## Stephanie Byrum

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3249331/publications.pdf

Version: 2024-02-01

279798 233421 2,511 87 23 45 citations g-index h-index papers 94 94 94 3854 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	DNA methylation on N6-adenine in mammalian embryonic stem cells. Nature, 2016, 532, 329-333.	27.8	554
2	Phase separation drives aberrant chromatin looping and cancer development. Nature, 2021, 595, 591-595.	27.8	197
3	ChAP-MS: A Method for Identification of Proteins and Histone Posttranslational Modifications at a Single Genomic Locus. Cell Reports, 2012, 2, 198-205.	6.4	110
4	ATXN7L3 and ENY2 Coordinate Activity of Multiple H2B Deubiquitinases Important for Cellular Proliferation and Tumor Growth. Molecular Cell, 2016, 62, 558-571.	9.7	106
5	Multi-omics data integration considerations and study design for biological systems and disease. Molecular Omics, 2021, 17, 170-185.	2.8	85
6	Anti-PD-1/L1 lead-in before MAPK inhibitor combination maximizes antitumor immunity and efficacy. Cancer Cell, 2021, 39, 1375-1387.e6.	16.8	78
7	A CRISPR-based approach for proteomic analysis of a single genomic locus. Epigenetics, 2014, 9, 1207-1211.	2.7	71
8	Indicators of responsiveness to immune checkpoint inhibitors. Scientific Reports, 2017, 7, 807.	3.3	70
9	proteiNorm – A User-Friendly Tool for Normalization and Analysis of TMT and Label-Free Protein Quantification. ACS Omega, 2020, 5, 25625-25633.	3.5	53
10	Purification of a specific native genomic locus for proteomic analysis. Nucleic Acids Research, 2013, 41, e195-e195.	14.5	49
11	A PWWP Domain-Containing Protein Targets the NuA3 Acetyltransferase Complex via Histone H3 Lysine 36 trimethylation to Coordinate Transcriptional Elongation at Coding Regions. Molecular and Cellular Proteomics, 2014, 13, 2883-2895.	3.8	48
12	Impact of <i>sarA</i> and Phenol-Soluble Modulins on the Pathogenesis of Osteomyelitis in Diverse Clinical Isolates of Staphylococcus aureus. Infection and Immunity, 2016, 84, 2586-2594.	2.2	46
13	Loss of E-Cadherin Inhibits CD103 Antitumor Activity and Reduces Checkpoint Blockade Responsiveness in Melanoma. Cancer Research, 2019, 79, 1113-1123.	0.9	45
14	Durable Suppression of Acquired MEK Inhibitor Resistance in Cancer by Sequestering MEK from ERK and Promoting Antitumor T-cell Immunity. Cancer Discovery, 2021, 11, 714-735.	9.4	45
15	Quantitative Proteomics Identifies Activation of Hallmark Pathways of Cancer in Patient Melanoma. Journal of Proteomics and Bioinformatics, 2013, 06, 43-50.	0.4	43
16	Impact of Spaceflight and Artificial Gravity on the Mouse Retina: Biochemical and Proteomic Analysis. International Journal of Molecular Sciences, 2018, 19, 2546.	4.1	41
17	Spaceflight induces oxidative damage to bloodâ€brain barrier integrity in a mouse model. FASEB Journal, 2020, 34, 15516-15530.	0.5	39
18	Analysis of Stable and Transient Protein–Protein Interactions. Methods in Molecular Biology, 2012, 833, 143-152.	0.9	36

