

Friederike Zeeh

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

130
citations

1684188

5
h-index

1199594

12
g-index

17
all docs

17
docs citations

17
times ranked

123
citing authors

#	ARTICLE	IF	CITATIONS
1	Diversity of <i>Mycoplasma hyopneumoniae</i> in pig farms revealed by direct molecular typing of clinical material. <i>Veterinary Research</i> , 2007, 38, 391-398.	3.0	51
2	Transposon-associated lincosamide resistance <i>lnu</i> (C) gene identified in <i>Brachyspira hyodysenteriae</i> ST83. <i>Veterinary Microbiology</i> , 2018, 214, 51-55.	1.9	30
3	Predominance of a macrolide-lincosamide-resistant <i>Brachyspira hyodysenteriae</i> of sequence type 196 in Swiss pig herds. <i>Veterinary Microbiology</i> , 2018, 226, 97-102.	1.9	12
4	Occurrence of <i>Brachyspira hyodysenteriae</i> in multiplier pig herds in Switzerland. <i>Tierärztliche Praxis Ausgabe G: Grosstiere - Nutztiere</i> , 2016, 44, 13-18.	0.5	8
5	Risk factors for the infection with <i>Brachyspira hyodysenteriae</i> in pig herds. <i>Preventive Veterinary Medicine</i> , 2020, 174, 104819.	1.9	6
6	Isolation of <i>Brachyspira hyodysenteriae</i> from a crow (<i>Corvus corone</i>) in close proximity to commercial pigs. <i>Veterinary Journal</i> , 2018, 236, 111-112.	1.7	5
7	Seroprevalence of atypical porcine pestivirus in a closed pig herd with subclinical infection. <i>Transboundary and Emerging Diseases</i> , 2020, 67, 2770-2774.	3.0	5
8	Effect of soy on faecal dry matter content and excretion of <i>Brachyspira hyodysenteriae</i> in pigs. <i>Veterinary Record Open</i> , 2016, 3, e000159.	1.0	3
9	A review of methods used for studying the molecular epidemiology of <i>Brachyspira hyodysenteriae</i> . <i>Veterinary Microbiology</i> , 2017, 207, 181-194.	1.9	2
10	First identification and characterisation of <i>Brachyspira hyodysenteriae</i> in pigs in Hong Kong. <i>Porcine Health Management</i> , 2019, 5, 27.	2.6	2
11	Low occurrence of <i>Brachyspira hyodysenteriae</i> in Swiss pig herds with diarrhoea. <i>Schweizer Archiv Fur Tierheilkunde</i> , 2021, 163, 595-599.	0.8	2
12	Chronological clinical and pathological documentation of porcine ulcerative dermatitis. <i>Veterinary Dermatology</i> , 2016, 27, 315.	1.2	1
13	Complete Circular Genome Sequences of <i>Brachyspira hyodysenteriae</i> Isolates of the Four Different Sequence Types Causing Swine Dysentery in Switzerland. <i>Microbiology Resource Announcements</i> , 2021, 10, e0084721.	0.6	1
14	Distribution, genetic heterogeneity, and antimicrobial susceptibility of <i>Brachyspira pilosicoli</i> in Swiss pig herds. <i>Veterinary Microbiology</i> , 2022, 269, 109421.	1.9	1
15	Whole-genome analyses reveal a novel prophage and cgSNPs-derived sublineages of <i>Brachyspira hyodysenteriae</i> ST196. <i>BMC Genomics</i> , 2022, 23, 131.	2.8	0
16	Factors driving pig owners' motivation and satisfaction to perform eradications from Swine dysentery. <i>Preventive Veterinary Medicine</i> , 2022, 204, 105638.	1.9	0