

Darina Cejkova

List of Publications by Citations

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37
papers

983
citations

20
h-index

31
g-index

38
ext. papers

1,341
ext. citations

4
avg, IF

3.85
L-index

#	Paper	IF	Citations
37	Succession and replacement of bacterial populations in the caecum of egg laying hens over their whole life. <i>PLoS ONE</i> , 2014 , 9, e115142	3.7	99
36	Whole genome sequences of three <i>Treponema pallidum</i> ssp. <i>pertenue</i> strains: yaws and syphilis treponemes differ in less than 0.2% of the genome sequence. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1471	4.8	86
35	Whole genome sequence of <i>Treponema pallidum</i> ssp. <i>pallidum</i> , strain Mexico A, suggests recombination between yaws and syphilis strains. <i>PLoS Neglected Tropical Diseases</i> , 2012 , 6, e1832	4.8	59
34	Whole genome sequencing and function prediction of 133 gut anaerobes isolated from chicken caecum in pure cultures. <i>BMC Genomics</i> , 2018 , 19, 561	4.5	54
33	Resequencing of <i>Treponema pallidum</i> ssp. <i>pallidum</i> strains Nichols and SS14: correction of sequencing errors resulted in increased separation of syphilis treponeme subclusters. <i>PLoS ONE</i> , 2013 , 8, e74319	3.7	53
32	Complete genome sequence of <i>Treponema paraluisuniculi</i> , strain Cuniculi A: the loss of infectivity to humans is associated with genome decay. <i>PLoS ONE</i> , 2011 , 6, e20415	3.7	51
31	Characterization of egg laying hen and broiler fecal microbiota in poultry farms in Croatia, Czech Republic, Hungary and Slovenia. <i>PLoS ONE</i> , 2014 , 9, e110076	3.7	50
30	Contact with adult hen affects development of caecal microbiota in newly hatched chicks. <i>PLoS ONE</i> , 2019 , 14, e0212446	3.7	49
29	Genome analysis of <i>Treponema pallidum</i> subsp. <i>pallidum</i> and subsp. <i>pertenue</i> strains: most of the genetic differences are localized in six regions. <i>PLoS ONE</i> , 2010 , 5, e15713	3.7	48
28	Whole genome sequence of the <i>Treponema</i> Fribourg-Blanc: unspecified simian isolate is highly similar to the yaws subspecies. <i>PLoS Neglected Tropical Diseases</i> , 2013 , 7, e2172	4.8	45
27	Whole genome sequence of the <i>Treponema pallidum</i> subsp. <i>endemicum</i> strain Bosnia A: the genome is related to yaws treponemes but contains few loci similar to syphilis treponemes. <i>PLoS Neglected Tropical Diseases</i> , 2014 , 8, e3261	4.8	39
26	Complete genome sequence of <i>Treponema pallidum</i> strain DAL-1. <i>Standards in Genomic Sciences</i> , 2012 , 7, 12-21		32
25	Effects of host genetics and environmental conditions on fecal microbiota composition of pigs. <i>PLoS ONE</i> , 2018 , 13, e0201901	3.7	28
24	Characterization of Antibiotic Resistance Gene Abundance and Microbiota Composition in Feces of Organic and Conventional Pigs from Four EU Countries. <i>PLoS ONE</i> , 2015 , 10, e0132892	3.7	28
23	Structure of <i>rrn</i> operons in pathogenic non-cultivable treponemes: sequence but not genomic position of intergenic spacers correlates with classification of <i>Treponema pallidum</i> and <i>Treponema paraluisuniculi</i> strains. <i>Journal of Medical Microbiology</i> , 2013 , 62, 196-207	3.2	25
22	Complete sequence of low-copy-number plasmid MccC7-H22 of probiotic <i>Escherichia coli</i> H22 and the prevalence of <i>mcc</i> genes among human <i>E. coli</i> . <i>Plasmid</i> , 2008 , 59, 1-10	3.3	21
21	Microcin determinants are associated with B2 phylogroup of human fecal <i>Escherichia coli</i> isolates. <i>MicrobiologyOpen</i> , 2016 , 5, 490-8	3.4	21