#	Article	IF	Citations
19	Mitotic phosphorylation of histone H3 threonine 80. Cell Cycle, 2014, 13, 440-452.	2.6	32
20	Characterization of mouse ocular response to a 35-day spaceflight mission: Evidence of blood-retinal barrier disruption and ocular adaptations. Scientific Reports, 2019, 9, 8215.	3.3	30
21	A NSD3-targeted PROTAC suppresses NSD3 and cMyc oncogenic nodes in cancer cells. Cell Chemical Biology, 2022, 29, 386-397.e9.	5.2	30
22	ZMYND11-MBTD1 induces leukemogenesis through hijacking NuA4/TIP60 acetyltransferase complex and a PWWP-mediated chromatin association mechanism. Nature Communications, 2021, 12, 1045.	12.8	27
23	Quantitative Histone Mass Spectrometry Identifies Elevated Histone H3 Lysine 27 (Lys27) Trimethylation in Melanoma. Molecular and Cellular Proteomics, 2016, 15, 765-775.	3.8	26
24	Metaproteomics reveals potential mechanisms by which dietary resistant starch supplementation attenuates chronic kidney disease progression in rats. PLoS ONE, 2019, 14, e0199274.	2.5	25
25	Time- and radiation-dose dependent changes in the plasma proteome after total body irradiation of non-human primates: Implications for biomarker selection. PLoS ONE, 2017, 12, e0174771.	2.5	25
26	Circulating Exosomal microRNAs as Predictive Biomarkers of Neoadjuvant Chemotherapy Response in Breast Cancer. Current Oncology, 2022, 29, 613-630.	2.2	24
27	A Proteomic Study of Human Merkel Cell Carcinoma. Journal of Proteomics and Bioinformatics, 2013, 06, 275-282.	0.4	23
28	Cistrome analysis of YY1 uncovers a regulatory axis of YY1:BRD2/4-PFKP during tumorigenesis of advanced prostate cancer. Nucleic Acids Research, 2021, 49, 4971-4988.	14.5	22
29	ProteoViz: a tool for the analysis and interactive visualization of phosphoproteomics data. Molecular Omics, 2020, 16, 316-326.	2.8	19
30	Epigenetic Control of <i>Cdkn2a.Arf</i> Protects Tumor-Infiltrating Lymphocytes from Metabolic Exhaustion. Cancer Research, 2020, 80, 4707-4719.	0.9	19
31	1,3-Butadiene-induced mitochondrial dysfunction is correlated with mitochondrial CYP2E1 activity in Collaborative Cross mice. Toxicology, 2017, 378, 114-124.	4.2	18
32	Label-Free Proteomic Approach to Characterize Protease-Dependent and -Independent Effects of <i>sarA</i> Inactivation on the <i>Staphylococcus aureus</i> Exoproteome. Journal of Proteome Research, 2018, 17, 3384-3395.	3.7	18
33	Formula Diet Alters the Ileal Metagenome and Transcriptome at Weaning and during the Postweaning Period in a Porcine Model. MSystems, 2020, 5, .	3.8	18
34	Discovery of a dual WDR5 and Ikaros PROTAC degrader as an anti-cancer therapeutic. Oncogene, 2022, 41, 3328-3340.	5.9	18
35	Proteomic analysis of bone cancer: a review of current and future developments. Expert Review of Proteomics, 2007, 4, 371-378.	3.0	17
36	Proteomics-Based Identification of Differentially Abundant Proteins from Human Keratinocytes Exposed to Arsenic Trioxide. Journal of Proteomics and Bioinformatics, 2014, 07, 166-178.	0.4	17

