Isabel GarcÃ-a FernÃ;ndez

List of Publications by Year in descending order

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104 papers 3,558 citations

126858 33 h-index 54 g-index

107 all docs

107 docs citations

107 times ranked

3531 citing authors

#	Article	IF	CITATIONS
1	Fatal cases of bovine anaplasmosis in a herd infected with different Anaplasma marginale genotypes in southern Spain. Ticks and Tick-borne Diseases, 2022, 13, 101864.	1.1	7
2	Bagaza Virus in Wild Birds, Portugal, 2021. Emerging Infectious Diseases, 2022, 28, 1504-1506.	2.0	4
3	Bagaza virus and <i>Plasmodium</i> spp. coinfection in redâ€legged partridges (<i>Alectoris rufa</i>), in Southern Spain 2019. Transboundary and Emerging Diseases, 2022, 69, .	1.3	7
4	The antibody response to the glycan αâ€Gal correlates with COVIDâ€19 disease symptoms. Journal of Medical Virology, 2021, 93, 2065-2075.	2.5	25
5	Detection of new Crimean–Congo haemorrhagic fever virus genotypes in ticks feeding on deer and wild boar, Spain. Transboundary and Emerging Diseases, 2021, 68, 993-1000.	1.3	30
6	Microbial community of Hyalomma lusitanicum is dominated by Francisella-like endosymbiont. Ticks and Tick-borne Diseases, 2021, 12, 101624.	1.1	7
7	Detection of environmental SARSâ€CoVâ€2 RNA in a high prevalence setting in Spain. Transboundary and Emerging Diseases, 2021, 68, 1487-1492.	1.3	38
8	Characterization of the anti-α-Gal antibody profile in association with Guillain-Barré syndrome, implications for tick-related allergic reactions. Ticks and Tick-borne Diseases, 2021, 12, 101651.	1.1	7
9	Characterization by Quantitative Serum Proteomics of Immune-Related Prognostic Biomarkers for COVID-19 Symptomatology. Frontiers in Immunology, 2021, 12, 730710.	2.2	30
10	Multi-level analysis of exposure to triazole fungicides through treated seed ingestion in the red-legged partridge. Environmental Research, 2020, 189, 109928.	3.7	23
11	Tick and Host Derived Compounds Detected in the Cement Complex Substance. Biomolecules, 2020, 10, 555.	1.8	32
12	Comparative Proteomic Analysis of Rhipicephalus sanguineus sensu lato (Acari: Ixodidae) Tropical and Temperate Lineages: Uncovering Differences During Ehrlichia canis Infection. Frontiers in Cellular and Infection Microbiology, 2020, 10, 611113.	1.8	6
13	A dataset for the analysis of antibody response to glycan alpha-Gal in individuals with immune-mediated disorders. F1000Research, 2020, 9, 1366.	0.8	3
14	A dataset for the analysis of antibody response to glycan alpha-Gal in individuals with immune-mediated disorders. F1000Research, 2020, 9, 1366.	0.8	4
15	Clinical gamasoidosis and antibody response in two patients infested with Ornithonyssus bursa (Acari: Gamasida: Macronyssidae). Experimental and Applied Acarology, 2019, 78, 555-564.	0.7	12
16	A metaproteomics approach reveals changes in mandibular lymph node microbiota of wild boar naturally exposed to an increasing trend of Mycobacterium tuberculosis complex infection. Tuberculosis, 2019, 114, 103-112.	0.8	2
17	Molecular identification of spotted fever group Rickettsia in ticks collected from dogs and small ruminants in Greece. Experimental and Applied Acarology, 2019, 78, 421-430.	0.7	9
18	Serum haptoglobin response in red deer naturally infected with tuberculosis. Comparative Immunology, Microbiology and Infectious Diseases, 2019, 64, 25-30.	0.7	7

