## Matthias Filter

List of Publications by Year in descending order

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Version: 2024-02-01

567247 454934 49 956 15 30 citations h-index g-index papers 57 57 57 1243 citing authors docs citations times ranked all docs

#	Article	IF	Citations
1	Virulence genotype of Pasteurella multocida strains isolated from different hosts with various disease status. Veterinary Microbiology, 2006, 114, 304-317.	1.9	180
2	Thermal Stability of Hepatitis E Virus as Estimated by a Cell Culture Method. Applied and Environmental Microbiology, 2016, 82, 4225-4231.	3.1	92
3	Thermal stability of hepatitis E virus assessed by a molecular biological approach. Virology Journal, 2011, 8, 487.	3.4	77
4	Impact of the probiotic bacteria i>Enterococcus faecium i>NCIMB 10415 (SF68) and i>Bacillus cereus i>var. toyoi NCIMB 40112 on the development of serum IgG and faecal IgA of sows and their piglets. Archives of Animal Nutrition, 2007, 61, 223-234.	1.8	64
5	Isolation and Characterization of Intestinal <i>Escherichia coli</i> li> Clones from Wild Boars in Germany. Applied and Environmental Microbiology, 2009, 75, 695-702.	3.1	53
6	Survival of Brucella spp. in mineral water, milk and yogurt. International Journal of Food Microbiology, 2011, 145, 326-330.	4.7	50
7	ExPECâ€typical virulenceâ€associated genes correlate with successful colonization by intestinal <i>E.â€fcoli</i> in a small piglet group. Environmental Microbiology, 2008, 10, 1742-1751.	3.8	47
8	Identification of Noncanonical Melanoma-Associated T Cell Epitopes for Cancer Immunotherapy. Journal of Immunology, 2005, 174, 6716-6724.	0.8	32
9	A one health glossary to support communication and information exchange between the human health, animal health and food safety sectors. One Health, 2021, 13, 100263.	3.4	28
10	FoodChain-Lab: A Trace-Back and Trace-Forward Tool Developed and Applied during Food-Borne Disease Outbreak Investigations in Germany and Europe. PLoS ONE, 2016, 11, e0151977.	2.5	27
11	Effects of Bacillus cereus var. toyoi on immune parameters of pregnant sows. Veterinary Immunology and Immunopathology, 2009, 127, 26-37.	1.2	24
12	Harmonized terms, concepts and metadata for microbiological risk assessment models: The basis for knowledge integration and exchange. Microbial Risk Analysis, 2018, 10, 3-12.	2.3	24
13	A Likelihood-Based Approach to Identifying Contaminated Food Products Using Sales Data: Performance and Challenges. PLoS Computational Biology, 2014, 10, e1003692.	3.2	21
14	Towards transparent and consistent exchange of knowledge for improved microbiological food safety. Current Opinion in Food Science, 2018, 19, 129-137.	8.0	21
15	FSK-Lab – An open source food safety model integration tool. Microbial Risk Analysis, 2018, 10, 13-19.	2.3	21
16	STEM: An Open Source Tool for Disease Modeling. Health Security, 2019, 17, 291-306.	1.8	16
17	Predictive models for thermal inactivation of human norovirus and surrogates in strawberry puree. Food Control, 2019, 96, 87-97.	5.5	16
18	Studies on the effect of an <i>Enterococcus faecium</i> probiotic on T cell populations in peripheral blood and intestinal epithelium and on the susceptibility to <i>Salmonella</i> during a challenge infection with <i>Salmonella</i> Typhimurium in piglets. Archives of Animal Nutrition, 2011, 65, 415-430.	1.8	15

