

Susan Bullman

List of Publications by Year in descending order

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39
papers

10,398
citations

304743
22
h-index

315739
38
g-index

43
all docs

43
docs citations

43
times ranked

15748
citing authors

#	ARTICLE	IF	CITATIONS
1	The Immune Landscape of Cancer. <i>Immunity</i> , 2018, 48, 812-830.e14.	14.3	3,706
2	Comprehensive Molecular Characterization of Muscle-Invasive Bladder Cancer. <i>Cell</i> , 2017, 171, 540-556.e25.	28.9	1,742
3	Analysis of <i>Fusobacterium</i> persistence and antibiotic response in colorectal cancer. <i>Science</i> , 2017, 358, 1443-1448.	12.6	983
4	<i>Fusobacterium nucleatum</i> in colorectal carcinoma tissue and patient prognosis. <i>Gut</i> , 2016, 65, 1973-1980.	12.1	718
5	Commensal Microbiota Promote Lung Cancer Development via $\gamma\delta$ T Cells. <i>Cell</i> , 2019, 176, 998-1013.e16.	28.9	592
6	<i>Fusobacterium nucleatum</i> and T Cells in Colorectal Carcinoma. <i>JAMA Oncology</i> , 2015, 1, 653.	7.1	498
7	Comparative Molecular Analysis of Gastrointestinal Adenocarcinomas. <i>Cancer Cell</i> , 2018, 33, 721-735.e8.	16.8	396
8	Genomic, Pathway Network, and Immunologic Features Distinguishing Squamous Carcinomas. <i>Cell Reports</i> , 2018, 23, 194-212.e6.	6.4	245
9	Association of Dietary Patterns With Risk of Colorectal Cancer Subtypes Classified by <i>Fusobacterium nucleatum</i> in Tumor Tissue. <i>JAMA Oncology</i> , 2017, 3, 921.	7.1	243
10	<i>Fusobacterium nucleatum</i> in Colorectal Carcinoma Tissue According to Tumor Location. <i>Clinical and Translational Gastroenterology</i> , 2016, 7, e200.	2.5	225
11	<i>Fusobacterium nucleatum</i> in Colorectal Cancer Relates to Immune Response Differentially by Tumor Microsatellite Instability Status. <i>Cancer Immunology Research</i> , 2018, 6, 1327-1336.	3.4	127
12	Diets That Promote Colon Inflammation Associate With Risk of Colorectal Carcinomas That Contain <i>Fusobacterium nucleatum</i> . <i>Clinical Gastroenterology and Hepatology</i> , 2018, 16, 1622-1631.e3.	4.4	103
13	GATK PathSeq: a customizable computational tool for the discovery and identification of microbial sequences in libraries from eukaryotic hosts. <i>Bioinformatics</i> , 2018, 34, 4287-4289.	4.1	70
14	The Amount of Bifidobacterium Genus in Colorectal Carcinoma Tissue in Relation to Tumor Characteristics and Clinical Outcome. <i>American Journal of Pathology</i> , 2018, 188, 2839-2852.	3.8	51
15	<i>Campylobacter ureolyticus</i> : an emerging gastrointestinal pathogen?. <i>FEMS Immunology and Medical Microbiology</i> , 2011, 61, 228-230.	2.7	42
16	Emerging dynamics of human campylobacteriosis in Southern Ireland. <i>FEMS Immunology and Medical Microbiology</i> , 2011, 63, 248-253.	2.7	39
17	Association of <i>Fusobacterium nucleatum</i> with Specific T-cell Subsets in the Colorectal Carcinoma Microenvironment. <i>Clinical Cancer Research</i> , 2021, 27, 2816-2826.	7.0	36
18	Molecular diagnostics. <i>Bioengineered</i> , 2012, 3, 1-7.	3.2	28

