

Samer Angelone

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7597124/publications.pdf>

Version: 2024-02-01

77
papers

1,585
citations

257450

24
h-index

377865

34
g-index

78
all docs

78
docs citations

78
times ranked

1584
citing authors

#	ARTICLE	IF	CITATIONS
1	Sarcoptic mange: An emerging zoonotic in wildlife. <i>Transboundary and Emerging Diseases</i> , 2022, 69, 927-942.	3.0	56
2	Unintentional Recovery of Parasitic Diversity Following Restoration of Red Deer (<i>Cervus elaphus</i>) in North-Western Italy. <i>Animals</i> , 2022, 12, 1433.	2.3	0
3	Biology and management of sarcoptic mange in wild Caprinae populations. <i>Mammal Review</i> , 2021, 51, 82-94.	4.8	18
4	Sarcoptic mange in wild ruminants in Spain: solving the epidemiological enigma using microsatellite markers. <i>Parasites and Vectors</i> , 2021, 14, 171.	2.5	20
5	Demography reveals populational expansion of a recently extinct Iberian ungulate. <i>Zoosystematics and Evolution</i> , 2021, 97, 211-221.	1.1	3
6	First report of interspecific transmission of sarcoptic mange from Iberian ibex to wild boar. <i>Parasites and Vectors</i> , 2021, 14, 481.	2.5	11
7	Filmmaking courses for scientists help promote richer alternatives to chronological narratives. <i>Studies in Higher Education</i> , 2020, 45, 2001-2010.	4.5	4
8	Patterns of helminth infection in Kenyan elephant populations. <i>Parasites and Vectors</i> , 2020, 13, 145.	2.5	4
9	Genetic diversity in natural range remnants of the critically endangered hirola antelope. <i>Zoological Journal of the Linnean Society</i> , 2020, 190, 384-395.	2.3	3
10	Modes of documentary films produced by the future generation of "scientists-as-filmmakers". <i>International Journal of Science Education, Part B: Communication and Public Engagement</i> , 2019, 9, 285-295.	1.5	1
11	Biogeography of Korea's top predator, the yellow-throated Marten: evolutionary history and population dynamics. <i>BMC Evolutionary Biology</i> , 2019, 19, 23.	3.2	2
12	Infection dynamics of gastrointestinal helminths in sympatric non-human primates, livestock and wild ruminants in Kenya. <i>PLoS ONE</i> , 2019, 14, e0217929.	2.5	14
13	A New Generation of Scientists-as-Filmmakers: Experiences Gained in Switzerland. <i>Science Communication</i> , 2019, 41, 369-377.	3.3	11
14	Molecular identification of Ehrlichia, Anaplasma, Babesia and Theileria in African elephants and their ticks. <i>PLoS ONE</i> , 2019, 14, e0226083.	2.5	5
15	Population genomics analyses of European ibex species show lower diversity and higher inbreeding in reintroduced populations. <i>Evolutionary Applications</i> , 2018, 11, 123-139.	3.1	62
16	Hidden MHC genetic diversity in the Iberian ibex (<i>Capra pyrenaica</i>). <i>BMC Genetics</i> , 2018, 19, 28.	2.7	12
17	International meeting on sarcoptic mange in wildlife, June 2018, Blacksburg, Virginia, USA. <i>Parasites and Vectors</i> , 2018, 11, 449.	2.5	33
18	Molecular survey of <i>Coxiella burnetii</i> in wildlife and ticks at wildlife-livestock interfaces in Kenya. <i>Experimental and Applied Acarology</i> , 2017, 72, 277-289.	1.6	20

