Miklos Gyuraneczz

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/5123523/publications.pdf

Version: 2024-02-01

106	2,222	218592	289141
papers	citations	26 h-index	40 g-index

107	107	107	2461
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Tularaemia: clinical aspects in Europe. Lancet Infectious Diseases, The, 2016, 16, 113-124.	4.6	187
2	Worldwide Phylogenetic Relationship of Avian Poxviruses. Journal of Virology, 2013, 87, 4938-4951.	1.5	112
3	Molecular detection of vector-borne bacteria in bat ticks (Acari: Ixodidae, Argasidae) from eight countries of the Old and New Worlds. Parasites and Vectors, 2019, 12, 50.	1.0	91
4	Phylogeography ofFrancisella tularensissubsp.holarctica, Europe. Emerging Infectious Diseases, 2012, 18, 290-293.	2.0	82
5	Melt Analysis of Mismatch Amplification Mutation Assays (Melt-MAMA): A Functional Study of a Cost-Effective SNP Genotyping Assay in Bacterial Models. PLoS ONE, 2012, 7, e32866.	1.1	73
6	Investigation of the Ecology ofFrancisella tularensisDuring an Inter-Epizootic Period. Vector-Borne and Zoonotic Diseases, 2011, 11, 1031-1035.	0.6	66
7	First detection of bartonellae in a broad range of bat ectoparasites. Veterinary Microbiology, 2012, 159, 541-543.	0.8	52
8	Mutations Associated with Decreased Susceptibility to Seven Antimicrobial Families in Field and Laboratory-Derived Mycoplasma bovis Strains. Antimicrobial Agents and Chemotherapy, 2017, 61, .	1.4	48
9	Synanthropic Birds Associated with High Prevalence of Tick-Borne Rickettsiae and with the First Detection ofRickettsia aeschlimanniiin Hungary. Vector-Borne and Zoonotic Diseases, 2013, 13, 77-83.	0.6	46
10	Detection of Brucella Canisâ€“Induced Reproductive Diseases in a Kennel. Journal of Veterinary Diagnostic Investigation, 2011, 23, 143-147.	0.5	45
11	First isolation and characterization of Brucella microti from wild boar. BMC Veterinary Research, 2015, 11, 147.	0.7	43
12	Phylogeography of Francisella tularensis subspecies holarctica from the country of Georgia. BMC Microbiology, 2011, 11, 139.	1.3	42
13	Antibiotic susceptibility profiles of Mycoplasma synoviae strains originating from Central and Eastern Europe. BMC Veterinary Research, 2017, 13, 342.	0.7	42
14	Tularemia of European Brown Hare (Lepus europaeus). Veterinary Pathology, 2010, 47, 958-963.	0.8	41
15	Screening of bat faeces for arthropod-borne apicomplexan protozoa: Babesia canis and Besnoitia besnoiti-like sequences from Chiroptera. Parasites and Vectors, 2015, 8, 441.	1.0	40
16	Prevalence ofCoxiella burnetiiin Hungary: Screening of Dairy Cows, Sheep, Commercial Milk Samples, and Ticks. Vector-Borne and Zoonotic Diseases, 2012, 12, 650-653.	0.6	39
17	Unique genomic organization of a novel Avipoxvirus detected in turkey (Meleagris gallopavo). Infection, Genetics and Evolution, 2015, 35, 221-229.	1.0	39
18	Genotyping of Coxiella burnetiifrom domestic ruminants and human in Hungary: indication of various genotypes. BMC Veterinary Research, 2014, 10, 107.	0.7	34

