Joana Revez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Assessment of the feed additive consisting of Lactococcus lactis NCIMB 30160 for all animal species for the renewal of its authorisation (Lactosan GmbH & Co KG). EFSA Journal, 2022, 20, e06975.	1.8	0
2	Safety and efficacy of the feed additive consisting of Bacillus licheniformis DSM 28710 (Bâ€Act®) for laying hens, minor poultry species for laying, poultry species for breeding purposes and ornamental birds (HuvePharma N.V.). EFSA Journal, 2021, 19, e06449.	1.8	2
3	Assessment of the feed additive consisting of Lactiplantibacillus plantarum (formerly Lactobacillus) Tj ETQq1 1 0.	784314 rg 1.8	BT /Overlock 3
4	Assessment of the feed additive consisting of Lactiplantibacillus plantarum (formerly Lactobacillus) Tj ETQq0 0 0	rgBT /Over 1.8	lock 10 Tf 50 0
5	Assessment of the feed additive consisting of Lactiplantibacillus plantarum (formerly Lactobacillus) Tj ETQq1 1 0.	784314 rg 1.8	BT /Overlock 0
6	Safety and efficacy of the feed additive consisting of Bacillus velezensisÂCECT 5940 (Ecobiol®) for turkeys for fattening, turkeys reared for breeding, minor poultry species for fattening and reared for laying and ornamental birds (Evonik Operations GmbH). EFSA Journal, 2021, 19, e06620.	1.8	1
7	Assessment of the feed additive consisting of Pediococcus pentosaceusÂDSM 12834 for all animal species for the renewal of its authorisation (Lactosan GmbH & Co KG). EFSA Journal, 2021, 19, e06713.	1.8	1
8	Safety and efficacy of a feed additive consisting of Lactiplantibacillus plantarum (formerly) Tj ETQq0 0 0 rgBT /Ov	erlock 10 1.8	۲f 50 467 Td 3
9	Safety and efficacy of a feed additive consisting of Lactiplantibacillus plantarum (formerly) Tj ETQq1 1 0.784314	rgBT /Over 1.8	lock 10 Tf 50 2
10	Assessment of the feed additive consisting of Lentilactobacillus buchneri (formerly Lactobacillus) Tj ETQq0 0 0 rg	BT /Overlo 1.8	ck 10 Tf 50 3 19
11	Safety and efficacy of a feed additive consisting of Lacticaseibacillus rhamnosus (formerly) Tj ETQq1 1 0.784314	rgBT /Over 1.8	lock 10 Tf 50 2
12	Assessment of the feed additive consisting of Pediococcus acidilacticiÂDSM 16243 for all animal species for the renewal of its authorisation (Lactosan GmbH & Co.KG). EFSA Journal, 2021, 19, e06697.	1.8	1
13	Safety and efficacy of a feed additive consisting of Pediococcus pentosaceus IMI 507024 for all animal species (ALLâ€TECHNOLOGY (IRELAND) LIMITED [Alltech Ireland]). EFSA Journal, 2021, 19, e06701.	1.8	1
14	Safety and efficacy of a feed additive consisting of Lactiplantibacillus plantarum (formerly) Tj ETQq0 0 0 rgBT /Ov	erlock 10 1.8	Մf 50 227 Td 3
15	Safety and efficacy of a feed additive consisting of Pediococcus pentosaceus IMI 507025 for all animal species (ALLâ€TECHNOLOGY (IRELAND) LIMITED [Alltech Ireland]). EFSA Journal, 2021, 19, e06702.	1.8	1
16	Safety and efficacy of a feed additive consisting of Lactiplantibacillus plantarum (formerly) Tj ETQq0 0 0 rgBT /Ov e06898.	erlock 10 1.8	If 50 147 Td 0
17	Assessment of the feed additive consisting of Levilactobacillus brevis (formerly Lactobacillus brevis) DSM 12835 EU for all animal species for the renewal of its authorisation (Lactosan GmbH & Co KG). EFSA Journal, 2021, 19, e06900.	1.8	1