#	ARTICLE	IF	CITATIONS
19	Molecular Analyses Reveal Unexpected Genetic Structure in Iberian Ibex Populations. PLoS ONE, 2017, 12, e0170827.	2.5	14
20	A practical guideline to remote biopsy darting of wildebeests for genetic sampling. International Journal of Veterinary Science and Medicine, 2016, 4, 27-32.	2.2	4
21	Three Novel Haplotypes of <i>Theileria bicornis</i> in Black and White Rhinoceros in Kenya. Transboundary and Emerging Diseases, 2016, 63, e144-e150.	3.0	4
22	Influence of Massive and Long Distance Migration on Parasite Epidemiology: Lessons from the Great Wildebeest Migration. EcoHealth, 2016, 13, 708-719.	2.0	12
23	The threatening but unpredictable <i>Sarcoptes scabiei</i> : first deadly outbreak in the Himalayan lynx, <i>Lynx lynx isabellinus</i> , from Pakistan. Parasites and Vectors, 2016, 9, 402.	2.5	3
24	First report of <i>Setaria tundra</i> in roe deer (<i>Capreolus capreolus</i>) from the Iberian Peninsula inferred from molecular data: epidemiological implications. Parasites and Vectors, 2016, 9, 521.	2.5	10
25	COMPLETE GENOMIC SEQUENCE OF VIRULENT PIGEON PARAMYXOVIRUS IN LAUGHING DOVES (<i>Streptopelia senegalensis</i>) IN KENYA. Journal of Wildlife Diseases, 2016, 52, 599-608.	0.8	6
26	On the population biology of <i>Sarcoptes scabiei</i> infesting Iberian ibex (<i>Capra pyrenaica</i>). International Journal of Acarology, 2016, 42, 7-11.	0.7	10
27	Universal conventional and real-time PCR diagnosis tools for <i>Sarcoptes scabiei</i> . Parasites and Vectors, 2015, 8, 587.	2.5	39
28	Epidemiology of <i>Theileria bicornis</i> among black and white rhinoceros metapopulation in Kenya. BMC Veterinary Research, 2015, 11, 4.	1.9	9
29	Detusking Fence-Breaker Elephants as an Approach in Human-Elephant Conflict Mitigation. PLoS ONE, 2014, 9, e91749.	2.5	43
30	Advances in studies of disease-navigating webs: <i>Sarcoptes scabiei</i> as a case study. Parasites and Vectors, 2014, 7, 16.	2.5	16
31	Comparative analysis of microRNA profiles between adult <i>Ascaris lumbricoides</i> and <i>Ascaris suum</i> . BMC Veterinary Research, 2014, 10, 99.	1.9	49
32	War diseases revealed by the social media: massive leishmaniasis outbreak in the Syrian Spring. Parasites and Vectors, 2013, 6, 94.	2.5	17
33	Traumatic myiasis in free-ranging eland, reported from Kenya. Parasites and Vectors, 2013, 6, 89.	2.5	11
34	The use of radio-collars for monitoring wildlife diseases: a case study from Iberian ibex affected by <i>Sarcoptes scabiei</i> in Sierra Nevada, Spain. Parasites and Vectors, 2013, 6, 242.	2.5	20
35	The neglected navigating web of the incomprehensibly emerging and re-emerging <i>Sarcoptes</i> mite. Infection, Genetics and Evolution, 2013, 17, 253-259.	2.3	46
36	Genetic epidemiology of <i>Sarcoptes scabiei</i> in the Iberian wolf in Asturias, Spain. Veterinary Parasitology, 2013, 196, 453-459.	1.8	23

