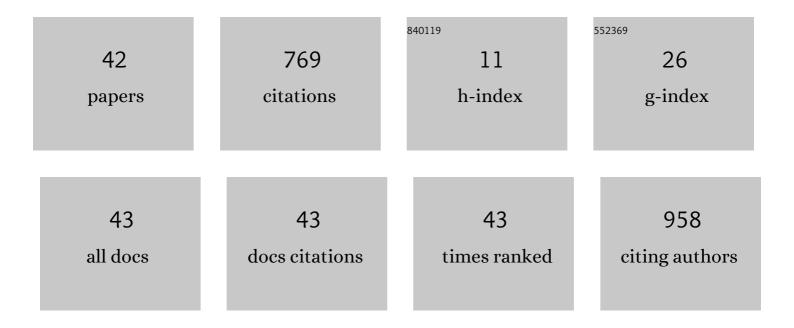
## Takehisa Yamamoto

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6562067/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Subgrouping and analysis of relationships between classical swine fever virus identified during the 2018–2020 epidemic in Japan by a novel approach using shared genomic variants. Transboundary and Emerging Diseases, 2022, 69, 1166-1177.	1.3	5
2	Phylogenetic and phylodynamic analysis of a classical swine fever virus outbreak in Japan (2018–2020). Transboundary and Emerging Diseases, 2022, 69, 1529-1538.	1.3	10
3	Pig farm vaccination against classical swine fever reduces the risk of transmission from wild boar. Preventive Veterinary Medicine, 2022, 198, 105554.	0.7	2
4	Epidemiological analysis of classical swine fever in wild boars in Japan. BMC Veterinary Research, 2021, 17, 188.	0.7	18
5	Genome variability of classical swine fever virus during the 2018–2020 epidemic in Japan. Veterinary Microbiology, 2021, 258, 109128.	0.8	1
6	Epidemiological verification of the mechanism of occurrence of atypical Lâ€ŧype bovine spongiform encephalopathy. Transboundary and Emerging Diseases, 2021, , .	1.3	0
7	Region-wise analysis of dairy cow movements in Japan. BMC Veterinary Research, 2021, 17, 305.	0.7	3
8	Mutations in the tumor suppressor gene p53 in cattle are associated with enzootic bovine leukosis. Veterinary Microbiology, 2021, 263, 109269.	0.8	3
9	Additive Bayesian network analysis of the relationship between bovine respiratory disease and management practices in dairy heifer calves at pre-weaning stage. BMC Veterinary Research, 2021, 17, 360.	0.7	3
10	Estimation of the Lethality Rate, Recovery Rate, and Case Fatality Ratio of Classical Swine Fever in Japanese Wild Boar: An Analysis of the Epidemics From September 2018 to March 2019. Frontiers in Veterinary Science, 2021, 8, 772995.	0.9	2
11	Estimation of infection risk on pig farms in infected wild boar areas—Epidemiological analysis for the reemergence of classical swine fever in Japan in 2018. Preventive Veterinary Medicine, 2020, 175, 104873.	0.7	44
12	Epidemiology of Classical Swine Fever in Japan—A Descriptive Analysis of the Outbreaks in 2018–2019. Frontiers in Veterinary Science, 2020, 7, 573480.	0.9	22
13	Transmission network reconstruction for foot-and-mouth disease outbreaks incorporating farm-level covariates. PLoS ONE, 2020, 15, e0235660.	1.1	11
14	COVID-19 Outbreak and Epidemiological Research in Japan - Part 2 Journal of Veterinary Epidemiology, 2020, 24, 127-133.	0.2	0
15	COVID-19 Outbreak and Epidemiological Researche in Japan - Part 1 Journal of Veterinary Epidemiology, 2020, 24, 21-28.	0.2	0
16	Evaluation of sampling methods for effective detection of infected pig farms during a disease outbreak. PLoS ONE, 2020, 15, e0241177.	1.1	2
17	Wild boars: A potential source ofErysipelothrix rhusiopathiaeinfection in Japan. Microbiology and Immunology, 2019, 63, 465-468.	0.7	9
18	Simultaneous evaluation of diagnostic marker utility for enzootic bovine leukosis. BMC Veterinary Research, 2019, 15, 406,	0.7	5

