

Marcia B Goldberg

List of Publications by Year in descending order

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Version: 2024-02-01

47
papers

2,972
citations

279701

23
h-index

330025

37
g-index

59
all docs

59
docs citations

59
times ranked

5397
citing authors

#	ARTICLE	IF	CITATIONS
1	Alveolar, Endothelial, and Organ Injury Marker Dynamics in Severe COVID-19. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2022, 205, 507-519.	2.5	56
2	Reply To: High Renin Levels in Severe COVID-19 are Indicative for a Hypo-Renin-Angiotensin-System State. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2022, , .	2.5	0
3	Fc γ R-mediated SARS-CoV-2 infection of monocytes activates inflammation. <i>Nature</i> , 2022, 606, 576-584.	13.7	314
4	The Kinetics of SARS-CoV-2 Antibody Development Is Associated with Clearance of RNAemia. <i>MBio</i> , 2022, 13, .	1.8	10
5	Viral Load Kinetics of Severe Acute Respiratory Syndrome Coronavirus 2 in Hospitalized Individuals With Coronavirus Disease 2019. <i>Open Forum Infectious Diseases</i> , 2021, 8, ofab153.	0.4	20
6	Longitudinal proteomic analysis of severe COVID-19 reveals survival-associated signatures, tissue-specific cell death, and cell-cell interactions. <i>Cell Reports Medicine</i> , 2021, 2, 100287.	3.3	183
7	Plasma from patients with bacterial sepsis or severe COVID-19 induces suppressive myeloid cell production from hematopoietic progenitors in vitro. <i>Science Translational Medicine</i> , 2021, 13, .	5.8	64
8	Plasma ACE2 predicts outcome of COVID-19 in hospitalized patients. <i>PLoS ONE</i> , 2021, 16, e0252799.	1.1	81
9	SARS-CoV-2 viremia is associated with distinct proteomic pathways and predicts COVID-19 outcomes. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	94
10	The type 3 secretion system requires actin polymerization to open translocon pores. <i>PLoS Pathogens</i> , 2021, 17, e1009932.	2.1	8
11	Early cross-coronavirus reactive signatures of humoral immunity against COVID-19. <i>Science Immunology</i> , 2021, 6, eabj2901.	5.6	67
12	Vasopressin infusion in COVID-19 critical illness is not associated with impaired viral clearance: a pilot study. <i>British Journal of Anaesthesia</i> , 2021, 127, e146-e148.	1.5	7
13	Plasma α -selectin is an early marker of thromboembolism in COVID-19. <i>American Journal of Hematology</i> , 2021, 96, E468-E471.	2.0	17
14	Topology and Contribution to the Pore Channel Lining of Plasma Membrane-Embedded <i>Shigella flexneri</i> Type 3 Secretion Translocase IpaB. <i>MBio</i> , 2021, 12, e0302121.	1.8	5
15	Viral epitope profiling of COVID-19 patients reveals cross-reactivity and correlates of severity. <i>Science</i> , 2020, 370, .	6.0	511
16	Direct binding of polymeric GBP1 to LPS disrupts bacterial cell envelope functions. <i>EMBO Journal</i> , 2020, 39, e104926.	3.5	103
17	<i>Shigella flexneri</i> Disruption of Cellular Tension Promotes Intercellular Spread. <i>Cell Reports</i> , 2020, 33, 108409.	2.9	20
18	An immune-cell signature of bacterial sepsis. <i>Nature Medicine</i> , 2020, 26, 333-340.	15.2	261

