Upinder Singh

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7482402/publications.pdf

Version: 2024-02-01

91 papers

4,973 citations

34 h-index 98622 67 g-index

188 all docs 188
docs citations

188 times ranked 5175 citing authors

#	Article	IF	Citations
1	The genome of the protist parasite Entamoeba histolytica. Nature, 2005, 433, 865-868.	13.7	783
2	New-onset IgG autoantibodies in hospitalized patients with COVID-19. Nature Communications, 2021, 12, 5417.	5.8	286
3	Distinct Distal Gut Microbiome Diversity and Composition in Healthy Children from Bangladesh and the United States. PLoS ONE, 2013, 8, e53838.	1.1	278
4	Proinflammatory IgG Fc structures in patients with severe COVID-19. Nature Immunology, 2021, 22, 67-73.	7.0	239
5	Toxoplasma gondii Asexual Development: Identification of Developmentally Regulated Genes and Distinct Patterns of Gene Expression. Eukaryotic Cell, 2002, 1, 329-340.	3.4	196
6	Gastrointestinal symptoms and fecal shedding of SARS-CoV-2 RNA suggest prolonged gastrointestinal infection. Med, 2022, 3, 371-387.e9.	2.2	165
7	Impact of intestinal colonization and invasion on the Entamoeba histolytica transcriptome. Molecular and Biochemical Parasitology, 2006, 147, 163-176.	0.5	153
8	Identification of developmentally regulated genes in Entamoeba histolytica: insights into mechanisms of stage conversion in a protozoan parasite. Cellular Microbiology, 2007, 9, 1426-1444.	1.1	128
9	Genetic analysis of tachyzoite to bradyzoite differentiation mutants in Toxoplasma gondii reveals a hierarchy of gene induction. Molecular Microbiology, 2002, 44, 721-733.	1.2	127
10	Patients With Uncomplicated Coronavirus Disease 2019 (COVID-19) Have Long-Term Persistent Symptoms and Functional Impairment Similar to Patients with Severe COVID-19: A Cautionary Tale During a Global Pandemic. Clinical Infectious Diseases, 2021, 73, e826-e829.	2.9	127
11	Identification of Differentially Expressed Genes in Virulent and Nonvirulent Entamoeba Species: Potential Implications for Amebic Pathogenesis. Infection and Immunity, 2006, 74, 340-351.	1.0	117
12	Interferon- \hat{I}^3 Release Assay for Accurate Detection of Severe Acute Respiratory Syndrome Coronavirus 2 T-Cell Response. Clinical Infectious Diseases, 2021, 73, e3130-e3132.	2.9	114
13	The genome and transcriptome of the enteric parasite Entamoeba invadens, a model for encystation. Genome Biology, 2013, 14, R77.	13.9	111
14	Peginterferon Lambda-1a for treatment of outpatients with uncomplicated COVID-19: a randomized placebo-controlled trial. Nature Communications, 2021, 12, 1967.	5.8	107
15	<i>Entamoeba histolytica</i> modulates a complex repertoire of novel genes in response to oxidative and nitrosative stresses: implications for amebic pathogenesis. Cellular Microbiology, 2009, 11, 51-69.	1.1	102
16	Antibodies elicited by SARS-CoV-2 infection or mRNA vaccines have reduced neutralizing activity against Beta and Omicron pseudoviruses. Science Translational Medicine, 2022, 14, eabn7842.	5.8	92
17	An <i>Entamoeba histolytica</i> rhomboid protease with atypical specificity cleaves a surface lectin involved in phagocytosis and immune evasion. Genes and Development, 2008, 22, 1636-1646.	2.7	84
18	Early non-neutralizing, afucosylated antibody responses are associated with COVID-19 severity. Science Translational Medicine, 2022, 14, eabm7853.	5.8	71