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19	Reconstructing a transmission network and identifying risk factors of secondary transmissions in the 2010 footâ€andâ€mouth disease outbreak in Japan. Transboundary and Emerging Diseases, 2019, 66, 2074-2086.	1.3	9
20	Control of paratuberculosis: who, why and how. A review of 48 countries. BMC Veterinary Research, 2019, 15, 198.	0.7	219
21	Reconstructing foot-and-mouth disease outbreaks: a methods comparison of transmission network models. Scientific Reports, 2019, 9, 4809.	1.6	32
22	Basic Reproduction Number as a Measure of the Rapidity of the Inter-farm Spread of Porcine Epidemic Diarrhea during the Initial Phase of the Epidemic in Japan in 2013-2014. Journal of Veterinary Epidemiology, 2019, 23, 111-118.	0.2	0
23	Matched case-control study of the influence of inland waters surrounding poultry farms on avian influenza outbreaks in Japan. Scientific Reports, 2018, 8, 3306.	1.6	11
24	Evaluation of fecal shedding and antibody response in dairy cattle infected with paratuberculosis using national surveillance data in Japan. Preventive Veterinary Medicine, 2018, 149, 38-46.	0.7	6
25	Meteorological factors affecting seroconversion of Akabane disease in sentinel calves in the subtropical Okinawa Islands of Japan. Tropical Animal Health and Production, 2018, 50, 209-215.	0.5	6
26	Mathematical modeling of porcine epidemic diarrhea virus dynamics within a farrow-to-finish swine farm to investigate the effects of control measures. Preventive Veterinary Medicine, 2018, 149, 115-124.	0.7	11
27	The effectiveness of colostral antibodies for preventing bovine leukemia virus (BLV) infection in vitro. BMC Veterinary Research, 2018, 14, 419.	0.7	20
28	Genomic Motifs as a Novel Indicator of the Relationship between Strains Isolated from the Epidemic of Porcine Epidemic Diarrhea in 2013-2014. PLoS ONE, 2016, 11, e0147994.	1.1	4
29	Evaluation of the Effect of Missing Data on the Estimation of the Analysis : A Simulation Example Using Epidemiological Survey Data. Journal of Veterinary Epidemiology, 2016, 20, 111-117.	0.2	1
30	Epidemiological analysis of bovine ephemeral fever in 2012–2013 in the subtropical islands of Japan. BMC Veterinary Research, 2016, 12, 47.	0.7	13
31	Fraction of bovine leukemia virus-infected dairy cattle developing enzootic bovine leukosis. Preventive Veterinary Medicine, 2016, 124, 96-101.	0.7	27
32	Source-Related Effects of Wastewater on Transcription Factor (AhR, CAR and PXR)-Mediated Induction of Gene Expression in Cultured Rat Hepatocytes and Their Association with the Prevalence of Antimicrobial-Resistant Escherichia coli. PLoS ONE, 2015, 10, e0138391.	1.1	5
33	Impact of wastewater from different sources on the prevalence of antimicrobial-resistant Escherichia coli in sewage treatment plants in South India. Ecotoxicology and Environmental Safety, 2015, 115, 203-208.	2.9	65
34	Potential risk associated with animal culling and disposal during the foot-and-mouth disease epidemic in Japan in 2010. Research in Veterinary Science, 2015, 102, 228-230.	0.9	8
35	Sampling Strategies in Antimicrobial Resistance Monitoring: Evaluating How Precision and Sensitivity Vary with the Number of Animals Sampled per Farm. PLoS ONE, 2014, 9, e87147.	1.1	6
36	Effectiveness of a short training session for improving pipetting accuracy. Accreditation and Quality Assurance, 2014, 19, 459-463.	0.4	1

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37	Foot-and-Mouth Disease : Infection and Transmission. Journal of Veterinary Epidemiology, 2014, 18, 46-55.	0.2	3
38	The 2010 Foot-and-Mouth Disease Epidemic in Japan. Journal of Veterinary Medical Science, 2012, 74, 399-404.	0.3	117
39	Simulation-based estimation of BSE infection in Japan. Preventive Veterinary Medicine, 2008, 84, 135-151.	0.7	10
40	Evaluation of surveillance strategies for bovine brucellosis in Japan using a simulation model. Preventive Veterinary Medicine, 2008, 86, 57-74.	0.7	28
41	Epidemiologic Indicators Associated with Within-farm Spread of Johne's Disease in Dairy Farms in Japan. Journal of Veterinary Medical Science, 2007, 69, 1255-1258.	0.3	11
42	A quantitative assessment of the risk of exposure to bovine spongiform encephalopathy via meat-and-bone meal in Japan. Preventive Veterinary Medicine, 2006, 75, 221-238.	0.7	16