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19	Isolated Cerebral Mucormycosis in Immunocompetent Adults who Inject Drugs: Case Reports and Systematic Review of the Literature. <i>Open Forum Infectious Diseases</i> , 2020, 7, ofaa552.	0.4	11
20	Activation of <i>Shigella flexneri</i> type 3 secretion requires a host-induced conformational change to the translocon pore. <i>PLoS Pathogens</i> , 2019, 15, e1007928.	2.1	20
21	1830. Single-cell Transcriptional Profiling Reveals an Immune Cell State Signature of Bacterial Sepsis. <i>Open Forum Infectious Diseases</i> , 2019, 6, S42-S42.	0.4	1
22	Topological Analysis of the Type 3 Secretion System Translocon Pore Protein IpaC following Its Native Delivery to the Plasma Membrane during Infection. <i>MBio</i> , 2019, 10, .	1.8	17
23	Rapid identification and phylogenetic classification of diverse bacterial pathogens in a multiplexed hybridization assay targeting ribosomal RNA. <i>Scientific Reports</i> , 2019, 9, 4516.	1.6	11
24	Title is missing!. , 2019, 15, e1007928.		0
25	Title is missing!. , 2019, 15, e1007928.		0
26	Title is missing!. , 2019, 15, e1007928.		0
27	Title is missing!. , 2019, 15, e1007928.		0
28	Rapid Detection of Powassan Virus in a Patient With Encephalitis by Metagenomic Sequencing. <i>Clinical Infectious Diseases</i> , 2018, 66, 789-792.	2.9	41
29	868. Prospective Pathogen Detection in Patients With Central Nervous System Inflammation Using Metagenomic Sequencing. <i>Open Forum Infectious Diseases</i> , 2018, 5, S23-S23.	0.4	0
30	A53â€¢Systematic application of metagenomics NGS to identify and sequence viral pathogens in infections of the central nervous system. <i>Virus Evolution</i> , 2018, 4, .	2.2	0
31	Identification of interactions among host and bacterial proteins and evaluation of their role early during <i>Shigella flexneri</i> infection. <i>Microbiology (United Kingdom)</i> , 2018, 164, 540-550.	0.7	12
32	Intermediate filaments enable pathogen docking to trigger type 3 effector translocation. <i>Nature Microbiology</i> , 2016, 1, 16025.	5.9	58
33	<i>Shigella</i> Effector OspB Activates mTORC1 in a Manner That Depends on IQGAP1 and Promotes Cell Proliferation. <i>PLoS Pathogens</i> , 2015, 11, e1005200.	2.1	32
34	<i>Shigella flexneri</i> Regulation of ARF6 Activation during Bacterial Entry via an IpgD-Mediated Positive Feedback Loop. <i>MBio</i> , 2015, 6, e02584.	1.8	30
35	Vasodilator-stimulated phosphoprotein restricts cell-to-cell spread of <i>Shigella flexneri</i> at the cell periphery. <i>Microbiology (United Kingdom)</i> , 2015, 161, 2149-2160.	0.7	4
36	Systematic Analysis of Bacterial Effector-Postsynaptic Density 95/Disc Large/Zonula Occludens-1 (PDZ) Domain Interactions Demonstrates <i>Shigella</i> OspE Protein Promotes Protein Kinase C Activation via PDLIM Proteins. <i>Journal of Biological Chemistry</i> , 2014, 289, 30101-30113.	1.6	20

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37	Host and Bacterial Proteins That Repress Recruitment of LC3 to Shigella Early during Infection. PLoS ONE, 2014, 9, e94653.	1.1	54
38	Requirement for Formin-Induced Actin Polymerization during Spread of <i>Shigella flexneri</i> . Infection and Immunity, 2010, 78, 193-203.	1.0	56
39	Probing bacterial surfaces using 4Pi spectral self-interference fluorescence microscopy. , 2008, , .		0
40	Polar Localization of the Autotransporter Family of Large Bacterial Virulence Proteins. Journal of Bacteriology, 2006, 188, 4841-4850.	1.0	73
41	Regulation of IcsP, the Outer Membrane Protease of the Shigella Actin Tail Assembly Protein IcsA, by Virulence Plasmid Regulators VirF and VirB. Journal of Bacteriology, 2004, 186, 699-705.	1.0	47
42	The making of a gradient: IcsA (VirG) polarity in <i>Shigella flexneri</i> . Molecular Microbiology, 2002, 41, 861-872.	1.2	93
43	N-WASP deficiency reveals distinct pathways for cell surface projections and microbial actin-based motility. Nature Cell Biology, 2001, 3, 897-904.	4.6	308
44	Periplasmic Transit and Disulfide Bond Formation of the Autotransported Shigella Protein IcsA. Journal of Bacteriology, 2001, 183, 951-958.	1.0	85
45	Characterization of a <i>Vibrio cholerae</i> virulence factor homologous to the family of TonB-dependent proteins. Molecular Microbiology, 1992, 6, 2407-2418.	1.2	58
46	The Spectrum of Salmonella Infection. Infectious Disease Clinics of North America, 1988, 2, 571-598.	1.9	75
47	The <i>Shigella</i> Spp. Type III Effector Protein OspB Is a Cysteine Protease. MBio, 0, , .	1.8	1