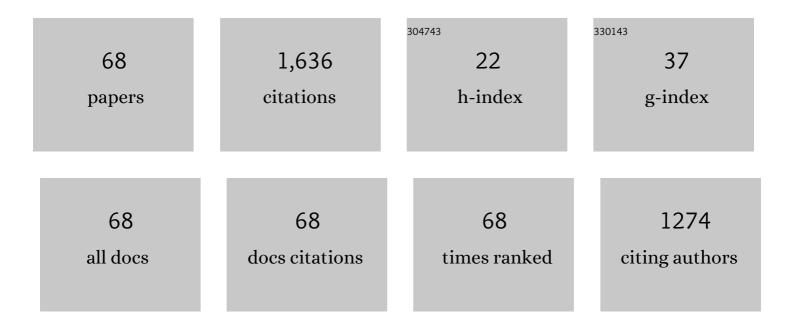
List of Publications by Year in descending order

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LOZSEE SOKI

#	Article	IF	CITATIONS
1	The prevalence of antibiotic resistance genes in Bacteroides fragilis group strains isolated in different European countries. Anaerobe, 2013, 21, 43-49.	2.1	123
2	Differentiation of division I (cfiA-negative) and division II (cfiA-positive) Bacteroides fragilis strains by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry. Journal of Medical Microbiology, 2011, 60, 1584-1590.	1.8	111
3	Community-Acquired Clostridium difficile Diarrhea Caused by Binary Toxin, Toxin A, and Toxin B Gene-Positive Isolates in Hungary. Journal of Clinical Microbiology, 2004, 42, 4316-4318.	3.9	85
4	Molecular characterization of imipenem-resistant, cfiA-positive Bacteroides fragilis isolates from the USA, Hungary and Kuwait. Journal of Medical Microbiology, 2004, 53, 413-419.	1.8	77
5	Molecular investigation of genetic elements contributing to metronidazole resistance in Bacteroides strains. Journal of Antimicrobial Chemotherapy, 2006, 57, 212-220.	3.0	75
6	Investigation of the prevalence of tetQ, tetX and tetX1 genes in Bacteroides strains with elevated tigecycline minimum inhibitory concentrations. International Journal of Antimicrobial Agents, 2011, 38, 522-525.	2.5	75
7	Multidrug-resistant Bacteroides fragilis group on the rise in Europe?. Journal of Medical Microbiology, 2012, 61, 1784-1788.	1.8	68
8	Comparison of a Rapid Molecular Method, the BD GeneOhm Cdiff Assay, to the Most Frequently Used Laboratory Tests for Detection of Toxin-Producing <i>Clostridium difficile</i> in Diarrheal Feces. Journal of Clinical Microbiology, 2009, 47, 3478-3481.	3.9	55
9	Examination of cfiA-mediated carbapenem resistance in Bacteroides fragilis strains from a European antibiotic susceptibility survey. International Journal of Antimicrobial Agents, 2006, 28, 497-502.	2.5	53
10	Molecular analysis of the carbapenem and metronidazole resistance mechanisms of Bacteroides strains reported in a Europe-wide antibiotic resistance survey. International Journal of Antimicrobial Agents, 2013, 41, 122-125.	2.5	52
11	Emergence and evolution of an international cluster of MDR <i>Bacteroides fragilis</i> isolates. Journal of Antimicrobial Chemotherapy, 2016, 71, 2441-2448.	3.0	47
12	Identification of antimicrobial resistance genes in multidrug-resistant clinical Bacteroides fragilis isolates by whole genome shotgun sequencing. Anaerobe, 2015, 31, 59-64.	2.1	42
13	First Hungarian case of an infection caused by multidrug-resistant Bacteroides fragilis strain. Anaerobe, 2015, 31, 55-58.	2.1	39
14	Instant screening and verification of carbapenemase activity in Bacteroides fragilis in positive blood culture, using matrix-assisted laser desorption ionization–time of flight mass spectrometry. Journal of Medical Microbiology, 2014, 63, 1105-1110.	1.8	37
15	To resist and persist: Important factors in the pathogenesis of Bacteroides fragilis. Microbial Pathogenesis, 2020, 149, 104506.	2.9	36
16	Use of MALDI-TOF/MS for routine detection of cfiA gene-positive Bacteroides fragilis strains. International Journal of Antimicrobial Agents, 2014, 44, 474-475.	2.5	29
17	Susceptibility profiles and resistance genes for carbapenems (cfiA) and metronidazole (nim) among Bacteroides species in a Turkish University Hospital. Anaerobe, 2012, 18, 169-171.	2.1	28
18	Detection of carbapenemase activities of Bacteroides fragilis strains with matrix-assisted laser desorption ionization – Time of flight mass spectrometry (MALDI-TOF MS). Anaerobe, 2014, 26, 49-52.	2.1	28

