

# Ghislaine Le Gall-ReculÃ©

## List of Publications by Year in descending order

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Version: 2024-02-01

18  
papers

823  
citations

759233

12  
h-index

839539

18  
g-index

18  
all docs

18  
docs citations

18  
times ranked

469  
citing authors

#	ARTICLE	IF	CITATIONS
1	Emergence of a new lagovirus related to rabbit haemorrhagic disease virus. <i>Veterinary Research</i> , 2013, 44, 81.	3.0	180
2	Proposal for a unified classification system and nomenclature of lagoviruses. <i>Journal of General Virology</i> , 2017, 98, 1658-1666.	2.9	148
3	The new French 2010 Rabbit Hemorrhagic Disease Virus causes an RHD-like disease in the Sardinian Cape hare ( <i>Lepus capensis mediterraneus</i> ). <i>Veterinary Research</i> , 2013, 44, 96.	3.0	113
4	Characterisation of a non-pathogenic and non-protective infectious rabbit lagovirus related to RHDV. <i>Virology</i> , 2011, 410, 395-402.	2.4	67
5	Double introduction of highly pathogenic H5N1 avian influenza virus into France in early 2006. <i>Avian Pathology</i> , 2008, 37, 15-23.	2.0	54
6	Large-scale lagovirus disease outbreaks in European brown hares ( <i>Lepus europaeus</i> ) in France caused by RHDV2 strains spatially shared with rabbits ( <i>Oryctolagus cuniculus</i> ). <i>Veterinary Research</i> , 2017, 48, 70.	3.0	54
7	Muscovy duck reovirus $\gamma$ C protein is atypically encoded by the smallest genome segment. <i>Journal of General Virology</i> , 2002, 83, 1189-1200.	2.9	40
8	Recombination at the emergence of the pathogenic rabbit haemorrhagic disease virus <i>Lagovirus europaeus</i> /GI.2. <i>Scientific Reports</i> , 2020, 10, 14502.	3.3	36
9	Emergence of Pathogenicity in Lagoviruses: Evolution from Pre-existing Nonpathogenic Strains or through a Species Jump?. <i>PLoS Pathogens</i> , 2015, 11, e1005087.	4.7	31
10	Molecular evolution and antigenic variation of European brown hare syndrome virus (EBHSV). <i>Virology</i> , 2014, 468-470, 104-112.	2.4	21
11	Immunocapture-RT-PCR assay for detection and molecular epidemiology studies of Rabbit Haemorrhagic Disease and European Brown Hare Syndrome viruses. <i>Journal of Virological Methods</i> , 2001, 97, 49-57.	2.1	18
12	Complete coding sequences of European brown hare syndrome virus (EBHSV) strains isolated in 1982 in Sweden. <i>Archives of Virology</i> , 2013, 158, 2193-2196.	2.1	14
13	First Complete Genome Sequence of a Hare Calicivirus Strain Isolated from <i>Lepus europaeus</i> . <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.6	12
14	First complete genome sequence of a European non-pathogenic rabbit calicivirus (lagovirus GI.3). <i>Archives of Virology</i> , 2018, 163, 2921-2924.	2.1	12
15	Retrospective Analysis Shows That Most RHDV GI.1 Strains Circulating Since the Late 1990s in France and Sweden Were Recombinant GI.3/GI.1d Strains. <i>Genes</i> , 2020, 11, 910.	2.4	12
16	Genetic diversity and evolution of Hare Calicivirus (HaCV), a recently identified lagovirus from <i>Lepus europaeus</i> . <i>Infection, Genetics and Evolution</i> , 2020, 82, 104310.	2.3	6
17	Rabbit haemorrhagic disease virus <i>Lagovirus europaeus</i> /GI.1d strain: genome sequencing, in vivo virus replication kinetics, and viral dose effect. <i>BMC Veterinary Research</i> , 2021, 17, 257.	1.9	3
18	Development and Evaluation of a Duplex Lateral Flow Assay for the Detection and Differentiation between Rabbit Haemorrhagic Disease Virus <i>Lagovirus europaeus</i> /GI.1 and /GI.2. <i>Biology</i> , 2022, 11, 401.	2.8	2