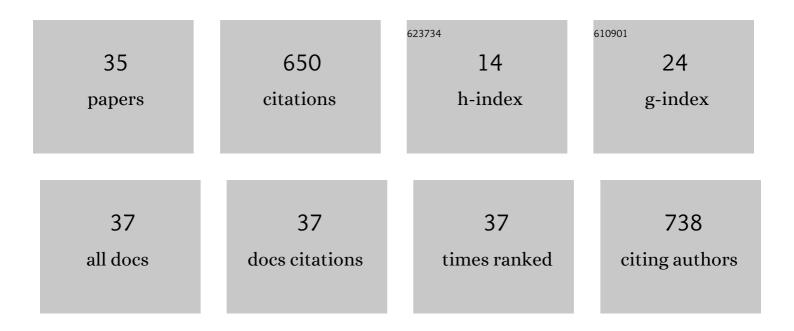
Kazumasa Fukuda

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Combined Radiographic Features and Age Can Distinguish Mycoplasma pneumoniae Pneumonia from Other Bacterial Pneumonias: Analysis Using the 16S rRNA Gene Sequencing Data. Journal of Clinical Medicine, 2022, 11, 2201.	2.4	2
2	A human respiratory tract-associated bacterium with an extremely small genome. Communications Biology, 2021, 4, 628.	4.4	3
3	Correlation between renal distribution of leptospires during the acute phase and chronic renal dysfunction in a hamster model of infection with Leptospira interrogans. PLoS Neglected Tropical Diseases, 2021, 15, e0009410.	3.0	5
4	Association between obligatory anaerobes and empyema caused by Streptococcus anginosus group bacteria. Respiratory Investigation, 2021, 59, 686-690.	1.8	4
5	Refinement of microbiota analysis of specimens from patients with respiratory infections using next-generation sequencing. Scientific Reports, 2021, 11, 19534.	3.3	9
6	Poor oral hygiene is associated with the detection of obligate anaerobes in pneumonia. Journal of Periodontology, 2020, 91, 65-73.	3.4	27
7	Clinical characteristics of patients with bacterial pleuritis in the presence of Streptococcus anginosus group and obligate anaerobes detected by clone library analysis. Clinical Respiratory Journal, 2020, 14, 267-276.	1.6	4
8	Pulmonary <i>Mycobacterium parascrofulaceum</i> Infection in a Patient with Chronic Progressive Pulmonary Aspergillosis: A Case Report and Literature Review. Internal Medicine, 2020, 59, 1417-1422.	0.7	3
9	The optimal duration of antimicrobial therapy for lower respiratory tract infection in patients with neuromuscular disorders based on a clone library analysis of the bacterial 16S rRNA gene sequence. International Journal of Infectious Diseases, 2020, 100, 396-401.	3.3	4
10	Bacterial composition of nasal discharge in children based on highly accurate 16S rRNA gene sequencing analysis. Scientific Reports, 2020, 10, 20193.	3.3	11
11	Acquisition of genetic mutations in Group A Streptococci at infection site and subsequent systemic dissemination of the mutants with lethal mutations in a streptococcal toxic shock syndrome mouse model. Microbial Pathogenesis, 2020, 143, 104116.	2.9	3
12	Changes of intravaginal microbiota and inflammation after selfâ€replacement ring pessary therapy compared to continuous ring pessary usage for pelvic organ prolapse. Journal of Obstetrics and Gynaecology Research, 2020, 46, 931-938.	1.3	6
13	Coinfection With Multiple Nontuberculous Mycobacteria as a Possible Exacerbating Factor in Pulmonary Nontuberculous Mycobacteriosis. Chest, 2020, 158, 2304-2313.	0.8	4
14	The microbiological characteristics of lower respiratory tract infection in patients with neuromuscular disorders: An investigation based on a multiplex polymerase chain reaction to detect viruses and a clone library analysis of the bacterial 16S rRNA gene sequence in sputum samples. Journal of Microbiology, Immunology and Infection, 2019, 52, 827-830.	3.1	8
15	Dynamics of microbiota during mechanical ventilation in aspiration pneumonia. BMC Pulmonary Medicine, 2019, 19, 260.	2.0	16
16	Transbronchial Invasion and Proliferation of Leptospira interrogans in Lung without Inflammatory Cell Infiltration in a Hamster Model. Infection and Immunity, 2019, 87, .	2.2	9