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19	Molecular analysis of the effector mechanisms of cefoxitin resistance among Bacteroides strains. Journal of Antimicrobial Chemotherapy, 2011, 66, 2492-2500.	3.0	27
20	A Europe-wide assessment of antibiotic resistance rates in Bacteroides and Parabacteroides isolates from intestinal microbiota of healthy subjects. Anaerobe, 2020, 62, 102182.	2.1	26
21	Extended role for insertion sequence elements in the antibiotic resistance of <i>Bacteroides</i> . World Journal of Clinical Infectious Diseases, 2013, 3, 1.	0.2	26
22	The Place of Molecular Genetic Methods in the Diagnostics of Human Pathogenic Anaerobic Bacteria. Acta Microbiologica Et Immunologica Hungarica, 2006, 53, 183-194.	0.8	24
23	A study on Nim expression in Bacteroides fragilis. Microbiology (United Kingdom), 2014, 160, 616-622.	1.8	24
24	Prevalence of the carbapenemase gene (cfiA) among clinical and normal flora isolates of Bacteroides species in Hungary. Journal of Medical Microbiology, 2000, 49, 427-430.	1.8	22
25	Bacteroides fragilis: A whole MALDI-based workflow from identification to confirmation of carbapenemase production for routine laboratories. Anaerobe, 2018, 54, 246-253.	2.1	21
26	High prevalence of division II (cfiA positive) isolates among blood stream Bacteroides fragilis in Slovenia as determined by MALDI-TOF MS. Anaerobe, 2019, 58, 30-34.	2.1	20
27	Analysis of Romanian Bacteroides isolates for antibiotic resistance levels and the corresponding antibiotic resistance genes. Anaerobe, 2015, 31, 11-14.	2.1	19
28	Occurrence of metronidazole and imipenem resistance among bacteroides fragilis group clinical isolates in Hungary. Acta Biologica Hungarica, 2001, 52, 271-280.	0.7	18
29	Screening of isolates from faeces for carbapenem-resistant Bacteroides strains; existence of strains with novel types of resistance mechanisms. International Journal of Antimicrobial Agents, 2004, 24, 450-454.	2.5	18
30	Distribution of Clostridium difficile PCR ribotypes in regions of Hungary. Journal of Medical Microbiology, 2006, 55, 279-282.	1.8	18
31	A multicentre survey of the antibiotic susceptibility of clinical <i>Bacteroides</i> species from Hungary. Infectious Diseases, 2018, 50, 372-380.	2.8	18
32	Prevalence and Characterization of nim Genes of Bacteroides spp. Isolated in Hungary. Anaerobe, 2002, 8, 175-179.	2.1	17
33	Assessment of changes in the epidemiology of Clostridium difficile isolated from diarrheal patients in Hungary. Anaerobe, 2009, 15, 237-240.	2.1	17
34	Molecular characterisation of multidrug-resistant Bacteroides isolates from Hungarian clinical samples. Journal of Global Antimicrobial Resistance, 2018, 13, 65-69.	2.2	17
35	Chicken or the Egg: Microbial Alterations in Biopsy Samples of Patients with Oral Potentially Malignant Disorders. Pathology and Oncology Research, 2019, 25, 1023-1033.	1.9	14
36	PCR ribotyping of clinically important Clostridium difficile strains from Hungary. Journal of Medical Microbiology, 2001, 50, 1082-1086.	1.8	14