#	ARTICLE	IF	CITATIONS
37	Molecular Phylogenetics of the Possibly Extinct Martinique Ground Snake. <i>Herpetologica</i> , 2013, 69, 227.	0.4	3
38	Noninvasive molecular and morphological evidences for an undiscovered population of snow vole in Southern Spain. <i>Mitochondrial DNA</i> , 2013, 24, 596-601.	0.6	2
39	Social and Population Structure in the Ant <i>Cataglyphis emmae</i> . <i>PLoS ONE</i> , 2013, 8, e72941.	2.5	20
40	First reported case of fatal tuberculosis in a wild African elephant with past human-wildlife contact. <i>Epidemiology and Infection</i> , 2013, 141, 1476-1480.	2.1	31
41	Spatio-Temporal Distribution of Injured Elephants in Masai Mara and the Putative Negative and Positive Roles of the Local Community. <i>PLoS ONE</i> , 2013, 8, e71179.	2.5	18
42	Sarcoptic mange and cheetah conservation in Masai Mara (Kenya): epidemiological study in a wildlife/livestock system. <i>Parasitology</i> , 2012, 139, 1587-1595.	1.5	30
43	Sarcoptic-mange detector dogs used to identify infected animals during outbreaks in wildlife. <i>BMC Veterinary Research</i> , 2012, 8, 110.	1.9	40
44	Applicability of major histocompatibility complex DRB1 alleles as markers to detect vertebrate hybridization: a case study from Iberian ibex – domestic goat in southern Spain. <i>Acta Veterinaria Scandinavica</i> , 2012, 54, 56.	1.6	19
45	Putative filariosis outbreak in white and black rhinoceros at Meru National Park in Kenya. <i>Parasites and Vectors</i> , 2012, 5, 206.	2.5	13
46	Applicability of molecular markers to determine parasitic infection origins in the animal trade: a case study from <i>Sarcoptes</i> mites in wildebeest. <i>Forensic Science, Medicine, and Pathology</i> , 2012, 8, 280-284.	1.4	17
47	Microsatellite-based genotyping of MHC class II DRB1 gene in Iberian and Alpine ibex. <i>European Journal of Wildlife Research</i> , 2012, 58, 743-748.	1.4	14
48	The opportunistic <i>Sarcoptes scabiei</i> : A new episode from giraffe in the drought-suffering Kenya. <i>Veterinary Parasitology</i> , 2012, 185, 359-363.	1.8	21
49	Common names of species, the curious case of <i>Capra pyrenaica</i> and the concomitant steps towards the "wild-to-domestic"™ transformation of a flagship species and its vernacular names. <i>Biodiversity and Conservation</i> , 2012, 21, 1-12.	2.6	25
50	Phylogenetic study of <i>Setaria cervi</i> based on mitochondrial <i>cox1</i> gene sequences. <i>Parasitology Research</i> , 2012, 110, 281-285.	1.6	14
51	Single-tube HotSHOT technique for the collection, preservation and PCR-ready DNA preparation of faecal samples: the threatened Cabrera's vole as a model. <i>European Journal of Wildlife Research</i> , 2012, 58, 345-350.	1.4	4
52	Knowledge of Mange among Masai Pastoralists in Kenya. <i>PLoS ONE</i> , 2012, 7, e43342.	2.5	20
53	Siberian tiger's recent population bottleneck in the Russian Far East revealed by microsatellite markers. <i>Mammalian Biology</i> , 2011, 76, 722-726.	1.5	18
54	Applicability of mitochondrial DNA for the identification of Arvicolid species from faecal samples: a case study from the threatened Cabrera's™ vole. <i>Molecular Ecology Resources</i> , 2011, 11, 409-414.	4.8	9

