SÃ;ndor Hornok

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/215065/publications.pdf

Version: 2024-02-01

100 papers 2,660 citations

30 h-index 233421 45 g-index

102 all docs

 $\begin{array}{c} 102 \\ \\ \text{docs citations} \end{array}$

102 times ranked

2375 citing authors

#	Article	IF	CITATIONS
1	Birds as potential reservoirs of tick-borne pathogens: first evidence of bacteraemia with Rickettsia helvetica. Parasites and Vectors, 2014, 7, 128.	2.5	95
2	Molecular Investigations of <i>Rickettsia helvetica</i> Infection in Dogs, Foxes, Humans, and <i>Ixodes</i> Ticks. Applied and Environmental Microbiology, 2009, 75, 3230-3237.	3.1	93
3	Molecular investigation of hard ticks (Acari: Ixodidae) and fleas (Siphonaptera: Pulicidae) as potential vectors of rickettsial and mycoplasmal agents. Veterinary Microbiology, 2010, 140, 98-104.	1.9	92
4	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.	2.5	91
5	First Molecular Evidence of <i>Anaplasma ovis </i> and <i>Rickettsia </i> spp. in Keds (Diptera:) Tj ETQq1 1 0.784314	4 rgBT /O\ 	verlgck 10 Tf
6	First serological and molecular evidence on the endemicity of Anaplasma ovis and A. marginale in Hungary. Veterinary Microbiology, 2007, 122, 316-322.	1.9	81
7	Molecular identification of Anaplasma marginale and rickettsial endosymbionts in blood-sucking flies (Diptera: Tabanidae, Muscidae) and hard ticks (Acari: Ixodidae). Veterinary Parasitology, 2008, 154, 354-359.	1.8	77
8	Emerging horizons for tick-borne pathogens: from the â€~one pathogen–one disease' vision to the pathobiome paradigm. Future Microbiology, 2015, 10, 2033-2043.	2.0	67
9	First molecular evidence of Hepatozoon canis infection in red foxes and golden jackals from Hungary. Parasites and Vectors, 2014, 7, 303.	2.5	63
10	First report of adult Hyalomma marginatum rufipes (vector of Crimean-Congo haemorrhagic fever) Tj ETQq0 0 0 r	gBT_/Over	lock 10 Tf 50
11	DNA of Piroplasms of Ruminants and Dogs in Ixodid Bat Ticks. PLoS ONE, 2016, 11, e0167735.	2.5	56
12	High prevalence of Hepatozoon-infection among shepherd dogs in a region considered to be free of Rhipicephalus sanguineus. Veterinary Parasitology, 2013, 196, 189-193.	1.8	54
13	Bovine besnoitiosis emerging in Central-Eastern Europe, Hungary. Parasites and Vectors, 2014, 7, 20.	2.5	54
14	Eco-epidemiology of Novel Bartonella Genotypes from Parasitic Flies of Insectivorous Bats. Microbial Ecology, 2018, 76, 1076-1088.	2.8	50
15	Synanthropic Birds Associated with High Prevalence of Tick-Borne Rickettsiae and with the First Detection of <i>Rickettsia aeschlimannii </i> in Hungary. Vector-Borne and Zoonotic Diseases, 2013, 13, 77-83.	1.5	46
16	Tick- and fly-borne bacteria in ungulates: the prevalence of Anaplasma phagocytophilum, haemoplasmas and rickettsiae in water buffalo and deer species in Central Europe, Hungary. BMC Veterinary Research, 2018, 14, 98.	1.9	46
17	Molecular characterization of two different strains of haemotropic mycoplasmas from a sheep flock with fatal haemolytic anaemia and concomitant Anaplasma ovis infection. Veterinary Microbiology, 2009, 136, 372-377.	1.9	43
18	Occurrence of ticks and prevalence of Anaplasma phagocytophilum and Borrelia burgdorferi s.l. in three types of urban biotopes: Forests, parks and cemeteries. Ticks and Tick-borne Diseases, 2014, 5, 785-789.	2.7	42