#	Article	IF	CITATIONS
19	Identification and characterization of differentiation mutants in the protozoan parasite Toxoplasma gondii. Molecular Microbiology, 2002, 44, 735-747.	1.2	68
20	Small RNAs with 5′-Polyphosphate Termini Associate with a Piwi-Related Protein and Regulate Gene Expression in the Single-Celled Eukaryote Entamoeba histolytica. PLoS Pathogens, 2008, 4, e1000219.	2.1	65
21	Downregulation of an Entamoeba histolytica Rhomboid Protease Reveals Roles in Regulating Parasite Adhesion and Phagocytosis. Eukaryotic Cell, 2010, 9, 1283-1293.	3.4	65
22	Robust gene silencing mediated by antisense small RNAs in the pathogenic protist Entamoeba histolytica. Nucleic Acids Research, 2013, 41, 9424-9437.	6.5	63
23	Identification of an <i>Entamoeba histolytica</i> Serine-, Threonine-, and Isoleucine-Rich Protein with Roles in Adhesion and Cytotoxicity. Eukaryotic Cell, 2007, 6, 2139-2146.	3.4	55
24	A developmentally regulated Myb domain protein regulates expression of a subset of stage-specific genes in <i>Entamoeba histolytica</i> Cellular Microbiology, 2009, 11, 898-910.	1.1	54
25	New insights into Entamoeba histolytica pathogenesis. Current Opinion in Infectious Diseases, 2008, 21, 489-494.	1.3	53
26	Nucleus-localized Antisense Small RNAs with 5′-Polyphosphate Termini Regulate Long Term Transcriptional Gene Silencing in Entamoeba histolytica G3 Strain. Journal of Biological Chemistry, 2011, 286, 44467-44479.	1.6	51
27	Comparative Genomic Hybridizations of Entamoeba Strains Reveal Unique Genetic Fingerprints That Correlate with Virulence. Eukaryotic Cell, 2005, 4, 504-515.	3.4	50
28	Regulation of H2O2 Stress-responsive Genes through a Novel Transcription Factor in the Protozoan Pathogen Entamoeba histolytica. Journal of Biological Chemistry, 2013, 288, 4462-4474.	1.6	48
29	A Detoxifying Oxygen Reductase in the Anaerobic Protozoan Entamoeba histolytica. Eukaryotic Cell, 2012, 11, 1112-1118.	3.4	47
30	Oxidative stress resistance genes contribute to the pathogenic potential of the anaerobic protozoan parasite, Entamoeba histolytica. International Journal for Parasitology, 2012, 42, 1007-1015.	1.3	43
31	Trichostatin A effects on gene expression in the protozoan parasite Entamoeba histolytica. BMC Genomics, 2007, 8, 216.	1.2	41
32	Identification of putative transcriptional regulatory networks in Entamoeba histolytica using Bayesian inference. Nucleic Acids Research, 2007, 35, 2141-2152.	6.5	40
33	Short hairpin RNA-mediated knockdown of protein expression in Entamoeba histolytica. BMC Microbiology, 2009, 9, 38.	1.3	39
34	High-Throughput Screening of Entamoeba Identifies Compounds Which Target Both Life Cycle Stages and Which Are Effective Against Metronidazole Resistant Parasites. Frontiers in Cellular and Infection Microbiology, 2018, 8, 276.	1.8	39
35	Dimethylated H3K27 Is a Repressive Epigenetic Histone Mark in the Protist Entamoeba histolytica and Is Significantly Enriched in Genes Silenced via the RNAi Pathway. Journal of Biological Chemistry, 2015, 290, 21114-21130.	1.6	36
36	Growth of the protozoan parasite Entamoeba histolytica in 5-azacytidine has limited effects on parasite gene expression. BMC Genomics, 2007, 8, 7.	1.2	35

