David J Gonzalez

List of Publications by Year in descending order

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279798 128289 5,403 61 23 60 citations h-index g-index papers 72 72 72 9314 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The $\hat{l}\pm$ -Arrestin ARRDC3 Is an Emerging Multifunctional Adaptor Protein in Cancer. Antioxidants and Redox Signaling, 2022, 36, 1066-1079.	5.4	8
2	Multi-omics of human plasma reveals molecular features of dysregulated inflammation and accelerated aging in schizophrenia. Molecular Psychiatry, 2022, 27, 1217-1225.	7.9	30
3	Multi-omics analyses of the ulcerative colitis gut microbiome link Bacteroides vulgatus proteases with disease severity. Nature Microbiology, 2022, 7, 262-276.	13.3	110
4	Comparative Analysis of T-Cell Spatial Proteomics and the Influence of HIV Expression. Molecular and Cellular Proteomics, 2022, 21, 100194.	3.8	2
5	Phosphoproteomic analysis of thrombin- and p38 MAPK-regulated signaling networks in endothelial cells. Journal of Biological Chemistry, 2022, 298, 101801.	3.4	8
6	The Host-Microbiome Response to Hyperbaric Oxygen Therapy in Ulcerative Colitis Patients. Cellular and Molecular Gastroenterology and Hepatology, 2022, 14, 35-53.	4.5	10
7	Case Series of Successful Intravenous Immunoglobulin (IVIG) Treatment in 4 Pregnant Patients with Severe COVID-19-Induced Hypoxia. American Journal of Case Reports, 2022, 23, e936734.	0.8	3
8	Dissociation of DNA damage sensing by endoglycosidase HPSE. IScience, 2021, 24, 102242.	4.1	7
9	Urinary Exosomes Identify Inflammatory Pathways in Vancomycin Associated Acute Kidney Injury. International Journal of Molecular Sciences, 2021, 22, 2784.	4.1	17
10	The tumor suppressor kinase DAPK3 drives tumor-intrinsic immunity through the STING–IFN-β pathway. Nature Immunology, 2021, 22, 485-496.	14.5	45
11	CLK1 reorganizes the splicing factor U1-70K for early spliceosomal protein assembly. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118 , .	7.1	13
12	mTORC2 controls the activity of PKC and Akt by phosphorylating a conserved TOR interaction motif. Science Signaling, 2021, 14, .	3.6	64
13	Disruption of innate defense responses by endoglycosidase HPSE promotes cell survival. JCI Insight, 2021, 6, .	5.0	14
14	A Cdk4/6-dependent phosphorylation gradient regulates the early to late G1 phase transition. Scientific Reports, 2021, 11, 14736.	3.3	5
15	Heat shock protein 27 activity is linked to endothelial barrier recovery after proinflammatory GPCR-induced disruption. Science Signaling, 2021, 14, eabc1044.	3.6	23
16	Unique virulence role of post-translocational chaperone PrsA in shaping <i>Streptococcus pyogenes</i> secretome. Virulence, 2021, 12, 2633-2647.	4.4	6
17	Antimicrobials from a feline commensal bacterium inhibit skin infection by drug-resistant S. pseudintermedius. ELife, 2021, 10, .	6.0	14
18	The S Protein of Group B Streptococcus Is a Critical Virulence Determinant That Impacts the Cell Surface Virulome. Frontiers in Microbiology, 2021, 12, 729308.	3.5	4

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19	A combined EM and proteomic analysis places HIV-1 Vpu at the crossroads of retromer and ESCRT complexes: PTPN23 is a Vpu-cofactor. PLoS Pathogens, 2021, 17, e1009409.	4.7	0
20	AssessORF: combining evolutionary conservation and proteomics to assess prokaryotic gene predictions. Bioinformatics, 2020, 36, 1022-1029.	4.1	10
21	Evaluating Organism-Wide Changes in the Metabolome and Microbiome following a Single Dose of Antibiotic. MSystems, 2020, 5, .	3.8	6
22	Kinetic profiling of metabolic specialists demonstrates stability and consistency of in vivo enzyme turnover numbers. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 23182-23190.	7.1	65
23	Development of muscle atrophy and loss of function in a Gulf-War illness model: underlying mechanisms. Scientific Reports, 2020, 10, 14526.	3.3	6
24	Multidimensional Proteome Profiling of Blood-Brain Barrier Perturbation by Group B <i>Streptococcus</i> . MSystems, 2020, 5, .	3.8	7
25	Mortality Risk Profiling of Staphylococcus aureus Bacteremia by Multi-omic Serum Analysis Reveals Early Predictive and Pathogenic Signatures. Cell, 2020, 182, 1311-1327.e14.	28.9	58
26	Exposure of Mycobacterium abscessus to Environmental Stress and Clinically Used Antibiotics Reveals Common Proteome Response among Pathogenic Mycobacteria. Microorganisms, 2020, 8, 698.	3.6	18
27	Molecular dissection of Chagas induced cardiomyopathy reveals central disease associated and druggable signaling pathways. PLoS Neglected Tropical Diseases, 2020, 14, e0007980.	3.0	9
28	Fermentable fiber-induced hepatocellular carcinoma in mice recapitulates gene signatures found in human liver cancer. PLoS ONE, 2020, 15, e0234726.	2.5	4
29	Fermentable fibers induce rapid macro- and micronutrient depletion in Toll-like receptor 5-deficient mice. American Journal of Physiology - Renal Physiology, 2020, 318, G955-G965.	3.4	3
30	Phosphoproteomic analysis of protease-activated receptor-1 biased signaling reveals unique modulators of endothelial barrier function. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 5039-5048.	7.1	25
31	Organ-level protein networks as a reference for the host effects of the microbiome. Genome Research, 2020, 30, 276-286.	5. 5	6
32	Identification of a Human Skin Commensal Bacterium that Selectively Kills CutibacteriumÂacnes. Journal of Investigative Dermatology, 2020, 140, 1619-1628.e2.	0.7	47
33	A Hetero-Multimeric Chitinase-Containing Plasmodium falciparum and Plasmodium gallinaceum Ookinete-Secreted Protein Complex Involved in Mosquito Midgut Invasion. Frontiers in Cellular and Infection Microbiology, 2020, 10, 615343.	3.9	4
34	Functional and Proteomic Analysis of Streptococcus pyogenes Virulence Upon Loss of Its Native Cas9 Nuclease. Frontiers in Microbiology, 2019, 10, 1967.	3.5	11
35	Associations of the Fecal Microbial Proteome Composition and Proneness to Diet-induced Obesity. Molecular and Cellular Proteomics, 2019, 18, 1864-1879.	3.8	19
36	Multiantigenic Nanotoxoids for Antivirulence Vaccination against Antibiotic-Resistant Gram-Negative Bacteria. Nano Letters, 2019, 19, 4760-4769.	9.1	63