#	Article	IF	CITATIONS
37	RNA-Seq Analysis of Spinal Cord Tissues from hPFN1G118V Transgenic Mouse Model of ALS at Pre-symptomatic and End-Stages of Disease. Scientific Reports, 2018, 8, 13737.	3.3	16
38	Quantitative Analysis of Histone Exchange during Chromatin Purification. Journal of Integrated OMICS, 2011, 1, 61-65.	0.5	15
39	SarA plays a predominant role in controlling the production of extracellular proteases in the diverse clinical isolates of <i>Staphylococcus aureus </i> LAC and UAMS-1. Virulence, 2020, 11, 1738-1762.	4.4	15
40	A conserved BAH module within mammalian BAHD1 connects H3K27me3 to Polycomb gene silencing. Nucleic Acids Research, 2021, 49, 4441-4455.	14.5	15
41	The promise of bone cancer proteomics. Annals of the New York Academy of Sciences, 2010, 1192, 222-229.	3.8	14
42	Exploiting Correlations between Protein Abundance and the Functional Status of <i>saeRS</i> and <i>sarA</i> To Identify Virulence Factors of Potential Importance in the Pathogenesis of <i>Staphylococcus aureus</i> Osteomyelitis. ACS Infectious Diseases, 2020, 6, 237-249.	3.8	14
43	Activation-induced cytidine deaminase localizes to G-quadruplex motifs at mutation hotspots in lymphoma. NAR Cancer, 2020, 2, zcaa029.	3.1	14
44	Assessing the Effects of Redox Modifier MnTnBuOE-2-PyP 5+ on Cognition and Hippocampal Physiology Following Doxorubicin, Cyclophosphamide, and Paclitaxel Treatment. International Journal of Molecular Sciences, 2020, 21, 1867.	4.1	14
45	Quantitative analysis of histone exchange for transcriptionally active chromatin. Journal of Clinical Bioinformatics, 2011, 1, 17.	1.2	13
46	Differences in cell death in methionine versus cysteine depletion. Environmental and Molecular Mutagenesis, 2021, 62, 216-226.	2.2	13
47	Raman Spectroscopy and Machine Learning Reveals Early Tumor Microenvironmental Changes Induced by Immunotherapy. Cancer Research, 2021, 81, 5745-5755.	0.9	13
48	Proteomics Indicates Lactate Dehydrogenase Is Prognostic in Acetaminophen-Induced Acute Liver Failure Patients and Reveals Altered Signaling Pathways. Toxicological Sciences, 2022, 187, 25-34.	3.1	13
49	Identification of Viral and Host Proteins That Interact with Murine Gammaherpesvirus 68 Latency-Associated Nuclear Antigen during Lytic Replication: a Role for Hsc70 in Viral Replication. Journal of Virology, 2016, 90, 1397-1413.	3.4	12
50	Proteomic identification of histone post-translational modifications and proteins enriched at a DNA double-strand break. Nucleic Acids Research, 2017, 45, 10923-10940.	14.5	12
51	Local and Relayed Effects of Deep Brain Stimulation of the Pedunculopontine Nucleus. Brain Sciences, 2019, 9, 64.	2.3	12
52	Respiratory defects in the <i>Crtap</i> KO mouse model of osteogenesis imperfecta. American Journal of Physiology - Lung Cellular and Molecular Physiology, 2020, 318, L592-L605.	2.9	11
53	Continuous Developmental and Early Life Trichloroethylene Exposure Promoted DNA Methylation Alterations in Polycomb Protein Binding Sites in Effector/Memory CD4+ T Cells. Frontiers in Immunology, 2019, 10, 2016.	4.8	10
54	Inhibition of tryptophan 2,3-dioxygenase impairs DNA damage tolerance and repair in glioma cells. NAR Cancer, 2021, 3, zcab014.	3.1	10

#	Article	IF	CITATIONS
55	Cognitive impairment resulting from treatment with docetaxel, doxorubicin, and cyclophosphamide. Brain Research, 2021, 1760, 147397.	2.2	10
56	Accurate and Sensitive Quantitation of the Dynamic Heat Shock Proteome Using Tandem Mass Tags. Journal of Proteome Research, 2020, 19, 1183-1195.	3.7	9
57	Multi-omics data integration reveals correlated regulatory features of triple negative breast cancer. Molecular Omics, 2021, 17, 677-691.	2.8	9
58	Purification of Specific Chromatin Loci for Proteomic Analysis. Methods in Molecular Biology, 2015, 1228, 83-92.	0.9	9
59	Control of the Anterior Pituitary Cell Lineage Regulator POU1F1 by the Stem Cell Determinant Musashi. Endocrinology, 2021, 162, .	2.8	9
60	Ionizing Radiation Activates Mitochondrial Function in Osteoclasts and Causes Bone Loss in Young Adult Male Mice. International Journal of Molecular Sciences, 2022, 23, 675.	4.1	9
61	In vivo transcriptional analysis of mice infected with Leishmania major unveils cellular heterogeneity and altered transcriptomic profiling at single-cell resolution. PLoS Neglected Tropical Diseases, 2022, 16, e0010518.	3.0	9
62	A Quantitative Proteomic Analysis of Urine from Gamma-Irradiated Non-Human Primates. Journal of Proteomics and Bioinformatics, 2014, 01, .	0.4	7
63	Effect of Sulforaphane and 5-Aza-2'-Deoxycytidine on Melanoma Cell Growth. Medicines (Basel,) Tj ETQq1 1	0.784314 1.4	rgBT  Overlo
64	Cranial irradiation impairs juvenile social memory and modulates hippocampal physiology. Brain Research, 2020, 1748, 147095.	2.2	7
65	Neonatal Diet Impacts Circulatory miRNA Profile in a Porcine Model. Frontiers in Immunology, 2020, 11, 1240.	4.8	6
66	Proteogenomic analysis of melanoma brain metastases from distinct anatomical sites identifies pathways of metastatic progression. Acta Neuropathologica Communications, 2020, 8, 157.	<b>5.</b> 2	5
67	PHF19 inhibition as a therapeutic target in multiple myeloma. Current Research in Translational Medicine, 2021, 69, 103290.	1.8	5
68	The Effects of 5-Fluorouracil/Leucovorin Chemotherapy on Cognitive Function in Male Mice. Frontiers in Molecular Biosciences, 2021, 8, 762116.	3 <b>.</b> 5	5
69	Proteomic characterization of the arsenic response locus in S. cerevisiae. Epigenetics, 2019, 14, 130-145.	2.7	4
70	Cold Storage Increases Albumin and Advanced Glycation-End Product-Albumin Levels in Kidney Transplants: A Possible Cause for Exacerbated Renal Damage. Transplantation Direct, 2019, 5, e454.	1.6	4
71	PTMViz: a tool for analyzing and visualizing histone post translational modification data. BMC Bioinformatics, 2021, 22, 275.	2.6	4
72	Deficiency of N-glycanase 1 perturbs neurogenesis and cerebral development modeled by human organoids. Cell Death and Disease, 2022, 13, 262.	6.3	4