#	Article	IF	CITATIONS
19	East and west separation of Rhipicephalus sanguineus mitochondrial lineages in the Mediterranean Basin. Parasites and Vectors, 2017, 10, 39.	2.5	42
20	Synanthropic rodents and their ectoparasites as carriers of a novel haemoplasma and vector-borne, zoonotic pathogens indoors. Parasites and Vectors, 2015, 8, 27.	2.5	41
21	Screening of bat faeces for arthropod-borne apicomplexan protozoa: Babesia canis and Besnoitia besnoiti-like sequences from Chiroptera. Parasites and Vectors, 2015, 8, 441.	2.5	40
22	Contributions to the phylogeny of Ixodes (Pholeoixodes) canisuga, I. (Ph.) kaiseri, I. (Ph.) hexagonus and a simple pictorial key for the identification of their females. Parasites and Vectors, 2017, 10, 545.	2.5	40
23	Seroprevalence of canine babesiosis in Hungary suggesting breed predisposition. Parasitology Research, 2006, 99, 638-642.	1.6	39
24	Allochronic seasonal peak activities of Dermacentor and Haemaphysalis spp. under continental climate in Hungary. Veterinary Parasitology, 2009, 163, 366-369.	1.8	39
25	Prevalence of i>Coxiella burnetii i>in Hungary: Screening of Dairy Cows, Sheep, Commercial Milk Samples, and Ticks. Vector-Borne and Zoonotic Diseases, 2012, 12, 650-653.	1.5	39
26	Bat ticks revisited: Ixodes ariadnae sp. nov. and allopatric genotypes of I. vespertilionis in caves of Hungary. Parasites and Vectors, 2014, 7, 202.	2.5	38
27	Phylogenies from mitochondrial genomes of 120 species of ticks: Insights into the evolution of the families of ticks and of the genus Amblyomma. Ticks and Tick-borne Diseases, 2021, 12, 101577.	2.7	38
28	Molecular evidence of Ehrlichia canis and Rickettsia massiliae in ixodid ticks of carnivores from South Hungary. Acta Veterinaria Hungarica, 2013, 61, 42-50.	0.5	37
29	First report on Babesia cf. microti infection of red foxes (Vulpes vulpes) from Hungary. Parasites and Vectors, 2015, 8, 55.	2.5	37
30	Sarcocystis-infection of cattle in Hungary. Parasites and Vectors, 2015, 8, 69.	2.5	36
31	Bats and ticks: host selection and seasonality of bat-specialist ticks in eastern Europe. Parasites and Vectors, 2019, 12, 605.	2.5	35
32	Molecular investigations of the bat tick Argas vespertilionis (Ixodida: Argasidae) and Babesia vesperuginis (Apicomplexa: Piroplasmida) reflect "bat connection―between Central Europe and Central Asia. Experimental and Applied Acarology, 2017, 72, 69-77.	1.6	33
33	Seroprevalence of Toxoplasma gondii and Neospora caninum infection of cats in Hungary. Acta Veterinaria Hungarica, 2008, 56, 81-88.	0.5	32
34	Babesia vesperuginis, a neglected piroplasmid: new host and geographical records, and phylogenetic relations. Parasites and Vectors, 2017, 10, 598.	2.5	31
35	Bartonella DNA in heart tissues of bats in central and eastern Europe and a review of phylogenetic relations of bat-associated bartonellae. Parasites and Vectors, 2018, 11, 489.	2.5	31
36	Re-emergence of bovine piroplasmosis in Hungary: has the etiological role of Babesia divergens been taken over by B. major and Theileria buffeli?. Parasites and Vectors, 2014, 7, 434.	2.5	30

