

Liese Van Gompel

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5355991/publications.pdf>

Version: 2024-02-01

19
papers

770
citations

687363

13
h-index

839539

18
g-index

21
all docs

21
docs citations

21
times ranked

913
citing authors

#	ARTICLE	IF	CITATIONS
1	Risk factors for the abundance of antimicrobial resistance genes <i>aph(3ϵ)-III</i> , <i>erm(B)</i> , <i>sul2</i> and <i>tet(W)</i> in pig and broiler faeces in nine European countries. <i>Journal of Antimicrobial Chemotherapy</i> , 2022, 77, 969-978.	3.0	5
2	Determinants for antimicrobial resistance genes in farm dust on 333 poultry and pig farms in nine European countries. <i>Environmental Research</i> , 2022, 208, 112715.	7.5	21
3	Antimicrobial resistance genes <i>aph(3ϵ)-III</i> , <i>erm(B)</i> , <i>sul2</i> and <i>tet(W)</i> abundance in animal faeces, meat, production environments and human faeces in Europe. <i>Journal of Antimicrobial Chemotherapy</i> , 2022, 77, 1883-1893.	3.0	6
4	Risk Factors for Antimicrobial Resistance in Turkey Farms: A Cross-Sectional Study in Three European Countries. <i>Antibiotics</i> , 2021, 10, 820.	3.7	8
5	Assigning Defined Daily/Course Doses for Antimicrobials in Turkeys to Enable a Cross-Country Quantification and Comparison of Antimicrobial Use. <i>Antibiotics</i> , 2021, 10, 971.	3.7	1
6	Occupational Exposure and Carriage of Antimicrobial Resistance Genes (<i>tetW</i> , <i>ermB</i>) in Pig Slaughterhouse Workers. <i>Annals of Work Exposures and Health</i> , 2020, 64, 125-137.	1.4	16
7	Description and determinants of the faecal resistome and microbiome of farmers and slaughterhouse workers: A metagenome-wide cross-sectional study. <i>Environment International</i> , 2020, 143, 105939.	10.0	33
8	Association of antimicrobial usage with faecal abundance of <i>aph(3ϵ)-III</i> , <i>ermB</i> , <i>sul2</i> and <i>tetW</i> resistance genes in veal calves in three European countries. <i>International Journal of Antimicrobial Agents</i> , 2020, 56, 106131.	2.5	8
9	Farm dust resistomes and bacterial microbiomes in European poultry and pig farms. <i>Environment International</i> , 2020, 143, 105971.	10.0	66
10	Addressing Learning Needs on the Use of Metagenomics in Antimicrobial Resistance Surveillance. <i>Frontiers in Public Health</i> , 2020, 8, 38.	2.7	11
11	Antimicrobial Usage and Resistance in Companion Animals: A Cross-Sectional Study in Three European Countries. <i>Antibiotics</i> , 2020, 9, 87.	3.7	72
12	Metagenomics-Based Approach to Source-Attribution of Antimicrobial Resistance Determinants – Identification of Reservoir Resistome Signatures. <i>Frontiers in Microbiology</i> , 2020, 11, 601407.	3.5	29
13	Associations between antimicrobial use and the faecal resistome on broiler farms from nine European countries. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 2596-2604.	3.0	49
14	Dynamics of faecal shedding of ESBL- or AmpC-producing <i>Escherichia coli</i> on dairy farms. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 1531-1538.	3.0	24
15	Quantitative and qualitative analysis of antimicrobial usage patterns in 180 selected farrow-to-finish pig farms from nine European countries based on single batch and purchase data. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 807-816.	3.0	64
16	The antimicrobial resistome in relation to antimicrobial use and biosecurity in pig farming, a metagenome-wide association study in nine European countries. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 865-876.	3.0	63
17	Quantitative and qualitative analysis of antimicrobial usage at farm and flock level on 181 broiler farms in nine European countries. <i>Journal of Antimicrobial Chemotherapy</i> , 2019, 74, 798-806.	3.0	45
18	Abundance and diversity of the faecal resistome in slaughter pigs and broilers in nine European countries. <i>Nature Microbiology</i> , 2018, 3, 898-908.	13.3	230

#	ARTICLE	IF	CITATIONS
19	O16-1â€¦ESBL carriage in slaughterhouse workers is associated with occupational exposure. , 2016, , .		0