20	Curcuma and Scutellaria plant extracts protect chickens against inflammation and Salmonella Enteritidis infection. <i>Poultry Science</i> , 2015 , 94, 2049-58	3.9	20
19	Complete genome sequences of two strains of <i>Treponema pallidum</i> subsp. <i>pertenue</i> from Ghana, Africa: Identical genome sequences in samples isolated more than 7 years apart. <i>PLoS Neglected Tropical Diseases</i> , 2017 , 11, e0005894	4.8	20
18	Use of 16S rRNA gene sequencing for prediction of new opportunistic pathogens in chicken ileal and cecal microbiota. <i>Poultry Science</i> , 2019 , 98, 2347-2353	3.9	20
17	Genomic analysis of <i>Escherichia coli</i> strains isolated from diseased chicken in the Czech Republic. <i>BMC Veterinary Research</i> , 2020 , 16, 189	2.7	16
16	Antibiotic Resistance, Core-Genome and Protein Expression in IncHI1 Plasmids in <i>Salmonella</i> Typhimurium. <i>Genome Biology and Evolution</i> , 2016 , 8, 1661-71	3.9	16
15	Directly Sequenced Genomes of Contemporary Strains of Syphilis Reveal Recombination-Driven Diversity in Genes Encoding Predicted Surface-Exposed Antigens. <i>Frontiers in Microbiology</i> , 2019 , 10, 1691	5.7	15
14	A Retrospective Study on Genetic Heterogeneity within <i>Treponema</i> Strains: Subpopulations Are Genetically Distinct in a Limited Number of Positions. <i>PLoS Neglected Tropical Diseases</i> , 2015 , 9, e0004110	4.8	15
13	Gut Anaerobes Capable of Chicken Caecum Colonisation. <i>Microorganisms</i> , 2019 , 7,	4.9	15
12	Complete genome sequences of two strains of <i>Treponema pallidum</i> subsp. <i>pertenue</i> from Indonesia: Modular structure of several treponemal genes. <i>PLoS Neglected Tropical Diseases</i> , 2018 , 12, e0006867	4.8	12
11	Human <i>Escherichia coli</i> strains of different geographical and time source: bacteriocin types and their gene sequences are population-specific. <i>Environmental Microbiology Reports</i> , 2012 , 4, 459-66	3.7	10
10	Sequencing of <i>Treponema pallidum</i> subsp. <i>pallidum</i> from isolate UZ1974 using Anti-Treponemal Antibodies Enrichment: First complete whole genome sequence obtained directly from human clinical material. <i>PLoS ONE</i> , 2018 , 13, e0202619	3.7	9
9	Two highly divergent lineages of exfoliative toxin B-encoding plasmids revealed in impetigo strains of <i>Staphylococcus aureus</i> . <i>International Journal of Medical Microbiology</i> , 2017 , 307, 291-296	3.7	8
8	Clinical Characteristics of Patients with Tick-Borne Encephalitis (TBE): A European Multicentre Study from 2010 to 2017. <i>Microorganisms</i> , 2021 , 9,	4.9	6
7	Whole genome sequence of the <i>Treponema pallidum</i> subsp. <i>endemicum</i> strain Iraq B: A subpopulation of bejel treponemes contains full-length <i>tprF</i> and <i>tprG</i> genes similar to those present in <i>T. p.</i> subsp. <i>pertenue</i> strains. <i>PLoS ONE</i> , 2020 , 15, e0230926	3.7	3
6	The distribution of antibiotic resistance genes in chicken gut microbiota commensals. <i>Scientific Reports</i> , 2021 , 11, 3290	4.9	3
5	Characterization of blaKPC-3-positive plasmids from an <i>Enterobacter aerogenes</i> isolated from a corvid in Canada. <i>Journal of Antimicrobial Chemotherapy</i> , 2018 , 73, 2573-2575	5.1	2
4	Whole Genome Analyses of Treponemes: New Targets for Strain- and Subspecies-Specific Molecular Diagnostics 2011 ,		2
3	Genome sequences of two Antarctic strains of <i>Pseudomonas prosekii</i> : insights into adaptation to extreme conditions. <i>Archives of Microbiology</i> , 2020 , 202, 447-454	3	2

2	Carbapenemase-Producing Gram-Negative Bacteria from American Crows in the United States. <i>Antimicrobial Agents and Chemotherapy</i> , 2020 , 65,	5.9	1
1	Pyocin-mediated antagonistic interactions in <i>Pseudomonas</i> spp. isolated in James Ross Island, Antarctica. <i>Environmental Microbiology</i> , 2021 ,	5.2	0