#	Article	IF	Citations
19	Factors Influencing Emergence of Tularemia, Hungary, 1984â€“2010. Emerging Infectious Diseases, 2012, 18, 1379-1381.	2.0	32
20	Antibiotic susceptibility profiles of Mycoplasma bovis strains isolated from cattle in Hungary, Central Europe. BMC Veterinary Research, 2014, 10, 256.	0.7	31
21	A Serosurvey of Flavivirus Infection in Horses and Birds in Slovakia. Vector-Borne and Zoonotic Diseases, 2018, 18, 206-213.	0.6	30
22	Prevalence ofFrancisella tularensisandFrancisella-Like Endosymbionts in the Tick Population of Hungary and the Genetic Variability ofFrancisella-Like Agents. Vector-Borne and Zoonotic Diseases, 2013, 13, 160-163.	0.6	29
23	Antimicrobial susceptibility of Francisella tularensis subsp. holarctica strains from Hungary, Central Europe. Journal of Antimicrobial Chemotherapy, 2013, 68, 370-373.	1.3	29
24	A Field Study of Plague and Tularemia in Rodents, Western Iran. Vector-Borne and Zoonotic Diseases, 2017, 17, 247-253.	0.6	29
25	Genetic relatedness of Brucella suis biovar 2 isolates from hares, wild boars and domestic pigs. Veterinary Microbiology, 2014, 172, 492-498.	0.8	28
26	Q fever epidemic in Hungary, April to July 2013. Eurosurveillance, 2014, 19, .	3.9	28
27	Detection ofBorrelia burgdorferiSensu Lato andAnaplasma phagocytophilumin Small Mammals and Ectoparasites in Hungary. Vector-Borne and Zoonotic Diseases, 2011, 11, 1499-1501.	0.6	26
28	Detection of Francisella-like endosymbiont in Hyalomma rufipes from Ethiopia. Ticks and Tick-borne Diseases, 2014, 5, 818-820.	1.1	26
29	Mycoplasma anserisalpingitidis sp. nov., isolated from European domestic geese (Anser anser) Tj ETQq1 1 0.784314 rgBT /Overlock 1 Microbiology, 2020, 70, 2369-2381.	0.8	26
30	Non-pet dogs as sentinels and potential synanthropic reservoirs of tick-borne and zoonotic bacteria. Veterinary Microbiology, 2013, 167, 700-703.	0.8	25
31	Phylogeny of Mycoplasma bovisisolates from Hungary based on multi locus sequence typing and multiple-locus variable-number tandem repeat analysis. BMC Veterinary Research, 2014, 10, 108.	0.7	25
32	Influence of the Biotope on the Tick Infestation of Cattle and on the Tick-Borne Pathogen Repertoire of Cattle Ticks in Ethiopia. PLoS ONE, 2014, 9, e106452.	1.1	24
33	Tularemia and plague survey in rodents in an earthquake zone in southeastern Iran. Epidemiology and Health, 2015, 37, e2015050.	0.8	22
34	Characterization of Francisella tularensis Strains, Comparing Their Carbon Source Utilization. Zoonoses and Public Health, 2010, 57, 417-422.	0.9	20
35	Antibiotic susceptibility profiles of Mycoplasma sp. 1220 strains isolated from geese in Hungary. BMC Veterinary Research, 2016, 12, 170.	0.7	20
36	Combination therapy of rabies-infected mice with inhibitors of pro-inflammatory host response, antiviral compounds and human rabies immunoglobulin. Vaccine, 2019, 37, 4724-4735.	1.7	20