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19	Physical and transcriptional map of the critical region for keratolytic winter erythema (KWE) on chromosome 8p22-p23 between D8S550 and D8S1759. European Journal of Human Genetics, 2002, 10, 17-25.	2.8	14
20	One Health Surveillance Codex: promoting the adoption of One Health solutions within and across European countries. One Health, 2021, 12, 100233.	3.4	13
21	Enterobacteriaceae populations during experimental Salmonella infection in pigs. Veterinary Microbiology, 2010, 142, 352-360.	1.9	12
22	Antimicrobial resistances do not affect colonization parameters of intestinal E. coli in a small piglet group. Gut Pathogens, 2009, $1,18$ .	3.4	11
23	Expert systems for food safety. Current Opinion in Food Science, 2015, 6, 61-65.	8.0	10
24	Influence of an <i>Enterococcus faecium</i> probiotic on the development of Peyer's patches B cells in piglets. Archives of Animal Nutrition, 2009, 63, 343-355.	1.8	9
25	A Strategy for the Identification of Canonical and Non-canonical MHCâ€I-binding Epitopes Using an ANN-based Epitope Prediction Algorithm. QSAR and Combinatorial Science, 2006, 25, 350-358.	1.4	7
26	Accelerating investigation of food-borne disease outbreaks using pro-active geospatial modeling of food supply chains. , $2012$ , , .		7
27	A Generic Open-Source Software Framework Supporting Scenario Simulations in Bioterrorist Crises. Biosecurity and Bioterrorism, 2013, 11, S134-S145.	1.2	7
28	Survival of Trichinella spiralis in cured meat products. Veterinary Parasitology, 2020, 287, 109260.	1.8	6
29	Open Science meets Food Modelling:Âlntroducing the Food Modelling Journal (FMJ). Food Modelling Journal, 0, 1, .	0.0	6
30	Towards efficient use of data, models and tools in food microbiology. Current Opinion in Food Science, 2022, 46, 100834.	8.0	5
31	Development of a Comparative Risk Ranking System for Agents Posing a Bioterrorism Threat to Human or Animal Populations. Biosecurity and Bioterrorism, 2013, 11, S3-S16.	1.2	4
32	Realizing virtual research environments for the agriâ€food community: The AGINFRA PLUS experience. Concurrency Computation Practice and Experience, 2021, 33, e6087.	2.2	4
33	Minimum Information Required to Annotate Food Safety Risk Assessment Models (MIRARAM). Food Research International, 2021, 139, 109952.	6.2	4
34	Big Data in Agricultural and Food Research: Challenges and Opportunities of an Integrated Big Data E-infrastructure. Studies in Big Data, 2019, , 129-150.	1.1	4
35	Towards a Food Safety Knowledge Base Applicable in Crisis Situations and Beyond. BioMed Research International, 2015, 2015, 1-11.	1.9	3
36	Towards Community Driven Food Safety Model Repositories. Procedia Food Science, 2016, 7, 105-108.	0.6	3

#	Article	IF	CITATIONS
37	Project DEMETER: Concept Note for an Emerging Risks Knowledge Exchange Platform (ERKEP) Framework. EFSA Supporting Publications, 2018, 15, 1524E.	0.7	3
38	Establishment of a prototypic Quantitative Microbial Risk Assessment (QMRA) food and feed safety model repository. EFSA Supporting Publications, 2019, 16, 1701E.	0.7	3
39	Determination and Metrics for Emerging Risks Identification DEMETER: Final Report. EFSA Supporting Publications, 2020, 17, 1889E.	0.7	3
40	FoodChain-lab: Tracing Software Supporting Foodborne Disease Outbreak Investigations. Procedia Food Science, 2016, 7, 101-104.	0.6	2
41	Exploitation of Commercial B2B Data for Risk Assessment Tasks in Foodborne Crisis Events. Communications in Computer and Information Science, 2012, , 471-474.	0.5	2
42	Bioinformatics: From Peptides to Profiled Leads. , 0, , 1771-1801.		2
43	An Open-Source Community Resource for Creating, Collecting, Sharing and Applying Predictive Microbial Models (PMM-Lab). Communications in Computer and Information Science, 2012, , 462-465.	0.5	1
44	Reprint of: Survival of Trichinella spiralis in cured meat products. Veterinary Parasitology, 2021, 297, 109544.	1.8	1
45	Temperature-Dependent Growth Characteristics of Bacillus thuringiensis in a Ratatouille Food Model. Journal of Food Protection, 2020, 83, 816-820.	1.7	1
46	The Glossaryfication Web Service $\hat{a}\in$ " an automated glossary creation tool to support One Health communication. ARPHA Conference Abstracts, 0, 4, .	0.0	0
47	Making Linked Data accessible for One Health Surveillance with the "One Health Linked Data Toolbox". ARPHA Conference Abstracts, 0, 4, .	0.0	О
48	DARWINIZER® â€" a Computer Based Method for Peptide and Peptidomimetics Design. , 2001, , 430-431.		0
49	One Health Consensus Report Annotation Checklist ( <scp>OHâ€CRAC</scp> ): A crossâ€sector checklist to support harmonized annotation of surveillance data in reports. Zoonoses and Public Health, 0, , .	2.2	0