

# Steven Van Borm

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4054306/publications.pdf>

Version: 2024-02-01

8  
papers

173  
citations

1684188  
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h-index

1588992  
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g-index

8  
all docs

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docs citations

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times ranked

291  
citing authors

#	ARTICLE	IF	CITATIONS
1	Has Epizootic Become Enzootic? Evidence for a Fundamental Change in the Infection Dynamics of Highly Pathogenic Avian Influenza in Europe, 2021. MBio, 2022, 13, .	4.1	64
2	Evaluation of convenient pretreatment protocols for RNA virus metagenomics in serum and tissue samples. Journal of Virological Methods, 2015, 222, 72-80.	2.1	48
3	Next-Generation Sequencing in Veterinary Medicine: How Can the Massive Amount of Information Arising from High-Throughput Technologies Improve Diagnosis, Control, and Management of Infectious Diseases?. Methods in Molecular Biology, 2015, 1247, 415-436.	0.9	33
4	Evaluation of a commercial exogenous internal process control for diagnostic RNA virus metagenomics from different animal clinical samples. Journal of Virological Methods, 2020, 283, 113916.	2.1	10
5	Serological and molecular epidemiology of foot-and-mouth disease viruses in agro-pastoralist livestock herds in the kachia grazing reserve, Nigeria. Transboundary and Emerging Diseases, 2019, 66, 1575-1586.	3.0	7
6	Increased viral read counts and metagenomic full genome characterization of porcine astrovirus 4 and Posavirus 1 in sows in a swine farm with unexplained neonatal piglet diarrhea. Virus Genes, 2020, 56, 696-704.	1.6	4
7	Metagenomic sequencing determines complete infectious bronchitis virus (avian Gamma coronavirus) vaccine strain genomes and associated viromes in chicken clinical samples. Virus Genes, 2021, 57, 529-540.	1.6	4
8	Complete Genome Sequences of Five Foot-and-Mouth Disease Viruses of Serotype A Isolated from Cattle in Nigeria between 2013 and 2015. Genome Announcements, 2018, 6, .	0.8	3