17	An infant case with hydrocephalus as the initial manifestation of Mycoplasma hominis -associated meningitis. Journal of Infection and Chemotherapy, 2017, 23, 713-716.	1.7	3
18	Frequency of detection of Chlamydophila pneumoniae using bronchoalveolar lavage fluid in patients with community-onset pneumonia. Respiratory Investigation, 2017, 55, 357-364.	1.8	8

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19	The first fatal case of Corynebacterium ulcerans infection in Japan. JMM Case Reports, 2017, 4, e005106.	1.3	17
20	Determining the Possible Etiology of Hospital-Acquired Pneumonia Using a Clone Library Analysis in Japan. Tohoku Journal of Experimental Medicine, 2017, 242, 9-17.	1.2	16
21	Bacteriological incidence in pneumonia patients with pulmonary emphysema: a bacterial floral analysis using the 16S ribosomal RNA gene in bronchoalveolar lavage fluid. International Journal of COPD, 2017, Volume 12, 2111-2120.	2.3	16
22	Molecular Approaches to Studying Microbial Communities: Targeting the 16S Ribosomal RNA Gene. Journal of UOEH, 2016, 38, 223-232.	0.6	39
23	The first report: An analysis of bacterial flora of the first voided urine specimens of patients with male urethritis using the 16S ribosomal RNA gene-based clone library method. Microbial Pathogenesis, 2016, 95, 95-100.	2.9	12
24	Clinical impact of methicillin-resistant staphylococcus aureus on bacterial pneumonia: cultivation and 16S ribosomal RNA gene analysis of bronchoalveolar lavage fluid. BMC Infectious Diseases, 2016, 16, 155.	2.9	26
25	Implantable cardioverter defibrillator infection due to Mycobacterium mageritense. Journal of Infection and Chemotherapy, 2016, 22, 180-183.	1.7	14
26	The significance of oral streptococci in patients with pneumonia with risk factors for aspiration: the bacterial floral analysis of 16S ribosomal RNA gene using bronchoalveolar lavage fluid. BMC Pulmonary Medicine, 2016, 16, 79.	2.0	60
27	Effects of pelvic organ prolapse ring pessary therapy on intravaginal microbial flora. International Urogynecology Journal, 2016, 27, 219-227.	1.4	16
28	Possible role of anaerobes in the pathogenesis of nontuberculous mycobacterial infection. Respirology, 2015, 20, 758-765.	2.3	23
29	Bacteriological Assessment of Healthcare-Associated Pneumonia Using a Clone Library Analysis. PLoS ONE, 2015, 10, e0124697.	2.5	45
30	An Unclassified Microorganism: Novel Pathogen Candidate Lurking in Human Airways. PLoS ONE, 2014, 9, e103646.	2.5	5
31	Ubiquitous distribution of phosphatidylinositol phosphate synthase and archaetidylinositol phosphate synthase in Bacteria and Archaea, which contain inositol phospholipid. Biochemical and Biophysical Research Communications, 2014, 443, 86-90.	2.1	33
32	Significance of Anaerobes and Oral Bacteria in Community-Acquired Pneumonia. PLoS ONE, 2013, 8, e63103.	2.5	112
33	Extrapolation of Group Proximity from Member Relations Using Embedding and Distribution Mapping. IEICE Transactions on Information and Systems, 2012, E95-D, 804-811.	0.7	1
34	A Higher Significance of Anaerobes. Chest, 2011, 139, 600-608.	0.8	62
35	Influence of Menstruation on the Microbiota of Healthy Women's Labia Minora as Analyzed Using a 16S rRNA Gene-Based Clone Library Method. Japanese Journal of Infectious Diseases, 2011, 64, 76-80.	1.2	23