18	Safety and efficacy of a feed additive consisting of Lacticaseibacillus rhamnosus (formerly) Tj ETQq0 0 0 rgBT /Ov (Lactosan GmbH & Co. KG). EFSA Journal, 2021, 19, e06901.	erlock 10 1.8	If 50 67 Td (3

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#	Article	IF	CITATIONS
19	Assessment of the feed additive consisting of Lacticaseibacillus paracasei (formerly Lactobacillus) Tj ETQq1 1 0	.784314 rgl 1.8	BT /Overlock 0
20	Safety and efficacy of a feed additive consisting of Bacillus subtilis strains CNCM lâ€4606, CNCM lâ€5043 and CNCM lâ€4607 and Lactococcus lactisÂCNCM lâ€4609 for all animal species (Nolivade). EFSA Journal, 2021, 19, e06907.	1.8	2
21	EFSA and ECDC technical report on the collection and analysis of whole genome sequencing data from foodâ€borne pathogens and other relevant microorganisms isolated from human, animal, food, feed and food/feed environmental samples in the joint ECDCâ€EFSA molecular typing database. EFSA Supporting Publications. 2019. 16. 1337E.	0.7	19
22	Survey on the Use of Whole-Genome Sequencing for Infectious Diseases Surveillance: Rapid Expansion of European National Capacities, 2015–2016. Frontiers in Public Health, 2017, 5, 347.	2.7	99
23	Large Sequence Diversity within the Biosynthesis Locus and Common Biochemical Features of Campylobacter coli Lipooligosaccharides. Journal of Bacteriology, 2016, 198, 2829-2840.	2.2	13
24	Genome analysis of Campylobacter jejuni strains isolated from a waterborne outbreak. BMC Genomics, 2014, 15, 768.	2.8	40
25	Genomic Variation between Campylobacter jejuni Isolates Associated with Milk-Borne-Disease Outbreaks. Journal of Clinical Microbiology, 2014, 52, 2782-2786.	3.9	47
26	Comparative genomics of unintrogressed Campylobacter coli clades 2 and 3. BMC Genomics, 2014, 15, 129.	2.8	36
27	Novel Microbiological and Spatial Statistical Methods to Improve Strength of Epidemiological Evidence in a Community-Wide Waterborne Outbreak. PLoS ONE, 2014, 9, e104713.	2.5	35
28	Contingency nature of Helicobacter bizzozeronii oxygen-insensitive NAD(P)H-nitroreductase (HBZC1_00960) and its role in metronidazole resistance. Veterinary Research, 2013, 44, 56.	3.0	9
29	Effect of ciprofloxacin exposure on DNA repair mechanisms in Campylobacter jejuni. Microbiology (United Kingdom), 2013, 159, 2513-2523.	1.8	5
30	Genetic heterogeneity of Campylobacter jejuni NCTC 11168 upon human infection. Infection, Genetics and Evolution, 2013, 16, 305-309.	2.3	29
31	Arcobacter Species and Their Pulsed-Field Gel Electrophoresis Genotypes in Finnish Raw Milk during Summer 2011. Journal of Food Protection, 2013, 76, 1630-1632.	1.7	13
32	Pathogenic Bacteria in Finnish Bulk Tank Milk. Foodborne Pathogens and Disease, 2013, 10, 99-106.	1.8	57
33	Association of Campylobacter jejuni Metabolic Traits with Multilocus Sequence Types. Applied and Environmental Microbiology, 2012, 78, 5550-5554.	3.1	34
34	Complete Genome Sequence of a Variant of Campylobacter jejuni NCTC 11168. Journal of Bacteriology, 2012, 194, 6298-6299.	2.2	11
35	Evidence for Conserved Function of γ–Glutamyltranspeptidase in Helicobacter Genus. PLoS ONE, 2012, 7, e30543.	2.5	28
36	Finnish Campylobacter jejuni Strains of Multilocus Sequence Type ST-22 Complex Have Two Lineages with Different Characteristics. PLoS ONE, 2011, 6, e26880.	2.5	26