Pam Tolomeo

List of Publications by Year
in descending order

47 papers	621 citations	623734 14 h-index	642732 23 g-index
48 all docs	48 docs citations	48 times ranked	935 citing authors

#	Article	IF	Citations
1	The shared microbiota of humans and companion animals as evaluated from Staphylococcus carriage sites. Microbiome, 2015, 3, 2.	11.1	95
2	Association of 30-Day Mortality With Oral Step-Down vs Continued Intravenous Therapy in Patients Hospitalized With Enterobacteriaceae Bacteremia. JAMA Internal Medicine, 2019, 179, 316.	5.1	94
3	Use of a Combination Biomarker Algorithm To Identify Medical Intensive Care Unit Patients with Suspected Sepsis at Very Low Likelihood of Bacterial Infection. Antimicrobial Agents and Chemotherapy, 2015, 59, 6494-6500.	3.2	32
4	Combined Biomarkers Predict Acute Mortality Among Critically Ill Patients With Suspected Sepsis*. Critical Care Medicine, 2018, 46, 1106-1113.	0.9	27
5	Whole-Genome Sequencing To Identify Drivers of Carbapenem-Resistant Klebsiella pneumoniae Transmission within and between Regional Long-Term Acute-Care Hospitals. Antimicrobial Agents and Chemotherapy, 2019, 63, .	3.2	24
6	Risk factors for multidrug‐resistant organisms among deceased organ donors. American Journal of Transplantation, 2019, 19, 2468-2478.	4.7	24
7	Risk Factors for Extended-Spectrum Î²-lactamase‐Producing Enterobacterales Bloodstream Infection Among Solid-Organ Transplant Recipients. Clinical Infectious Diseases, 2021, 72, 953-960.	5.8	22
8	The role of extended-spectrum cephalosporin-resistance in recurrent community-onset Enterobacteriaceae urinary tract infections: a retrospective cohort study. BMC Infectious Diseases, 2019, 19, 163.	2.9	21
9	The Effect of Total Household Decolonization on Clearance of Colonization With Methicillin-Resistant <i>Staphylococcus aureus</i>. Infection Control and Hospital Epidemiology, 2016, 37, 1226-1233.	1.8	19
10	Combined biomarkers discriminate a low likelihood of bacterial infection among surgical intensive care unit patients with suspected sepsis. Diagnostic Microbiology and Infectious Disease, 2016, 85, 109-115.	1.8	19
11	Genome sequencing reveals strain dynamics of methicillin-resistant Staphylococcus aureus in the same household in the context of clinical disease in a person and a dog. Veterinary Microbiology, 2015, 180, 304-307.	1.9	18
12	Quantifying the Impact of Nasopharyngeal Specimen Quality on Severe Acute Respiratory Syndrome Coronavirus 2 Test Performance. Open Forum Infectious Diseases, 2021, 8, ofab235.	0.9	18
13	Poor clinical outcomes associated with community-onset urinary tract infections due to extended-spectrum cephalosporin-resistant Enterobacteriaceae. Infection Control and Hospital Epidemiology, 2018, 39, 1431-1435.	1.8	17
14	Impact of donor multidrug‐resistant organisms on solid organ transplant recipient outcomes. Transplant Infectious Disease, 2022, 24, e13783.	1.7	17
15	Risk Factors for Recurrent Colonization With Methicillin-ResistantStaphylococcus aureusin Community-Dwelling Adults and Children. Infection Control and Hospital Epidemiology, 2015, 36, 786-793.	1.8	16
16	Implementation of a Pragmatic Biomarker-Driven Algorithm to Guide Antibiotic Use in the Pediatric Intensive Care Unit: the Optimizing Antibiotic Strategies in Sepsis (OASIS) II Study. Journal of the Pediatric Infectious Diseases Society, 2020, 9, 36-43.	1.3	15
17	Impact of deceased donor multidrug-resistant bacterial organisms on organ utilization. American Journal of Transplantation, 2020, 20, 2559-2566.	4.7	14
18	Clinical and Molecular Characterization of Community-Onset Urinary Tract Infections Due to Extended-Spectrum Cephalosporin-Resistant Enterobacteriaceae. Infection Control and Hospital Epidemiology, 2016, 37, 1433-1439.	1.8	13