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19	Characterization of the bacterial microbiota in wild-caught Ixodes ventalloi. Ticks and Tick-borne Diseases, 2019, 10, 336-343.	1.1	19
20	Red deer in Iberia: Molecular ecological studies in a southern refugium and inferences on European postglacial colonization history. PLoS ONE, 2019, 14, e0210282.	1.1	29
21	Molecular identification of tick-borne pathogens in ticks collected from dogs and small ruminants from Greece. Experimental and Applied Acarology, 2018, 74, 443-453.	0.7	18
22	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
23	Draft Genome Sequences of Anaplasma phagocytophilum , A.Âmarginale , and A.Âovis Isolates from Different Hosts. Genome Announcements, 2018, 6, .	0.8	6
24	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.	0.7	46
25	Identification and molecular characterization of spotted fever group rickettsiae in ticks collected from farm ruminants in Lebanon. Ticks and Tick-borne Diseases, 2018, 9, 104-108.	1.1	18
26	Biotic and abiotic factors shape the microbiota of wildâ€caught populations of the arbovirus vector <i>Culicoides imicola</i> . Insect Molecular Biology, 2018, 27, 847-861.	1.0	18
27	Combination of RT-PCR and proteomics for the identification of Crimean-Congo hemorrhagic fever virus in ticks. Heliyon, 2017, 3, e00353.	1.4	10
28	Molecular survey of Rickettsial organisms in ectoparasites from a dog shelter in Northern Mexico. Veterinary Parasitology: Regional Studies and Reports, 2017, 10, 143-148.	0.3	2
29	Anaplasma phagocytophilum MSP4 and HSP70 Proteins Are Involved in Interactions with Host Cells during Pathogen Infection. Frontiers in Cellular and Infection Microbiology, 2017, 7, 307.	1.8	44
30	Vaccinomics Approach to the Identification of Candidate Protective Antigens for the Control of Tick Vector Infestations and Anaplasma phagocytophilum Infection. Frontiers in Cellular and Infection Microbiology, 2017, 7, 360.	1.8	34
31	Tick-host conflict: immunoglobulin E antibodies to tick proteins in patients with anaphylaxis to tick bite. Oncotarget, 2017, 8, 20630-20644.	0.8	54
32	Genotypes of <i>Coxiella burnetii</i> in wildlife: disentangling the molecular epidemiology of a multiâ€host pathogen. Environmental Microbiology Reports, 2016, 8, 708-714.	1.0	11
33	Molecular screening for Anaplasmataceae in ticks and tsetse flies from Ethiopia. Acta Veterinaria Hungarica, 2016, 64, 65-70.	0.2	8
34	Evidence of co-infection with Mycobacterium bovis and tick-borne pathogens in a naturally infected sheep flock. Ticks and Tick-borne Diseases, 2016, 7, 384-389.	1.1	4
35	Molecular identification and characterization of Anaplasma platys and Ehrlichia canis in dogs in Mexico. Ticks and Tick-borne Diseases, 2016, 7, 276-283.	1.1	49
36	Expression of Early Growth Response Gene-2 and Regulated Cytokines Correlates with Recovery from Guillain–Barré Syndrome. Journal of Immunology, 2016, 196, 1102-1107.	0.4	15

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37	Molecular detection of vector-borne pathogens in wild and domestic carnivores and their ticks at the human–wildlife interface. Ticks and Tick-borne Diseases, 2016, 7, 284-290.	1.1	77
38	Regulation of the Immune Response to \hat{l}_{\pm} -Gal and Vector-borne Diseases. Trends in Parasitology, 2015, 31, 470-476.	1.5	34
39	Infection and exposure to vector-borne pathogens in rural dogs and their ticks, Uganda. Parasites and Vectors, 2015, 8, 306.	1.0	33
40	Long-Term Dynamics of Coxiella burnetii in Farmed Red Deer (Cervus elaphus). Frontiers in Veterinary Science, 2015, 2, 74.	0.9	12
41	Identification and characterization of a novel tick-borne flavivirus subtype in goats (Capra hircus) in Spain. Journal of General Virology, 2015, 96, 1676-1681.	1.3	21
42	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.	1.0	25
43	Molecular and immunological characterization of three strains of Anaplasma marginale grown in cultured tick cells. Ticks and Tick-borne Diseases, 2015, 6, 522-529.	1.1	9
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.	1.0	23
45	Oral Vaccination with Heat Inactivated Mycobacterium bovis Activates the Complement System to Protect against Tuberculosis. PLoS ONE, 2014, 9, e98048.	1.1	52
46	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.	1.0	30
47	Piroplasmosis in wildlife: Babesia and Theileria affecting free-ranging ungulates and carnivores in the Italian Alps. Parasites and Vectors, 2014, 7, 70.	1.0	92
48	Control of tick infestations and pathogen prevalence in cattle and sheep farms vaccinated with the recombinant Subolesin-Major Surface Protein 1a chimeric antigen. Parasites and Vectors, 2014, 7, 10.	1.0	36
49	Isolation and characterization of Babesia pecorum sp. nov. from farmed red deer (Cervus elaphus). Veterinary Research, 2014, 45, 78.	1.1	9
50	Use of Percoll gradients to purify Anaplasma marginale (Rickettsiales: Anaplasmataceae) from tick cell cultures. Ticks and Tick-borne Diseases, 2014, 5, 511-515.	1.1	7
51	A Systems Biology Approach to the Characterization of Stress Response in Dermacentor reticulatus Tick Unfed Larvae. PLoS ONE, 2014, 9, e89564.	1.1	72
52	Lesser protein degradation machinery correlates with higher BM86 tick vaccine efficacy in Rhipicephalus annulatus when compared to Rhipicephalus microplus. Vaccine, 2013, 31, 4728-4735.	1.7	42
53	Non-pet dogs as sentinels and potential synanthropic reservoirs of tick-borne and zoonotic bacteria. Veterinary Microbiology, 2013, 167, 700-703.	0.8	25
54	High prevalence of Hepatozoon-infection among shepherd dogs in a region considered to be free of Rhipicephalus sanguineus. Veterinary Parasitology, 2013, 196, 189-193.	0.7	54