#	Article	IF	Citations
37	High mitochondrial sequence divergence in synanthropic flea species (Insecta: Siphonaptera) from Europe and the Mediterranean. Parasites and Vectors, 2018, 11, 221.	2.5	30
38	Effect of Climate and Land Use on the Spatio-Temporal Variability of Tick-Borne Bacteria in Europe. International Journal of Environmental Research and Public Health, 2018, 15, 732.	2.6	29
39	Argasid Ticks of Palearctic Bats: Distribution, Host Selection, and Zoonotic Importance. Frontiers in Veterinary Science, 2021, 8, 684737.	2.2	27
40	Non-pet dogs as sentinels and potential synanthropic reservoirs of tick-borne and zoonotic bacteria. Veterinary Microbiology, 2013, 167, 700-703.	1.9	25
41	Contributions to the morphology and phylogeny of the newly discovered bat tick species, Ixodes ariadnae in comparison with I. vespertilionis and I. simplex. Parasites and Vectors, 2015, 8, 47.	2.5	25
42	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.	2.5	24
43	Mitochondrial gene heterogeneity of the bat soft tick Argas vespertilionis (Ixodida: Argasidae) in the Palaearctic. Parasites and Vectors, 2017, 10, 109.	2.5	24
44	High degree of mitochondrial gene heterogeneity in the bat tick species Ixodes vespertilionis, I. ariadnae and I. simplex from Eurasia. Parasites and Vectors, 2015, 8, 457.	2.5	23
45	Genetic diversity, piroplasms and trypanosomes in Rhipicephalus microplus and Hyalomma anatolicum collected from cattle in northern Pakistan. Experimental and Applied Acarology, 2019, 79, 233-243.	1.6	23
46	Phylogenetic analyses of bat-associated bugs (Hemiptera: Cimicidae: Cimicinae and Cacodminae) indicate two new species close to Cimex lectularius. Parasites and Vectors, 2017, 10, 439.	2.5	22
47	Rickettsia raoultii and Rickettsia sibirica in ticks from the long-tailed ground squirrel near the China–Kazakhstan border. Experimental and Applied Acarology, 2019, 77, 425-433.	1.6	22
48	Molecular analysis of Ixodes rugicollis, Candidatus Neoehrlichia sp. (FU98) and a novel Babesia genotype from a European badger (Meles meles). Ticks and Tick-borne Diseases, 2017, 8, 41-44.	2.7	21
49	Rickettsiae in the common pipistrelle Pipistrellus pipistrellus (Chiroptera: Vespertilionidae) and the bat soft tick Argas vespertilionis (Ixodida: Argasidae). Parasites and Vectors, 2020, 13, 10.	2.5	21
50	Vector-Borne Agents Detected in Fleas of the Northern White-Breasted Hedgehog. Vector-Borne and Zoonotic Diseases, 2014, 14, 74-76.	1.5	20
51	Checklist of the hard tick (Acari: Ixodidae) fauna of Hungary with emphasis on host-associations and the emergence of Rhipicephalus sanguineus. Experimental and Applied Acarology, 2020, 80, 311-328.	1.6	20
52	Uneven seasonal distribution of Babesia canis and its two 18S rDNA genotypes in questing Dermacentor reticulatus ticks in urban habitats. Ticks and Tick-borne Diseases, 2016, 7, 694-697.	2.7	19
53	Diversity of Haemaphysalis-associated piroplasms of ruminants in Central-Eastern Europe, Hungary. Parasites and Vectors, 2015, 8, 627.	2.5	18
54	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.	1.7	18

