

# Georgina Milne

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6713644/publications.pdf>

Version: 2024-02-01

14  
papers

228  
citations

1163117

8  
h-index

1058476

14  
g-index

19  
all docs

19  
docs citations

19  
times ranked

250  
citing authors

#	ARTICLE	IF	CITATIONS
1	Bayesian latent class estimation of sensitivity and specificity parameters of diagnostic tests for bovine tuberculosis in chronically infected herds in Northern Ireland. <i>Veterinary Journal</i> , 2018, 238, 15-21.	1.7	67
2	Is There a Relationship Between Bovine Tuberculosis (bTB) Herd Breakdown Risk and Mycobacterium avium subsp. paratuberculosis Status? An Investigation in bTB Chronically and Non-chronically Infected Herds. <i>Frontiers in Veterinary Science</i> , 2019, 6, 30.	2.2	21
3	The influence of pigmentation patterning on bumblebee foraging from flowers of <i>Antirrhinum majus</i> . <i>Die Naturwissenschaften</i> , 2013, 100, 249-256.	1.6	20
4	Spatiotemporal analysis of prolonged and recurrent bovine tuberculosis breakdowns in Northern Irish cattle herds reveals a new infection hotspot. <i>Spatial and Spatio-temporal Epidemiology</i> , 2019, 28, 33-42.	1.7	20
5	Tracing a toad invasion: lack of mitochondrial DNA variation, haplotype origins, and potential distribution of introduced <i>Duttaphrynus melanostictus</i> in Madagascar. <i>Amphibia - Reptilia</i> , 2017, 38, 197-207.	0.5	18
6	Bovine tuberculosis breakdown duration in cattle herds: an investigation of herd, host, pathogen and wildlife risk factors. <i>PeerJ</i> , 2020, 8, e8319.	2.0	18
7	Variation in <i>Mycobacterium bovis</i> genetic richness suggests that inwards cattle movements are a more important source of infection in beef herds than in dairy herds. <i>BMC Microbiology</i> , 2019, 19, 154.	3.3	16
8	Seasonal variation of <i>Fasciola hepatica</i> antibodies in dairy herds in Northern Ireland measured by bulk tank milk ELISA. <i>Parasitology Research</i> , 2018, 117, 2725-2733.	1.6	13
9	Quantifying Land Fragmentation in Northern Irish Cattle Enterprises. <i>Land</i> , 2022, 11, 402.	2.9	9
10	<i>Mycobacterium bovis</i> Population Structure in Cattle and Local Badgers: Co-Localisation and Variation by Farm Type. <i>Pathogens</i> , 2020, 9, 592.	2.8	8
11	Liver fluke ( <i>Fasciola hepatica</i> ) co-infection with bovine tuberculosis in cattle: A prospective herd-level assessment of herd bTB risk in dairy enterprises. <i>Transboundary and Emerging Diseases</i> , 2019, 66, 1727-1736.	3.0	7
12	Quantifying intraherd cattle movement metrics: Implications for disease transmission risk. <i>Preventive Veterinary Medicine</i> , 2020, 185, 105203.	1.9	5
13	Investigating Farm Fragmentation as a Risk Factor for Bovine Tuberculosis in Cattle Herds: A Matched Case-Control Study from Northern Ireland. <i>Pathogens</i> , 2022, 11, 299.	2.8	4
14	Characteristics of Northern Irish cattle herds without bovine tuberculosis infection. <i>Veterinary Record</i> , 2019, 184, 772-772.	0.3	1