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37	Prevalence, nucleotide sequence and expression studies of two proteins of a 5.6kb, Class III, Bacteroides plasmid frequently found in clinical isolates from European countries. Plasmid, 2010, 63, 86-97.	1.4	13
38	Outbreak of septicaemic cases caused by Acinetobacter ursingii in a neonatal intensive care unit. International Journal of Medical Microbiology, 2010, 300, 338-340.	3.6	12
39	Investigation of the MICs of fidaxomicin and other antibiotics against Hungarian Clostridium difficile isolates. Anaerobe, 2015, 31, 47-49.	2.1	12
40	Rare extraintestinal infection caused by toxin-producing Clostridium difficile. Anaerobe, 2010, 16, 301-303.	2.1	11
41	A novel <i>Bacteroides</i> metallo-l²-lactamase (MBL) and its gene ( <i>crxA</i> ) in <i>Bacteroides xylanisolvens</i> revealed by genomic sequencing and functional analysis. Journal of Antimicrobial Chemotherapy, 2022, 77, 1553-1556.	3.0	11
42	Modulation of Iron Import and Metronidazole Resistance in Bacteroides fragilis Harboring a nimA Gene. Frontiers in Microbiology, 0, 13, .	3.5	11
43	Incidence, antibiotic resistance and clonal relations of MRSA strains isolated from a Romanian university hospital. Acta Microbiologica Et Immunologica Hungarica, 2008, 55, 1-13.	0.8	9
44	MALDI-TOF MS versus 16S rRNA sequencing: Minor discrepancy between tools in identification of Bacteroides isolates. Acta Microbiologica Et Immunologica Hungarica, 2017, 65, 173-181.	0.8	9
45	Correlation between antibiotic resistance and clinical outcome of anaerobic infections; mini-review. Anaerobe, 2021, 72, 102463.	2.1	9
46	Characterisation of a 5.5-kb cryptic plasmid present in different isolates of Bacteroides spp. originating from Hungary. Journal of Medical Microbiology, 1999, 48, 25-31.	1.8	8
47	Coincidence of bft and cfiA genes in a multi-resistant clinical isolate of Bacteroides fragilis. Journal of Medical Microbiology, 2007, 56, 1416-1418.	1.8	8
48	Lactate dehydrogenase activity in Bacteroides fragilis group strains with induced resistance to metronidazole. Journal of Global Antimicrobial Resistance, 2016, 5, 11-14.	2.2	8
49	Detection of enterotoxin and protease genes among Hungarian clinical Bacteroides fragilis isolates. Anaerobe, 2017, 48, 98-102.	2.1	8
50	Detection of beta-lactamase production in clinical Prevotella species by MALDI-TOF MS method. Anaerobe, 2020, 65, 102240.	2.1	8
51	Corynebacterium striatum—Cot Worse by a Pandemic?. Pathogens, 2022, 11, 685.	2.8	8
52	Occurrence and analysis of rare cfiA–bft doubly positive Bacteroides fragilis strains. Anaerobe, 2013, 23, 70-73.	2.1	7
53	Molecular characterization of metronidazole resistant Bacteroides strains from Kuwait. Anaerobe, 2021, 69, 102357.	2.1	7
54	Haemin deprivation renders <i>Bacteroides fragilis</i> hypersusceptible to metronidazole and cancels high-level metronidazole resistance. Journal of Antimicrobial Chemotherapy, 2022, 77, 1027-1031.	3.0	7

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55	Fungaemia caused byCandida pulcherrima. Medical Mycology, 2012, 50, 522-524.	0.7	6
56	Prevalence and antimicrobial susceptibility of enterotoxigenic extra-intestinal Bacteroides fragilis among 13-year collection of isolates in Kuwait. BMC Microbiology, 2020, 20, 14.	3.3	6
57	Phenotypic and Molecular Characterization of Carbapenem-Heteroresistant Bacteroides fragilis Strains. Antibiotics, 2022, 11, 590.	3.7	6
58	An update on ampicillin resistance and β-lactamase genes of Bacteroides spp Journal of Medical Microbiology, 2021, 70, .	1.8	5
59	Isolation and characterization of an imipenem-resistant Bacteroides fragilis strain from a prostate abscess in a dog. Veterinary Microbiology, 2002, 84, 187-190.	1.9	4
60	A clinical isolate of Bacteroides fragilis from Hungary with high-level resistance to imipenem. Journal of Medical Microbiology, 2001, 50, 107-107.	1.8	4
61	A comparison of the antimicrobial resistance of fecal Bacteroides isolates and assessment of the composition of the intestinal microbiotas of carbapenem-treated and non-treated persons from Belgium and Hungary. Anaerobe, 2022, 73, 102480.	2.1	4
62	The microbial composition of the initial insult can predict the prognosis of experimental sepsis. Scientific Reports, 2021, 11, 22772.	3.3	4
63	The first characterized carbapenem-resistant Bacteroides fragilis strain from Croatia and the case study for it. Acta Microbiologica Et Immunologica Hungarica, 2018, 65, 317-323.	0.8	3
64	Characterization of the components of the thioredoxin system in Bacteroides fragilis and evaluation of its activity during oxidative stress. Anaerobe, 2022, 73, 102507.	2.1	3
65	Overexpression and purification of enzymatically active recombinant integrase protein of rous sarcoma virus. Virus Genes, 1992, 6, 301-306.	1.6	2
66	Molecular Relatedness of the Prevalent Cryptic Plasmids of Bacteroides Species Isolated in Hungary. Anaerobe, 2000, 6, 179-185.	2.1	2
67	A new insertion sequence element containing a cfiA gene in the first imipenem-resistant Bacteroides fragilis strain isolated in Italy. International Journal of Antimicrobial Agents, 2009, 34, 608-609.	2.5	1
68	P189 Bacteremia caused by Achromobacter xylosoxidans in a patient with haematological malignancy. Blood Reviews, 2007, 21, S145.	5.7	0