#	Article	IF	CITATIONS
55	<i>Babesia vesperuginis</i> in Common Pipistrelle (<i>Pipistrellus pipistrellus</i>) and the Bat Soft Tick <i>Argas vespertilionis</i> in the People's Republic of China. Journal of Wildlife Diseases, 2018, 54, 419-421.	0.8	18
56	Molecular identification of badger-associated Babesia sp. DNA in dogs: updated phylogeny of piroplasms infecting Caniformia. Parasites and Vectors, 2018, 11, 235.	2.5	17
57	Molecular phylogeny of Amblyomma exornatum and Amblyomma transversale, with reinstatement of the genus Africaniella (Acari: Ixodidae) for the latter. Ticks and Tick-borne Diseases, 2020, 11, 101494.	2.7	17
58	Vector-borne transmission of Besnoitia besnoiti by blood-sucking and secretophagous flies: epidemiological and clinicopathological implications. Parasites and Vectors, 2015, 8, 450.	2.5	16
59	Rickettsiae in red fox (Vulpes vulpes), marbled polecat (Vormela peregusna) and their ticks in northwestern China. Parasites and Vectors, 2021, 14, 204.	2.5	15
60	First record of Ixodes ariadnae in Germany â€" Short communication. Acta Veterinaria Hungarica, 2015, 63, 347-351.	0.5	14
61	Phylogenetic analysis of Haemaphysalis erinacei Pavesi, 1884 (Acari: Ixodidae) from China, Turkey, Italy and Romania. Parasites and Vectors, 2016, 9, 643.	2.5	14
62	Molecular evidence of a badger-associated Ehrlichia sp., a Candidatus Neoehrlichia lotoris-like genotype and Anaplasma marginale in dogs. Ticks and Tick-borne Diseases, 2018, 9, 1302-1309.	2.7	14
63	Evidence for hostÂspecificity of Theileria capreoli genotypes in cervids. Parasites and Vectors, 2017, 10, 473.	2.5	13
64	DNA of free-living bodonids (Euglenozoa: Kinetoplastea) in bat ectoparasites: potential relevance to the evolution of parasitic trypanosomatids. Acta Veterinaria Hungarica, 2017, 65, 531-540.	0.5	13
65	DNA of Theileria orientalis, T. equi and T. capreoli in stable flies (Stomoxys calcitrans). Parasites and Vectors, 2020, 13, 186.	2.5	13
66	An unexpected advantage of insectivorism: insect moulting hormones ingested by song birds affect their ticks. Scientific Reports, 2016, 6, 23390.	3.3	12
67	Description of a new tick species, Ixodes collaris n. sp. (Acari: Ixodidae), from bats (Chiroptera:) Tj ETQq1 1 0.78	4314 rgBT 2.5	 Overlock 10 12
68	On the way between Africa and Europe: Molecular taxonomy of ticks collected from birds in Malta. Ticks and Tick-borne Diseases, 2022, 13, 102001.	2.7	12
69	First molecular identification of â€~ Candidatus Mycoplasma haemominutum' from a cat with fatal haemolytic anaemia in Hungary. Acta Veterinaria Hungarica, 2008, 56, 441-450.	0.5	11
70	Birds as disseminators of ixodid ticks and tick-borne pathogens: note on the relevance to migratory routes. Ornis Hungarica, 2012, 20, 86-89.	0.4	11
71	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.	2.7	11
72	Wide Distribution and Diversity of Malaria-Related Haemosporidian Parasites (Polychromophilus spp.) in Bats and Their Ectoparasites in Eastern Europe. Microorganisms, 2021, 9, 230.	3.6	11