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55	Optimizing the sampling effort to evaluate body condition in ungulates: A case study on red deer. Ecological Indicators, 2013, 30, 65-71.	2.6	20
56	Sex-related differences in body condition and serum biochemical parameters in red deer (Cervus) Tj ETQq0 0 0 r	gBT/Qver	lock 10 Tf 50 7
57	Molecular evidence of Ehrlichia canis and Rickettsia massiliae in ixodid ticks of carnivores from South Hungary. Acta Veterinaria Hungarica, 2013, 61, 42-50.	0.2	37
58	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.	0.6	46
59	Prevalence of Tick-Borne Pathogens in Adult <i>Dermacentor < li>spp. Ticks from Nine Collection Sites in France. Vector-Borne and Zoonotic Diseases, 2013, 13, 226-236.</i>	0.6	95
60	Sequencing of modern Lepus VDJ genes shows that the usage of VHn genes has been retained in both Oryctolagus and Lepus that diverged 12 million years ago. Immunogenetics, 2013, 65, 777-784.	1.2	18
61	Spotted Fever Group Rickettsiae in Questing Ticks, Central Spain. Emerging Infectious Diseases, 2013, 19, 1163-1165.	2.0	24
62	Usutu Virus in Migratory Song Thrushes, Spain. Emerging Infectious Diseases, 2013, 19, 1173-1175.	2.0	42
63	Proteomics Approach to the Study of Cattle Tick Adaptation to White Tailed Deer. BioMed Research International, 2013, 2013, 1-8.	0.9	17
64	Genetic characterization of Coxiella burnetii in Amblyomma varigatum ticks from North-central Nigeria: public health importance. Veterinary World, 2013, 6, 818-822.	0.7	6
65	Sex-biased differences in the effects of host individual, host population and environmental traits driving tick parasitism in red deer. Frontiers in Cellular and Infection Microbiology, 2013, 3, 23.	1.8	23
66	Factors Driving the Abundance of Ixodes ricinus Ticks and the Prevalence of Zoonotic I. ricinus-Borne Pathogens in Natural Foci. Applied and Environmental Microbiology, 2012, 78, 2669-2676.	1.4	69
67	Fatal bovine anaplasmosis in a herd with new genotypes of Anaplasma marginale, Anaplasma ovis and concurrent haemoplasmosis. Research in Veterinary Science, 2012, 92, 30-35.	0.9	39
68	Natural Bagaza virus infection in game birds in southern Spain. Veterinary Research, 2012, 43, 65.	1.1	38
69	<i>Rickettsia conorii</i> Indian Tick Typhus Strain and <i>R. slovaca</i> Infectious Diseases, 2012, 18, 1008-10.	2.0	23
70	Louping III in Goats, Spain, 2011. Emerging Infectious Diseases, 2012, 18, 976-978.	2.0	37
71	Diagnosis of Tuberculosis in Camelids: Old Problems, Current Solutions and Future Challenges. Transboundary and Emerging Diseases, 2012, 59, 1-10.	1.3	25
72	A database for animal tuberculosis (mycoDB.es) within the context of the Spanish national programme for eradication of bovine tuberculosis. Infection, Genetics and Evolution, 2012, 12, 877-882.	1.0	34