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19	Identification and genetic characterization of a novel picornavirus from chickens. Journal of General Virology, 2014, 95, 1094-1103.	2.9	28
20	Cancer as microenvironmental, systemic and environmental diseases: opportunity for transdisciplinary microbiomics science. Gut, 2022, 71, 2107-2122.	12.1	28
21	Association of autophagy status with amount of <i>Fusobacterium nucleatum</i> in colorectal cancer. Journal of Pathology, 2020, 250, 397-408.	4.5	27
22	The relationship between gastrointestinal cancers and the microbiota. The Lancet Gastroenterology and Hepatology, 2021, 6, 498-509.	8.1	25
23	Metagenomic Characterization of Microbial Communities In Situ Within the Deeper Layers of the Ileum in Crohn's Disease. Cellular and Molecular Gastroenterology and Hepatology, 2016, 2, 563-566.e5.	4.5	23
24	Patients with mesenchymal tumours and high <i>Fusobacteriales</i> prevalence have worse prognosis in colorectal cancer (CRC). Gut, 2021, , gutjnl-2021-325193.	12.1	23
25	Genomic Investigation into Strain Heterogeneity and Pathogenic Potential of the Emerging Gastrointestinal Pathogen <i>Campylobacter ureolyticus</i> . PLoS ONE, 2013, 8, e71515.	2.5	21
26	Molecular and Pathology Features of Colorectal Tumors and Patient Outcomes Are Associated with <i>Fusobacterium nucleatum</i> and Its Subspecies <i>animalis</i> . Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 210-220.	2.5	19
27	Emerging Concepts and Technologies for the Discovery of Microorganisms Involved in Human Disease. Annual Review of Pathology: Mechanisms of Disease, 2017, 12, 217-244.	22.4	13
28	Molecular-based detection of the gastrointestinal pathogen <i>Campylobacter ureolyticus</i> in unpasteurized milk samples from two cattle farms in Ireland. Gut Pathogens, 2012, 4, 14.	3.4	12
29	Comprehensive metagenomic analysis of blastic plasmacytoid dendritic cell neoplasm. Blood Advances, 2020, 4, 1006-1011.	5.2	10
30	Harnessing the microbiome to restore immunotherapy response. Nature Cancer, 2021, 2, 1301-1304.	13.2	10
31	The tumour-associated microbiome. Nature Reviews Gastroenterology and Hepatology, 2022, 19, 347-348.	17.8	9
32	Epsilonproteobacteria in Humans, New Zealand. Emerging Infectious Diseases, 2012, 18, 1709-1710.	4.3	4
33	Mechanistic Insights into Transmissible Cancers of Mammals. Cancer Cell, 2018, 33, 543-544.	16.8	4
34	Draft Genome Sequence of <i>Campylobacter ureolyticus</i> Strain CIT007, the First Whole-Genome Sequence of a Clinical Isolate. Genome Announcements, 2014, 2, .	0.8	3
35	Draft Genome Sequence of <i>Campylobacter corcagiensis</i> Strain CIT045 T, a Representative of a Novel <i>Campylobacter</i> Species Isolated from Lion-Tailed Macaques (<i>Macaca silenus</i>). Genome Announcements, 2014, 2, .	0.8	3
36	Complete Genome Sequence of <i>Morganella morganii</i> CTX51T, Isolated from a Human Cecal Adenocarcinoma. Microbiology Resource Announcements, 2022, 11, e0006622.	0.6	2

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37	Complete Genome Sequence of <i>Clostridium cadaveris</i> IFB3C5, Isolated from a Human Colonic Adenocarcinoma. <i>Microbiology Resource Announcements</i> , 2022, 11, e0113521.	0.6	1
38	<i>Bifidobacterium</i> Genus in Colorectal Carcinoma Tissue in relation to Tumor Characteristics and Patient Survival. <i>FASEB Journal</i> , 2018, 32, 407.3.	0.5	0
39	Metagenomic analysis to identify novel infectious agents in systemic anaplastic large cell lymphoma. <i>Infectious Agents and Cancer</i> , 2021, 16, 65.	2.6	0