#	Article	IF	Citations
19	Comparison of Culture-Based Methods for Identification of Colonization with Methicillin-Resistant and Methicillin-Susceptible Staphylococcus aureus in the Context of Cocolonization. Journal of Clinical Microbiology, 2016, 54, 1907-1911.	3.9	11
20	Clinical Characteristics and Outcomes of Hematologic Malignancy Patients With Positive Clostridium difficile Toxin Immunoassay Versus Polymerase Chain Reaction Test Results. Infection Control and Hospital Epidemiology, 2018, 39, 863-866.	1.8	9
21	Improving Outpatient Antibiotic Prescribing for Respiratory Tract Infections in Primary Care: A Stepped-Wedge Cluster Randomized Trial. Clinical Infectious Diseases, 2022, 74, 947-956.	5.8	9
22	The Impact of Centers for Medicare & Medicaid Services SEP-1 Core Measure Implementation on Antibacterial Utilization: A Retrospective Multicenter Longitudinal Cohort Study With Interrupted Time-Series Analysis. Clinical Infectious Diseases, 2022, 75, 503-511.	5.8	9
23	Clinical and Molecular Epidemiology of Escherichia coli Sequence Type 131 among Hospitalized Patients Colonized Intestinally with Fluoroquinolone-Resistant E. coli. Antimicrobial Agents and Chemotherapy, 2014, 58, 7003-7006.	3.2	8
24	Epidemiology and characteristics of Escherichia coli sequence type 131 (ST131) from long-term care facility residents colonized intestinally with fluoroquinolone-resistant Escherichia coli. Diagnostic Microbiology and Infectious Disease, 2017, 87, 275-280.	1.8	8
25	Spatial and temporal effects on severe acute respiratory coronavirus virus 2 (SARS-CoV-2) contamination of the healthcare environment. Infection Control and Hospital Epidemiology, 2021, , 1-6.	1.8	8
26	Stopping Hospital Infections With Environmental Services (SHINE): A Cluster-randomized Trial of Intensive Monitoring Methods for Terminal Room Cleaning on Rates of Multidrug-resistant Organisms in the Intensive Care Unit. Clinical Infectious Diseases, 2022, 75, 1217-1223.	5.8	7
27	Pet Ownership Protects Against Recurrence of Clostridioides difficile Infection. Open Forum Infectious Diseases, 2020, 7, ofz541.	0.9	6
28	Risk Factors for Infection with <i>Escherichia coli</i> in Nursing Home Residents Colonized with Fluoroquinolone-Resistant <i>E. coli</i>. Infection Control and Hospital Epidemiology, 2015, 36, 575-577.	1.8	5
29	Clinical prediction tool for extended‐spectrum beta‐lactamase‐producing enterobacterales as the etiology of a bloodstream infection in solid organ transplant recipients. Transplant Infectious Disease, 2021, 23, e13599.	1.7	5
30	Antibiotic Utilization in Deceased Organ Donors. Clinical Infectious Diseases, 2021, 73, 1284-1287.	5.8	4
31	Healthcare microenvironments define multidrug-resistant organism persistence. Infection Control and Hospital Epidemiology, 2021, , 1-7.	1.8	4
32	Severe acute respiratory coronavirus virus 2 (SARS-CoV-2) surface contamination in staff common areas and impact on healthcare worker infection: Prospective surveillance during the coronavirus disease 2019 (COVID-19) pandemic. Infection Control and Hospital Epidemiology, 2021, , 1-4.	1.8	4
33	Postdischarge antibiotic use for prophylaxis following spinal fusion. Infection Control and Hospital Epidemiology, 2020, 41, 789-798.	1.8	3
34	1961. A Randomized Controlled Trial of the Effect of Accelerated Copper Textiles on Healthcare-Associated Infections and Multidrug-Resistant Organisms: The "Investigating Microbial Pathogen Activity of Copper Textiles" (IMPACT) Study. Open Forum Infectious Diseases, 2018, 5, S568-S568.	0.9	2
35	Surgeon choice in the use of postdischarge antibiotics for prophylaxis following mastectomy with and without breast reconstruction. Infection Control and Hospital Epidemiology, 2021, 42, 467-470.	1.8	2
36	SARS-CoV-2 RNA persists on surfaces following terminal disinfection of COVID-19 hospital isolation rooms. American Journal of Infection Control, 2022, 50, 462-464.	2.3	2

#	Article	IF	Citations
37	1336A Randomized Controlled Trial of the Effect of Total Household Decolonization on Termination of Colonization with Methicillin-Resistant Staphylococcus aureus. Open Forum Infectious Diseases, 2014, 1, S57-S57.	0.9	1
38	Use of a Precision Antibiotic Therapy (PAT) Prediction Model to Identify Multidrug-Resistant (MDR) Enterobacteriaceae. Open Forum Infectious Diseases, 2017, 4, S328-S328.	0.9	1
39	1588. Clinical Prediction Tool for Extended-Spectrum β Lactamase-Producing Enterobacteriaceae as the Etiology of Bacteremia in Solid Organ Transplant Recipients. Open Forum Infectious Diseases, 2018, 5, S497-S497.	0.9	1
40	Respiratory Microbiome Disruption and Risk for Ventilator-Associated Lower Respiratory Tract Infection. Clinical Infectious Diseases, 2021, , .	5.8	1
41	Comparison of Respiratory Microbiome Disruption Indices to Predict VAP and VAE risk at LTACH Admission. Infection Control and Hospital Epidemiology, 2020, 41, s179-s180.	1.8	1
42	Evaluation of a research use only luminex based assay for measurement of procalcitonin in serum samples. American Journal of Translational Research (discontinued), 2016, 8, 4362-4369.	0.0	1
43	Enhanced Environmental Cleaning to Reduce Rates of Clostridioides difficile Infection on Oncology Units. Infection Control and Hospital Epidemiology, 2020, 41, s213-s213.	1.8	1
44	1813. Development and Validation of Novel Ambulatory Antibiotic Stewardship Metrics. Open Forum Infectious Diseases, 2018, 5, S514-S515.	0.9	0
45	2128. Predictors of Post-Discharge Prophylactic Antibiotics Following Spinal Fusion. Open Forum Infectious Diseases, 2018, 5, S626-S626.	0.9	0
46	Using nasal povidone-iodine to prevent bloodstream infections and transmission of Staphylococcus aureus among haemodialysis patients: a stepped-wedge cluster randomised control trial protocol. BMJ Open, 2021, 11, e048830.	1.9	0
47	Impact of Diagnosed and Undiagnosed Respiratory <i>Pseudomonas</i> on VAP and VAE During Long-Term Acute Care. Infection Control and Hospital Epidemiology, 2020, 41, s258-s259.	1.8	0