Sergio Sanchez

List of Publications by Year
in descending order

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32 papers	824 citations	516710 16 h-index	501196 28 g-index
35 all docs	35 docs citations	35 times ranked	1084 citing authors

#	Article	IF	Citations
1	Methicillin resistant Staphylococcus aureus (MRSA) carriage in different free-living wild animal species in Spain. Veterinary Journal, 2013, 198, 127-130.	1.7	72
2	Prevalence, serotypes and virulence genes of Shiga toxin-producing Escherichia coli isolated from ovine and caprine milk and other dairy products in Spain. International Journal of Food Microbiology, 2006, 107, 212-217.	4.7	67
3	Detection and characterisation of Shiga toxin-producing Escherichia coli other than Escherichia coli O157:H7 in wild ruminants. Veterinary Journal, 2009, 180, 384-388.	1.7	67
4	Prevalence of Shiga toxin-producing Escherichia coli, Salmonella spp. and Campylobacter spp. in large game animals intended for consumption: Relationship with management practices and livestock influence. Veterinary Microbiology, 2013, 163, 274-281.	1.9	57
5	Detection and characterisation of O157:H7 and non-O157 Shiga toxin-producing Escherichia coli in wild boars. Veterinary Microbiology, 2010, 143, 420-423.	1.9	50
6	A new pathogenicity island carrying an allelic variant of the Subtilase cytotoxin is common among Shiga toxin producing Escherichia coli of human and ovine origin. Clinical Microbiology and Infection, 2013, 19, E149-E156.	6.0	50
7	Carriage of Staphylococcus aureus by Free-Living Wild Animals in Spain. Applied and Environmental Microbiology, 2014, 80, 4865-4870.	3.1	48
8	Eutrophication and Bacterial Pathogens as Risk Factors for Avian Botulism Outbreaks in Wetlands Receiving Effluents from Urban Wastewater Treatment Plants. Applied and Environmental Microbiology, 2014, 80, 4251-4259.	3.1	46
9	Presence of Shiga toxin-producing E. coli O157:H7 in a survey of wild artiodactyls. Veterinary Microbiology, 2007, 121, 373-377.	1.9	40
10	Characterization of an emergent clone of enteroinvasive Escherichia coli circulating in Europe. Clinical Microbiology and Infection, 2016, 22, 287.e11-287.e19.	6.0	38
11	Development of Three Multiplex PCR Assays Targeting the 21 Most Clinically Relevant Serogroups Associated with Shiga Toxin-Producing E. coli Infection in Humans. PLoS ONE, 2015, 10, e0117660.	2.5	30
12	Pheno-genotypic characterisation of Escherichia coli O157:H7 isolates from domestic and wild ruminants. Veterinary Microbiology, 2010, 142, 445-449.	1.9	25
13	Subtilase cytotoxin encoding genes are present in human, sheep and deer intimin-negative, Shiga toxin-producing Escherichia coli O128:H2. Veterinary Microbiology, 2012, 159, 531-535.	1.9	25
14	Detection and characterization of Shiga toxin-producing Escherichia coli in game meat and ready-to-eat meat products. International Journal of Food Microbiology, 2012, 160, 179-182.	4.7	24
15	Sorbitol-Fermenting, Î²-Glucuronidaseâ€“Positive, Shiga Toxinâ€“Negative Escherichia coli O157:H7 in Free-Ranging Red Deer in South-Central Spain. Foodborne Pathogens and Disease, 2011, 8, 1313-1315.	1.8	19
16	Variation in the prevalence of non-O157 Shiga toxin-producing Escherichia coli in four sheep flocks during a 12-month longitudinal study. Small Ruminant Research, 2010, 93, 144-148.	1.2	16
17	Occurrence of avian pathogenicEscherichia coliand antimicrobial-resistantE. coliin red-legged partridges (Alectoris rufa): sanitary concerns of farming. Avian Pathology, 2012, 41, 337-344.	2.0	16
18	Longitudinal Study of Shiga Toxin-Producing Escherichia coli Shedding in Sheep Feces: Persistence of Specific Clones in Sheep Flocks. Applied and Environmental Microbiology, 2009, 75, 1769-1773.	3.1	15

#	Article	IF	Citations
19	A zoonotic ringworm outbreak caused by a dysgonic strain of Microsporum canis from stray cats. Revista Iberoamericana De Micologia, 2010, 27, 62-65.	0.9	14
20	Outbreak of ringworm in a traditional Iberian pig farm in Spain. Mycoses, 2011, 54, 179-181.	4.0	12
21	The new allelic variant of the subtilase cytotoxin (subAB2) is common among Shiga toxin-producing Escherichia coli strains from large game animals and their meat and meat products. Veterinary Microbiology, 2013, 166, 645-649.	1.9	12
22	Detection and characterization of Shiga toxin-producing Escherichia coli (STEC) in bulk tank ewesâ€™ milk and sheep farm environment. Small Ruminant Research, 2017, 154, 110-114.	1.2	10
23	Shiga toxin-producing Escherichia coli O157:H7 from extensive cattle of the fighting bulls breed. Research in Veterinary Science, 2010, 88, 208-210.	1.9	9
24	Cluster investigation of mixed O76:H19 Shiga toxin-producingEscherichia coliand atypical enteropathogenicE. coliinfection in a Spanish household. Epidemiology and Infection, 2014, 142, 1029-1033.	2.1	9
25	Mucus-Activatable Shiga Toxin Genotypestx2dinEscherichia coliO157:H7. Emerging Infectious Diseases, 2017, 23, 1431-1433.	4.3	9
26	Helcococcus ovis isolated from a goat with purulent bronchopneumonia and pulmonary abscesses. Journal of Veterinary Diagnostic Investigation, 2012, 24, 235-237.	1.1	8
27	A Colibacillosis Outbreak in Farmed Red-Legged Partridges (Alectoris rufa). Avian Diseases, 2013, 57, 143-146.	1.0	8
28	Occurrence of verocytotoxin-producing Escherichia coli in the faeces of free-ranging wild lagomorphs in southwest Spain. European Journal of Wildlife Research, 2011, 57, 187-189.	1.4	7
29	Plasmid-Mediated Quinolone Resistance in Different Diarrheagenic Escherichia coli Pathotypes Responsible for Complicated, Noncomplicated, and Traveler's Diarrhea Cases. Antimicrobial Agents and Chemotherapy, 2016, 60, 1950-1951.	3.2	7
30	SalmonellaSpp. and Shiga Toxinâ€“ProducingEscherichia coliPrevalence in an Ocellated Lizard (Timon lepidus) Research Center in Spain. Foodborne Pathogens and Disease, 2011, 8, 1309-1311.	1.8	4
31	Evaluation of the SHIGA TOXIN QUIK CHEK after overnight enrichment as screening tool for Shiga toxinâ€“producing Escherichia coli detection in human fecal samples. Diagnostic Microbiology and Infectious Disease, 2019, 94, 218-222.	1.8	1
32	InfecciÃ³n mixta por cuatro patotipos diarreagÃ©nicos de Escherichia coli en un caso de diarrea del viajero: caracterizaciÃ³n de los aislados obtenidos mediante secuenciaciÃ³n del genoma completo. Enfermedades Infecciosas Y MicrobiologÃa ClÃnica, 2020, 38, 39-40.	0.5	0