#	Article	IF	Citations
37	New antimicrobial susceptibility data from monitoring of Mycoplasma bovis isolated in Europe. Veterinary Microbiology, 2019, 238, 108432.	0.8	20
38	Genotyping Mycoplasma gallisepticum by multilocus sequence typing. Veterinary Microbiology, 2019, 231, 191-196.	0.8	20
39	Comparison of virulence of Francisella tularensis ssp. holarctica genotypes B.12 and B.FTNF002-00. BMC Veterinary Research, 2016, 13, 46.	0.7	19
40	Antibiotic susceptibility testing of Mycoplasma hyopneumoniae field isolates from Central Europe for fifteen antibiotics by microbroth dilution method. PLoS ONE, 2018, 13, e0209030.	1.1	19
41	Brucellosis of the European Brown Hare (Lepus europaeus). Journal of Comparative Pathology, 2011, 145, 1-5.	0.1	18
42	Francisella tularensissubsp.tularensisGroup A.I, United States. Emerging Infectious Diseases, 2014, 20, 861-5.	2.0	18
43	Assessing bat droppings and predatory bird pellets for vector-borne bacteria: molecular evidence of bat-associated Neorickettsia sp. in Europe. Antonie Van Leeuwenhoek, 2018, 111, 1707-1717.	0.7	18
44	Antibiotic susceptibility profiles of Mycoplasma hyorhinis strains isolated from swine in Hungary. Veterinary Microbiology, 2019, 228, 196-201.	0.8	18
45	Occurrence of Coxiella burnetii and Chlamydiales species in abortions of domestic ruminants and in wild ruminants in Hungary, Central Europe. Journal of Veterinary Diagnostic Investigation, 2015, 27, 206-210.	0.5	17
46	Serologic Evidence of Crimean-Congo Hemorrhagic Fever Virus Infection in Hungary. Vector-Borne and Zoonotic Diseases, 2013, 13, 270-272.	0.6	16
47	Isolation of Mycoplasma anserisalpingitidis from swan goose (Anser cygnoides) in China. BMC Veterinary Research, 2020, 16, 178.	0.7	16
48	Susceptibility of the Common Hamster (Cricetus cricetus) to Francisella tularensis and Its Effect on the Epizootiology of Tularemia in an Area Where Both Are Endemic. Journal of Wildlife Diseases, 2010, 46, 1316-1320.	0.3	15
49	Identification of tick-borne encephalitis virus in ticks collected in southeastern Hungary. Ticks and Tick-borne Diseases, 2013, 4, 427-431.	1.1	15
50	Within-host evolution of Brucella canis during a canine brucellosis outbreak in a kennel. BMC Veterinary Research, 2013, 9, 76.	0.7	15
51	Genotyping of Brucella melitensis strains from dromedary camels (Camelus dromedarius) from the United Arab Emirates with multiple-locus variable-number tandem repeat analysis. Veterinary Microbiology, 2016, 186, 8-12.	0.8	15
52	Genotyping Mycoplasma hyopneumoniae isolates based on multi-locus sequence typing, multiple-locus variable-number tandem repeat analysis and analysing gene p146. Veterinary Microbiology, 2018, 222, 85-90.	0.8	15
53	Rapid, Simple and Cost-Effective Molecular Method to Differentiate the Temperature Sensitive (ts+) MS-H Vaccine Strain and Wild-Type Mycoplasma synoviae Isolates. PLoS ONE, 2015, 10, e0133554.	1.1	15
54	First isolation of Histophilus somni from goats. Veterinary Microbiology, 2009, 133, 383-386.	0.8	14

