <i>Genome Biology</i> , 2011, 12, R73.	13.9	97
14	Comparisons of predicted genetic modules: identification of co-expressed genes through module gene flow. <i>Genome Informatics</i> , 2004, 15, 221-8.	0.4	4