#	Article	IF	Citations
37	Standardized preservation, extraction and quantification techniques for detection of fecal SARS-CoV-2 RNA. Nature Communications, 2021, 12, 5753.	5.8	32
38	Transcriptional profiling of Entamoeba histolytica trophozoites. International Journal for Parasitology, 2005, 35, 533-542.	1.3	31
39	RNA interference in <i>Entamoeba histolytica</i> : implications for parasite biology and gene silencing. Future Microbiology, 2011, 6, 103-117.	1.0	31
40	Transient and stable transfection in the protozoan parasite Entamoeba invadens. Molecular and Biochemical Parasitology, 2012, 184, 59-62.	0.5	29
41	Characterization of Extracellular Vesicles from Entamoeba histolytica Identifies Roles in Intercellular Communication That Regulates Parasite Growth and Development. Infection and Immunity, 2020, 88, .	1.0	29
42	Coding and Noncoding Genomic Regions of Entamoeba histolytica Have Significantly Different Rates of Sequence Polymorphisms: Implications for Epidemiological Studies. Journal of Clinical Microbiology, 2005, 43, 4815-4819.	1.8	28
43	DNA microarrays in parasitology: strengths and limitations. Trends in Parasitology, 2003, 19, 470-476.	1.5	27
44	Small RNA pyrosequencing in the protozoan parasite Entamoeba histolytica reveals strain-specific small RNAs that target virulence genes. BMC Genomics, 2013, 14, 53.	1.2	27
45	Favipiravir for Treatment of Outpatients With Asymptomatic or Uncomplicated Coronavirus Disease 2019: A Double-Blind, Randomized, Placebo-Controlled, Phase 2 Trial. Clinical Infectious Diseases, 2022, 75, 1883-1892.	2.9	27
46	The Novel Core Promoter Element GAAC in the hgl5 Gene of Entamoeba histolytica Is Able to Direct a Transcription Start Site Independent of TATA or Initiator Regions. Journal of Biological Chemistry, 1998, 273, 21663-21668.	1.6	26
47	Functional Characterization of Spliceosomal Introns and Identification of U2, U4, and U5 snRNAs in the Deep-Branching Eukaryote Entamoeba histolytica. Eukaryotic Cell, 2007, 6, 940-948.	3.4	24
48	SARS-CoV-2 Subgenomic RNA Kinetics in Longitudinal Clinical Samples. Open Forum Infectious Diseases, 2021, 8, ofab310.	0.4	24
49	Context-dependent roles of the Entamoeba histolytica core promoter element GAAC in transcriptional activation and protein complex assembly. Molecular and Biochemical Parasitology, 2002, 120, 107-116.	0.5	23
50	Recent insights into Entamoeba development: Identification of transcriptional networks associated with stage conversion. International Journal for Parasitology, 2009, 39, 41-47.	1.3	23
51	Entamoeba histolytica rhomboid protease 1 has a role in migration and motility as validated by two independent genetic approaches. Experimental Parasitology, 2015, 154, 33-42.	0.5	22
52	Entamoeba histolytica: a snapshot of current research and methods for genetic analysis. Current Opinion in Microbiology, 2012, 15, 469-475.	2.3	21
53	Development of RNA Interference Trigger-Mediated Gene Silencing in Entamoeba invadens. Infection and Immunity, 2016, 84, 964-975.	1.0	21
54	An NAD+-dependent novel transcription factor controls stage conversion in Entamoeba. ELife, 2018, 7,	2.8	21

#	Article	IF	CITATIONS
55	Regulation of gene expression in the protozoan parasite Entamoeba invadens: identification of core promoter elements and promoters with stage-specific expression patterns. International Journal for Parasitology, 2014, 44, 837-845.	1.3	20
56	Identification of anisomycin, prodigiosin and obatoclax as compounds with broad-spectrum anti-parasitic activity. PLoS Neglected Tropical Diseases, 2020, 14, e0008150.	1.3	20
57	Identification of plicamycin, TG02, panobinostat, lestaurtinib, and GDC-0084 as promising compounds for the treatment of central nervous system infections caused by the free-living amebae Naegleria, Acanthamoeba and Balamuthia. International Journal for Parasitology: Drugs and Drug Resistance, 2019. 11. 80-94.	1.4	18
58	Nuclear Factor Y (NF-Y) Modulates Encystation in <i>Entamoeba</i> via Stage-Specific Expression of the NF-YB and NF-YC Subunits. MBio, 2019, 10, .	1.8	17
59	Technical advances in trigger-induced RNA interference gene silencing in the parasite Entamoeba histolytica. International Journal for Parasitology, 2016, 46, 205-212.	1.3	15
60	TNF-α+ CD4+ TÂcells dominate the SARS-CoV-2 specific T cell response in COVID-19 outpatients and are associated with durable antibodies. Cell Reports Medicine, 2022, 3, 100640.	3.3	15
61	Long-Term Accuracy of Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Interferon-Î ³ Release Assay and Its Application in Household Investigation. Clinical Infectious Diseases, 2022, 75, e314-e321.	2.9	14
62	Genomic DNA microarrays for Entamoeba histolytica: Applications for use in expression profiling and strain genotyping. Experimental Parasitology, 2005, 110, 196-202.	0.5	13
63	Recent advances in Entamoeba biology: RNA interference, drug discovery, and gut microbiome. F1000Research, 2016, 5, 2578.	0.8	13
64	Policy Recommendations for Optimizing the Infectious Diseases Physician-Scientist Workforce. Journal of Infectious Diseases, 2018, 218, S49-S54.	1.9	13
65	High Throughput Sequencing of Entamoeba 27nt Small RNA Population Reveals Role in Permanent Gene Silencing But No Effect on Regulating Gene Expression Changes during Stage Conversion, Oxidative, or Heat Shock Stress. PLoS ONE, 2015, 10, e0134481.	1.1	12
66	Loss of dsRNA-based gene silencing in Entamoeba histolytica: Implications for approaches to genetic analysis. Experimental Parasitology, 2008, 119, 296-300.	0.5	11
67	RNAi Pathway Genes Are Resistant to Small RNA Mediated Gene Silencing in the Protozoan Parasite Entamoeba histolytica. PLoS ONE, 2014, 9, e106477.	1.1	11
68	Functional Characterization of Entamoeba histolytica Argonaute Proteins Reveals a Repetitive DR-Rich Motif Region That Controls Nuclear Localization. MSphere, 2019, 4, .	1.3	10
69	Entamoeba stage conversion: progress and new insights. Current Opinion in Microbiology, 2020, 58, 62-68.	2.3	10
70	The NAD+ Responsive Transcription Factor ERM-BP Functions Downstream of Cellular Aggregation and Is an Early Regulator of Development and Heat Shock Response in Entamoeba. Frontiers in Cellular and Infection Microbiology, 2020, 10, 363.	1.8	10
71	A Single RNaselll Domain Protein from Entamoeba histolytica Has dsRNA Cleavage Activity and Can Help Mediate RNAi Gene Silencing in a Heterologous System. PLoS ONE, 2015, 10, e0133740.	1.1	10
72	SARS-CoV-2 Neutralizing Monoclonal Antibodies for the Treatment of COVID-19 in Kidney Transplant Recipients. Kidney360, 2022, 3, 10.34067/KID.0005732021.	0.9	9