#	Article	IF	Citations
55	Minimal inhibitory concentration of seven antimicrobials to Mycoplasma gallisepticum and Mycoplasma synoviae isolates from six European countries. Avian Pathology, 2021, 50, 161-173.	0.8	14
56	Molecular analysis and MIRU-VNTR typing of Mycobacterium avium subsp. paratuberculosis strains from various sources. Journal of Applied Microbiology, 2015, 118, 275-283.	1.4	12
57	Deciphering the protein interaction in adhesion of Francisella tularensis subsp. holarctica to the endothelial cells. Microbial Pathogenesis, 2015, 81, 6-15.	1.3	12
58	Development of Molecular Methods for Rapid Differentiation of Mycoplasma gallisepticum Vaccine Strains from Field Isolates. Journal of Clinical Microbiology, 2019, 57, .	1.8	12
59	Molecular Differentiation of Mycoplasma gallisepticum Outbreaks: A Last Decade Study on Italian Farms Using GTS and MLST. Vaccines, 2020, 8, 665.	2.1	12
60	Molecular analysis and MIRU-VNTR typing of Mycobacterium avium subsp. avium, â€˜hominissuisâ€™ and silvaticum strains of veterinary origin. Infection, Genetics and Evolution, 2016, 40, 192-199.	1.0	11
61	Babesia genotypes in Haemaphysalis concinna collected from birds in Hungary reflect phylogeographic connections with Siberia and the Far East. Ticks and Tick-borne Diseases, 2017, 8, 666-670.	1.1	11
62	Development of molecular methods for the rapid detection of antibiotic susceptibility of Mycoplasma bovis. Veterinary Microbiology, 2018, 213, 47-57.	0.8	11
63	Detection of Mycoplasma anatis, M. anseris, M. cloacale and Mycoplasma sp. 1220 in waterfowl using species-specific PCR assays. PLoS ONE, 2019, 14, e0219071.	1.1	11
64	Evaluation of in vitro inhibitory potential of type-I interferons and different antiviral compounds on rabies virus replication. Vaccine, 2019, 37, 4663-4672.	1.7	11
65	The core genome multi-locus sequence typing of Mycoplasma anserisalpingitidis. BMC Genomics, 2020, 21, 403.	1.2	11
66	Identification of Novel Coxiella burnetii Genotypes from Ethiopian Ticks. PLoS ONE, 2014, 9, e113213.	1.1	11
67	Development of mismatch amplification mutation assays for the differentiation of MS1 vaccine strain from wild-type Mycoplasma synoviae and MS-H vaccine strains. PLoS ONE, 2017, 12, e0175969.	1.1	11
68	Generalized Tularemia in a Vervet Monkey (Chlorocebus Aethiops) and a Patas Monkey (Erythrocebus Patas) in a Zoo. Journal of Veterinary Diagnostic Investigation, 2009, 21, 384-387.	0.5	9
69	Analyses of separate and concatenated cox1 and 18S rRNA gene sequences indicate that the bat piroplasm Babesia vesperuginis is phylogenetically close to Cytauxzoon felis and the â€˜prototheileridâ€™ Babesia conradae. Acta Veterinaria Hungarica, 2018, 66, 107-115.	0.2	9
70	Antimicrobial susceptibility of pathogenic mycoplasmas in chickens in Asia. Veterinary Microbiology, 2020, 250, 108840.	0.8	9
71	Molecular screening for Anaplasmataceae in ticks and tsetse flies from Ethiopia. Acta Veterinaria Hungarica, 2016, 64, 65-70.	0.2	8
72	Impact of a freeway on the dispersal of ticks and Ixodes ricinus-borne pathogens: forested resting areas may become Lyme disease hotspots. Acta Veterinaria Hungarica, 2017, 65, 242-252.	0.2	8

#	Article	IF	Citations
73	Genotyping Mycoplasma synoviae: Development of a multi-locus variable number of tandem-repeats analysis and comparison with current molecular typing methods. Veterinary Microbiology, 2018, 226, 41-49.	0.8	8
74	Prevalence of Coxiella burnetii in Central and Eastern European dairy herds. Comparative Immunology, Microbiology and Infectious Diseases, 2020, 72, 101489.	0.7	8
75	Aerosol infection of calves with Histophilus somni. Acta Veterinaria Hungarica, 2009, 57, 347-356.	0.2	7
76	A serological and molecular study on Francisella tularensis in rodents from Hamadan province, Western Iran. Comparative Immunology, Microbiology and Infectious Diseases, 2020, 68, 101379.	0.7	7
77	Genotyping Mycoplasma hyorhinis by multi-locus sequence typing and multiple-locus variable-number tandem-repeat analysis. Veterinary Microbiology, 2020, 249, 108836.	0.8	7
78	Anaplasmataceae closely related to Ehrlichia chaffeensis and Neorickettsia helminthoeca from birds in Central Europe, Hungary. Antonie Van Leeuwenhoek, 2020, 113, 1067-1073.	0.7	7
79	Decrease ofMycoplasma gallisepticumseroprevalence and introduction of new genotypes in Dutch commercial poultry during the years 2001â€“2018. Avian Pathology, 2021, 50, 52-60.	0.8	7
80	Seroprevalence of Francisella tularensis in Austrian Hunting Dogs. Vector-Borne and Zoonotic Diseases, 2018, 18, 117-119.	0.6	6
81	Mutations potentially associated with decreased susceptibility to fluoroquinolones, macrolides and lincomycin in Mycoplasma synoviae. Veterinary Microbiology, 2020, 248, 108818.	0.8	6
82	Multilocus sequence typing of the goose pathogen Mycoplasma anserisalpingitidis. Veterinary Microbiology, 2021, 254, 108972.	0.8	6
83	Complete Genome Sequences of Three Mycoplasma anserisalpingitis (Mycoplasma sp. 1220) Strains. Microbiology Resource Announcements, 2019, 8, .	0.3	6
84	Development of molecular biological tools for the rapid determination of antibiotic susceptibility of Mycoplasma hyopneumoniae isolates. Veterinary Microbiology, 2020, 245, 108697.	0.8	5
85	Diversity of tick species and associated pathogens on peri-urban wild boars â€“ First report of the zoonotic Babesia cf. crassa from Hungary. Ticks and Tick-borne Diseases, 2022, 13, 101936.	1.1	5
86	Antimicrobial susceptibility of Bacillus anthracis strains from Hungary. Acta Veterinaria Hungarica, 2016, 64, 141-147.	0.2	4
87	Complete Genome Sequences of Mycoplasma anatis, M. anseris, and M. cloacale Type Strains. Microbiology Resource Announcements, 2018, 7, .	0.3	4
88	Serological screening for Coxiella burnetii in the context of early pregnancy loss in dairy cows. Acta Veterinaria Hungarica, 2020, 68, 305-309.	0.2	4
89	Natural IS711 insertion causing omp31 gene inactivation in Brucella ovis. Journal of Veterinary Diagnostic Investigation, 2013, 25, 234-238.	0.5	3
90	Development of a molecular biological assay for the detection of markers related to decreased susceptibility to macrolides and lincomycin in Mycoplasma hyorhinis. Acta Veterinaria Hungarica, 2021, 69, 110-115.	0.2	3