#	ARTICLE	IF	CITATIONS
55	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 December 2010â€“31 January 2011. <i>Molecular Ecology Resources</i> , 2011, 11, 586-589.	4.8	38
56	Sarcoptes-World Molecular Network (Sarcoptes-WMN): integrating research on scabies. <i>International Journal of Infectious Diseases</i> , 2011, 15, e294-e297.	3.3	46
57	Genetic diversity and relatedness of <i>Fasciola</i> spp. isolates from different hosts and geographic regions revealed by analysis of mitochondrial DNA sequences. <i>Veterinary Parasitology</i> , 2011, 181, 329-334.	1.8	28
58	A fluorescence-based polymerase chain reaction-linked single-strand conformation polymorphism (F-PCR-SSCP) assay for the identification of <i>Fasciola</i> spp.. <i>Parasitology Research</i> , 2011, 108, 1513-1517.	1.6	11
59	Efficient identification of <i>Microtus cabreræ</i> excrements using noninvasive molecular analysis. <i>Conservation Genetics Resources</i> , 2011, 3, 127-129.	0.8	8
60	Genetic characterization, species differentiation and detection of <i>Fasciola</i> spp. by molecular approaches. <i>Parasites and Vectors</i> , 2011, 4, 101.	2.5	58
61	Temporal stability in the genetic structure of <i>Sarcoptes scabiei</i> under the host-taxon law: empirical evidences from wildlife-derived <i>Sarcoptes</i> mite in Asturias, Spain. <i>Parasites and Vectors</i> , 2011, 4, 151.	2.5	39
62	The curse of the prey: <i>Sarcoptes</i> mite molecular analysis reveals potential prey-to-predator parasitic infestation in wild animals from Masai Mara, Kenya. <i>Parasites and Vectors</i> , 2011, 4, 193.	2.5	40
63	A TaqMan real-time PCR-based assay for the identification of <i>Fasciola</i> spp.. <i>Veterinary Parasitology</i> , 2011, 179, 266-271.	1.8	41
64	Neatness depends on season, age, and sex in Iberian ibex <i>Capra pyrenaica</i> . <i>Behavioral Ecology</i> , 2011, 22, 1070-1078.	2.2	12
65	Host taxon-derived <i>Sarcoptes</i> mite in European wild animals revealed by microsatellite markers. <i>Biological Conservation</i> , 2010, 143, 1269-1277.	4.1	57
66	<i>Sarcoptes</i> mite from collection to DNA extraction: the lost realm of the neglected parasite. <i>Parasitology Research</i> , 2009, 104, 723-732.	1.6	31
67	Two simple techniques for the safe <i>Sarcoptes</i> collection and individual mite DNA extraction. <i>Parasitology Research</i> , 2009, 105, 1465-1468.	1.6	13
68	Effectiveness of the postponed isolation (post-frozen isolation) method for PCR-quality <i>Sarcoptes</i> mite gDNA. <i>Experimental and Applied Acarology</i> , 2009, 47, 173-178.	1.6	15
69	Is ITS-2 rDNA suitable marker for genetic characterization of <i>Sarcoptes</i> mites from different wild animals in different geographic areas?. <i>Veterinary Parasitology</i> , 2009, 159, 181-185.	1.8	51
70	Bronchopulmonary nematode infection of <i>Capra pyrenaica</i> in the Sierra Nevada massif, Spain. <i>Veterinary Parasitology</i> , 2009, 164, 340-343.	1.8	16
71	Epidemiology of fasciolosis affecting Iberian ibex (<i>Capra pyrenaica</i>) in southern Spain. <i>Parasitology Research</i> , 2008, 102, 751-755.	1.6	24
72	Genetic variability among <i>Fasciola hepatica</i> samples from different host species and geographical localities in Spain revealed by the novel SRAP marker. <i>Parasitology Research</i> , 2008, 103, 181-186.	1.6	22

#	ARTICLE	IF	CITATIONS
73	HotSHOT Plus ThermalSHOCK, a new and efficient technique for preparation of PCR-quality mite genomic DNA. Parasitology Research, 2008, 103, 1455-1457.	1.6	34
74	Skin-scale genetic structure of Sarcoptes scabiei populations from individual hosts: empirical evidence from Iberian ibex-derived mites. Parasitology Research, 2008, 104, 101-105.	1.6	32
75	Characterization of Fasciola samples from different host species and geographical localities in Spain by sequences of internal transcribed spacers of rDNA. Parasitology Research, 2007, 101, 1245-1250.	1.6	35
76	Storyboardgraphy. Visual Studies, 0, , 1-5.	0.5	0
77	<i>Donâ€™t Look Up</i>: Science Communication Revisited. Science Communication, 0, , 107554702210921.	3.3	0