#	Article	IF	Citations
73	Approaches to characterizing Entamoeba histolytica transcriptional regulation. Cellular Microbiology, 2010, 12, 1681-1690.	1.1	8
74	Enteric Amebiasis., 2011,, 614-622.		8
75	Destabilization domain approach adapted for regulated protein expression in the protozoan parasite Entamoeba histolytica. International Journal for Parasitology, 2014, 44, 729-735.	1.3	7
76	Development of a CRISPR/Cas9 system in Entamoeba histolytica: proof of concept. International Journal for Parasitology, 2021, 51, 193-200.	1.3	7
77	Ponatinib, Lestaurtinib, and mTOR/PI3K Inhibitors Are Promising Repurposing Candidates against Entamoeba histolytica. Antimicrobial Agents and Chemotherapy, 2022, 66, AAC0120721.	1.4	7
78	Transcriptional Regulatory Networks in Entamoeba histolytica. Current Drug Targets, 2008, 9, 931-937.	1.0	6
79	Identification of oligo-adenylated small RNAs in the parasite Entamoeba and a potential role for small RNA control. BMC Genomics, 2020, 21, 879.	1.2	6
80	The COVID-19 Outpatient Pragmatic Platform Study (COPPS): Study design of a multi-center pragmatic platform trial. Contemporary Clinical Trials, 2021, 108, 106509.	0.8	5
81	Inflammatory but not respiratory symptoms are associated with ongoing upper airway viral shedding in outpatients with uncomplicated COVID-19. Diagnostic Microbiology and Infectious Disease, 2022, 102, 115612.	0.8	3
82	Variation in Severe Acute Respiratory Syndrome Coronavirus 2 Bioaerosol Production in Exhaled Breath. Open Forum Infectious Diseases, 2022, 9, ofab600.	0.4	3
83	Investigating amoebic pathogenesis usingEntamoeba histolytica DNA microarrays. Journal of Biosciences, 2002, 27, 595-601.	0.5	2
84	DNA Content Analysis on Microarrays. , 2004, 270, 237-248.		2
85	RISC in Entamoeba histolytica: Identification of a Protein-Protein Interaction Network for the RNA Interference Pathway in a Deep-Branching Eukaryote. MBio, 2021, 12, e0154021.	1.8	1
86	Infectious Polymyositis. , 0, , 491-494.		0
87	Supporting Research Career Development of Physician-Scientists. Journal of Infectious Diseases, 2018, 218, S36-S39.	1.9	0
88	Title is missing!. , 2020, 14, e0008150.		0
89	Title is missing!. , 2020, 14, e0008150.		0
90	Title is missing!. , 2020, 14, e0008150.		0

ARTICLE IF CITATIONS
91 Title is missing!., 2020, 14, e0008150. 0