#	Article	IF	Citations
91	Brucella melitensis caused abortion in a serologically positive dromedary camel. Journal of Camel Practice and Research, 2019, 26, 1.	0.0	3
92	Identification and detection of mutations potentially associated with decreased susceptibility to macrolides and lincomycin in Mycoplasma anserisalpingitidis isolates. Veterinary Microbiology, 2022, 266, 109362.	0.8	3
93	Complement sensitivity and factor H binding of European Francisella tularensis ssp. holarctica strains in selected animal species. Acta Veterinaria Hungarica, 2015, 63, 275-284.	0.2	2
94	Screening of Hungarian cattle herds for seropositivity to Mycoplasma bovis. Acta Veterinaria Hungarica, 2017, 65, 166-172.	0.2	2
95	Antimicrobial susceptibility monitoring of Mycoplasma hyopneumoniae isolated from seven European countries during 2015â€“2016. Veterinary Microbiology, 2021, 253, 108973.	0.8	2
96	Serological survey of Coxiella burnetii infections in dairy cattle, sheep, goats and zoo animals in Hungary â€“ Short communication. Acta Veterinaria Hungarica, 2021, 69, 105-109.	0.2	2
97	The distribution of lagomorph syphilis caused by Treponema paraluisleporidarum in Europe. European Journal of Wildlife Research, 2021, 67, 1.	0.7	2
98	West Nile virus - a new infection in the Slovak Republic?. Central European Journal of Public Health, 2018, 26, S51-S55.	0.4	2
99	Evidence of Mycoplasma spp. transmission by migratory wild geese. Poultry Science, 2022, 101, 101526.	1.5	2
100	Development of molecular assays for the rapid and cost-effective determination of fluoroquinolone, macrolide and lincosamide susceptibility of Mycoplasma synoviae isolates. PLoS ONE, 2020, 15, e0241647.	1.1	2
101	Genetic Traces of the Francisella tularensis Colonization of Spain, 1998â€“2020. Microorganisms, 2020, 8, 1784.	1.6	1
102	Novel prophage-like sequences in Mycoplasma anserisalpingitidis. Infection, Genetics and Evolution, 2021, 92, 104886.	1.0	1
103	Laboratory Investigations after Eye Drop Immunisation of Dromedaries with Live Attenuated Brucellamelitensis Rev 1 Vaccine. Journal of Camel Practice and Research, 2017, 24, 9.	0.0	1
104	In vitro susceptibility of Mycoplasma iowae isolates to antimicrobial agents. Avian Pathology, 2022, 51, 374-380.	0.8	1
105	Development of mismatch amplification mutation assay for the rapid differentiation of Mycoplasma gallisepticum K vaccine strain from field isolates. Avian Pathology, 2020, 49, 317-324.	0.8	0
106	Mycoplasma species in the male reproductive organs and the fresh and frozen semen of the Hungarian native goose. Avian Pathology, 2021, 50, 1-19.	0.8	0