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37	EGFR is required for Wnt9a–Fzd9b signalling specificity in haematopoietic stem cells. Nature Cell Biology, 2019, 21, 721-730.	10.3	42
38	Neutrophilic proteolysis in the cystic fibrosis lung correlates with a pathogenic microbiome. Microbiome, 2019, 7, 23.	11.1	53
39	Quantitative Multiplex Substrate Profiling of Peptidases by Mass Spectrometry. Molecular and Cellular Proteomics, 2019, 18, 968a-981.	3.8	28
40	Evaluating Metagenomic Prediction of the Metaproteome in a 4.5-Year Study of a Patient with Crohn's Disease. MSystems, 2019, 4, .	3.8	40
41	Quantitative analysis of Mycobacterium avium subsp. hominissuis proteome in response to antibiotics and during exposure to different environmental conditions. Clinical Proteomics, 2019, 16, 39.	2.1	23
42	Group A Streptococcal S Protein Utilizes Red Blood Cells as Immune Camouflage and Is a Critical Determinant for Immune Evasion. Cell Reports, 2019, 29, 2979-2989.e15.	6.4	16
43	Essential metabolism for a minimal cell. ELife, 2019, 8, .	6.0	110
44	<i>PTMphinder</i> : an R package for PTM site localization and motif extraction from proteomic datasets. PeerJ, 2019, 7, e7046.	2.0	8
45	Context-Dependent and Disease-Specific Diversity in Protein Interactions within Stress Granules. Cell, 2018, 172, 590-604.e13.	28.9	672
46	Identification of Common and Rare Genetic Variation Associated With Plasma Protein Levels Using Whole-Exome Sequencing and Mass Spectrometry. Circulation Genomic and Precision Medicine, 2018, 11, e002170.	3.6	26
47	Optimization of carbon and energy utilization through differential translational efficiency. Nature Communications, 2018, 9, 4474.	12.8	35
48	Dysregulated Microbial Fermentation of Soluble Fiber Induces Cholestatic Liver Cancer. Cell, 2018, 175, 679-694.e22.	28.9	344
49	A Tyrosine Switch on NEDD4-2 E3 Ligase Transmits GPCR Inflammatory Signaling. Cell Reports, 2018, 24, 3312-3323.e5.	6.4	36
50	Defining Host Responses during Systemic Bacterial Infection through Construction of a Murine Organ Proteome Atlas. Cell Systems, 2018, 6, 579-592.e4.	6.2	23
51	Staphylococcus aureus Membrane-Derived Vesicles Promote Bacterial Virulence and Confer Protective Immunity in Murine Infection Models. Frontiers in Microbiology, 2018, 9, 262.	3.5	65
52	Group B Streptococcus Biofilm Regulatory Protein A Contributes to Bacterial Physiology and Innate Immune Resistance. Journal of Infectious Diseases, 2018, 218, 1641-1652.	4.0	38
53	Pseudopodium-enriched atypical kinase 1 mediates angiogenesis by modulating GATA2-dependent VEGFR2 transcription. Cell Discovery, 2018, 4, 26.	6.7	19
54	Identification of the S-transferase like superfamily bacillithiol transferases encoded by Bacillus subtilis. PLoS ONE, 2018, 13, e0192977.	2.5	8

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55	Contextâ€dependent and Diseaseâ€specific Diversity in Stress Granules Formed from Preâ€existing Protein Interactions. FASEB Journal, 2018, 32, 252.3.	0.5	2
56	Quantitative Temporal Viromics of an Inducible HIV-1 Model Yields Insight to Global Host Targets and Phospho-Dynamics Associated with Protein Vpr. Molecular and Cellular Proteomics, 2017, 16, 1447-1461.	3.8	60
57	Biomimetic Virulomics for Capture and Identification of Cell-Type Specific Effector Proteins. ACS Nano, 2017, 11, 11831-11838.	14.6	27
58	Mitochondrial H+-ATP synthase in human skeletal muscle: contribution to dyslipidaemia and insulin resistance. Diabetologia, 2017, 60, 2052-2065.	6.3	32
59	Phosphorylation of serine96 of histidine-rich calcium-binding protein by the Fam20C kinase functions to prevent cardiac arrhythmia. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 9098-9103.	7.1	43
60	Sharing and community curation of mass spectrometry data with Global Natural Products Social Molecular Networking. Nature Biotechnology, 2016, 34, 828-837.	17.5	2,802
61	Tamoxifen augments the innate immune function of neutrophils through modulation of intracellular ceramide. Nature Communications, 2015, 6, 8369.	12.8	98