#	Article	IF	CITATIONS
73	Infestation of Rhipicephalus sanguineus sensu lato on cats in Malta. Ticks and Tick-borne Diseases, 2018, 9, 1120-1124.	2.7	9
74	First record of Ixodes simplex found on a human host, with a review of cases of human infestation by bat tick species occurring in Europe. Ticks and Tick-borne Diseases, 2021, 12, 101722.	2.7	9
75	Molecular epidemiological study on ticks and tick-borne protozoan parasites (Apicomplexa:) Tj ETQq1 1 0.784 (Sciurus vulgaris) in central Europe, Hungary. Parasites and Vectors, 2022, 15, .	314 rgBT /C 2.5	Overlock 10 Tf 9
76	Molecular screening for Anaplasmataceae in ticks and tsetse flies from Ethiopia. Acta Veterinaria Hungarica, 2016, 64, 65-70.	0.5	8
77	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.	0.5	8
78	Morphological and molecular divergence of Rhipicephalus turanicus tick from Albania and China. Experimental and Applied Acarology, 2017, 73, 493-499.	1.6	7
79	Anaplasmataceae closely related to Ehrlichia chaffeensis and Neorickettsia helminthoeca from birds in Central Europe, Hungary. Antonie Van Leeuwenhoek, 2020, 113, 1067-1073.	1.7	7
80	The heart microbiome of insectivorous bats from Central and South Eastern Europe. Comparative Immunology, Microbiology and Infectious Diseases, 2021, 75, 101605.	1.6	7
81	Molecular survey of Babesia spp. in red foxes (Vulpes Vulpes), Asian badgers (Meles leucurus) and their ticks in China. Ticks and Tick-borne Diseases, 2021, 12, 101710.	2.7	7
82	Rickettsiaceae in two reptile-associated tick species, Amblyomma exornatum and Africaniella transversale: First evidence of Occidentia massiliensis in hard ticks (Acari: Ixodidae). Ticks and Tick-borne Diseases, 2022, 13, 101830.	2.7	7
83	Investigation on haplotypes of ixodid ticks and retrospective finding of Borrelia miyamotoi in bank vole (Myodes glareolus) in Switzerland. Ticks and Tick-borne Diseases, 2021, 13, 101865.	2.7	7
84	Description of the male and the larva of Ixodes ariadnae Hornok, 2014. Ticks and Tick-borne Diseases, 2016, 7, 1252-1255.	2.7	6
85	First record of Ixodes ariadnae in Western Europe, Belgium — Short communication. Acta Veterinaria Hungarica, 2016, 64, 467-471.	0.5	6
86	First records and molecular-phylogenetic analyses of three tick species (Ixodes kaiseri, Hyalomma) Tj ETQq0 0 (O rgBT /Ove	rlock 10 Tf 50
87	Morphology of Pholeoixodes species associated with carnivores in the western Palearctic: Pictorial key based on molecularly identified Ixodes (Ph.) canisuga, I. (Ph.) hexagonus and I. (Ph.) kaiseri males, nymphs and larvae. Ticks and Tick-borne Diseases, 2021, 12, 101715.	2.7	6
88	Babesia gibsoni emerging with high prevalence and co-infections in "fighting dogs―in Hungary. Current Research in Parasitology and Vector-borne Diseases, 2021, 1, 100048.	1.9	6
89	Illustrated redescription of the male of Ixodes simplex Neumann, 1906. Ticks and Tick-borne Diseases, 2018, 9, 1328-1330.	2.7	5
90	Diversity of tick species and associated pathogens on peri-urban wild boars – First report of the zoonotic Babesia cf. crassa from Hungary. Ticks and Tick-borne Diseases, 2022, 13, 101936.	2.7	5

#	Article	lF	CITATIONS
91	Haematospirillum and insect Wolbachia DNA in avian blood. Antonie Van Leeuwenhoek, 2018, 111, 479-483.	1.7	4
92	Ecdysteroids are present in the blood of wild passerine birds. Scientific Reports, 2019, 9, 17002.	3.3	4
93	Five ixodid tick species including two morphotypes of Rhipicephalus turanicus on nestlings of Eurasian eagle owl (Bubo bubo) from south-eastern Bulgaria. Parasites and Vectors, 2021, 14, 334.	2.5	4
94	Dogs are final hosts of Sarcocystis morae (Apicomplexa: Sarcocystidae): First report of this species in Hungary and its region – Short communication. Acta Veterinaria Hungarica, 2021, 69, 157-160.	0.5	4
95	First broad-range molecular screening of tick-borne pathogens in Ixodes (Pholeoixodes) kaiseri, with special emphasis on piroplasms. Acta Veterinaria Hungarica, 2020, 68, 30-33.	0.5	4
96	Urban emergence of Dermanyssus gallinae lineage L1 and Ornithonyssus sylviarum in Hungary: phylogenetic differentiation between the roles of migrating vs transported synanthropic birds. Parasites and Vectors, 2021, 14, 147.	2.5	3
97	Description of a new bat-associated bug species of the Cimex lectularius group from Vietnam. Acta Veterinaria Hungarica, 2018, 66, 607-612.	0.5	2
98	Description of the male and the larva of Ixodes collaris Hornok, 2016 with drawings of all stages. Parasites and Vectors, 2019, 12, 144.	2.5	1
99	Latrocimicinae completes the phylogeny of Cimicidae: meeting old morphologic data rather than modern host phylogeny. Parasites and Vectors, 2021, 14, 441.	2.5	1
100	Arthropod moulting hormones (ecdysteroids) are present in the blood of insectivorous bats. Mammal Review, 0, , .	4.8	0