#	Article	IF	CITATIONS
73	Proteomic profiling of tear fluid as a promising non-invasive screening test for colon cancer. American Journal of Surgery, 2022, 224, 19-24.	1.8	4
74	Misregulation of Rad50 expression inÂmelanoma cells. Journal of Cutaneous Pathology, 2012, 39, 680-684.	1.3	3
75	Dysbiotic stress increases the sensitivity of the tumor vasculature to radiotherapy and c-Met inhibitors. Angiogenesis, 2021, 24, 597-611.	7.2	3
76	Proteomic Technologies for the Study of Osteosarcoma. Sarcoma, 2012, 2012, 1-10.	1.3	2
77	Vulvar squamous cell carcinoma aggressiveness is associated with differential expression of collagen and STAT1. Clinical Proteomics, 2017, 14, 40.	2.1	2
78	Milk Formula Diet Alters Bacterial and Host Protein Profile in Comparison to Human Milk Diet in Neonatal Piglet Model. Nutrients, 2021, 13, 3718.	4.1	2
79	Epigenetically Enhanced MED12L in ETO2-GLIS2 Positive Pediatric Acute Megakaryoblastic Leukemia Is Associated with Resistance to the CDK8 Inhibitors. Blood, 2021, 138, 2208-2208.	1.4	2
80	Short-Term Metformin Treatment Enriches Bacteroides dorei in an Obese Liver Steatosis Zucker Rat Model. Frontiers in Microbiology, 2022, 13, 834776.	3.5	2
81	HIF-α Activation Impacts Macrophage Function during Murine Leishmania major Infection. Pathogens, 2021, 10, 1584.	2.8	2
82	Effect of excess weight and insulin resistance on DNA methylation in prepubertal children. Scientific Reports, 2022, 12, 8430.	3.3	2
83	Registered report protocol: Quantitative analysis of septin Cdc10-associated proteome in Cryptococcus neoformans. PLoS ONE, 2020, 15, e0242381.	2.5	1
84	PSII-29 Identification of serum proteins that interact with extracellular histones in feedlot cattle. Journal of Animal Science, 2019, 97, 246-246.	0.5	0
85	Phosphoproteomics Provides Novel Insights into the Response of Primary Acute Lymphoblastic Leukemia Cells to Microtubule Depolymerization in G1 Phase of the Cell Cycle. ACS Omega, 2021, 6, 24949-24959.	3.5	O
86	PTHrP(12-48) for the diagnosis of breast cancer bone metastasis Journal of Clinical Oncology, 2012, 30, e21039-e21039.	1.6	0
87	Methamphetamineâ€Induced Proteomic Changes Within the Neuroinflammatory TLR4 Pathway Persist After Longâ€Term Selfâ€Administration in Rats. FASEB Journal, 2022, 36, .	0.5	O