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73	No evidence that wild red deer (Cervus elaphus) on the Iberian Peninsula are a reservoir of Mycobacterium avium subspecies paratuberculosis infection. Veterinary Journal, 2012, 192, 544-546.	0.6	9
74	Unexpected high responses to tuberculin skin-test in farmed red deer: Implications for tuberculosis control. Preventive Veterinary Medicine, 2012, 104, 327-334.	0.7	14
75	Molecular identification of tick-borne pathogens in Nigerian ticks. Veterinary Parasitology, 2012, 187, 572-577.	0.7	62
76	First Molecular Evidence of <i>Anaplasma ovis </i> and <i>Rickettsia </i> spp. in Keds (Diptera:) Tj ETQq0 0 0 rgBT /	Overlock 1	.0 Тf 50 622 Т 83
77	The testing season affects red deer skinfold increase in response to phytohaemagglutinin. Preventive Veterinary Medicine, 2011, 100, 79-83.	0.7	8
78	Assessment of in vivo and in vitro tuberculosis diagnostic tests in Mycobacterium caprae naturally infected caprine flocks. Preventive Veterinary Medicine, 2011, 100, 187-192.	0.7	17
79	Eurasian wild boar response to skin-testing with mycobacterial and non-mycobacterial antigens. Preventive Veterinary Medicine, 2010, 96, 211-217.	0.7	20
80	Comparison of Three Immunological Diagnostic Tests for the Detection of Avian Tuberculosis in Naturally Infected Red Deer (Cervus Elaphus). Journal of Veterinary Diagnostic Investigation, 2009, 21, 102-107.	0.5	9
81	Rickettsia massiliaein the Canary Islands. Emerging Infectious Diseases, 2009, 15, 1869-1870.	2.0	24
82	Factors affecting red deer skin test responsiveness to bovine and avian tuberculin and to phytohaemagglutinin. Preventive Veterinary Medicine, 2009, 90, 119-126.	0.7	25
83	Reduced major histocompatibility complex class II polymorphism in a hunterâ€managed isolated Iberian red deer population. Journal of Zoology, 2009, 277, 157-170.	0.8	13
84	Impact of major histocompatibility complex class II polymorphisms on Iberian red deer parasitism and life history traits. Infection, Genetics and Evolution, 2009, 9, 1232-1239.	1.0	20
85	Epidemiological risk factors of Aujeszky's disease in wild boars (Sus scrofa) and domestic pigs in Spain. European Journal of Wildlife Research, 2008, 54, 549-555.	0.7	32
86	Differential expression of inflammatory and immune response genes in mesenteric lymph nodes of Iberian red deer (Cervus elaphus hispanicus) naturally infected with Mycobacterium bovis. Developmental and Comparative Immunology, 2008, 32, 85-91.	1.0	27
87	The effects of sex and age on phytohaemagglutinin skin-testing of deer. New Zealand Veterinary Journal, 2008, 56, 71-73.	0.4	12
88	Sequence analysis of the msp4 gene of Anaplasma ovis strains. Veterinary Microbiology, 2007, 119, 375-381.	0.8	152
89	The importance of parasite life history and host density in predicting the impact of infections in red deer. Oecologia, 2007, 152, 655-664.	0.9	60
90	Risk factors associated with the prevalence of tuberculosis-like lesions in fenced wild boar and red deer in south central Spain. Veterinary Research, 2007, 38, 451-464.	1.1	143

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91	Ixodid ticks parasitizing Iberian red deer (Cervus elaphus hispanicus) and European wild boar (Sus) Tj ETQq1 10.	784 <u>3</u> 14 rş	gBT ₁₀ 9verlock
92	Molecular Epidemiology of Human and Bovine Anaplasmosis in Southern Europe. Annals of the New York Academy of Sciences, 2006, 1078, 95-99.	1.8	29
93	Epidemiology and risk factors analysis of elaphostrongylosis in red deer (Cervus elaphus) from Spain. Parasitology Research, 2006, 98, 77-85.	0.6	27
94	Efficacy of in-feed-administered ivermectin on Elaphostrongylus cervi first-stage excretion in red deer (Cervus elaphus). Parasitology Research, 2006, 98, 176-178.	0.6	6
95	Effects of parasitic helminths and ivermectin treatment on clinical parameters in the European wild boar (Sus scrofa). Parasitology Research, 2006, 98, 582-587.	0.6	23
96	Optimal dose and timing in phytohaemagglutinin skin-testing of deer. New Zealand Veterinary Journal, 2006, 54, 357-359.	0.4	10
97	Wild boar and red deer display high prevalences of tuberculosis-like lesions in Spain. Veterinary Research, 2006, 37, 107-119.	1.1	165
98	Genetic resistance to bovine tuberculosis in the Iberian wild boar. Molecular Ecology, 2005, 14, 3209-3217.	2.0	114
99	Serosurvey of Aujeszky's disease virus infection in European wild boar in Spain. Veterinary Record, 2005, 156, 408-412.	0.2	55
100	Potential Vertebrate Reservoir Hosts and Invertebrate Vectors of Anaplasma marginale and A. phagocytophilumin Central Spain. Vector-Borne and Zoonotic Diseases, 2005, 5, 390-401.	0.6	119
101	Molecular characterization of Mycobacterium tuberculosis complex isolates from wild ungulates in south-central Spain. Veterinary Research, 2005, 36, 43-52.	1.1	109
102	infection in free-ranging Iberian red deer in the region of Castilla-La Mancha, Spain. Veterinary Microbiology, 2004, 100, 163-173.	0.8	72
103	Efficacy of an in-feed preparation of ivermectin against helminths in the European wild boar. Parasitology Research, 2004, 92, 133-136.	0.6	17
104	Wild boar helminths: risks in animal translocations. Veterinary Parasitology, 2003, 115, 335-341.	0.7	59