

Identification of a New Genotype of African Swine Fever Ethiopia

Transboundary and Emerging Diseases

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Citation Report

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1	Polymerase cross-linking spiral reaction (PCLSR) for detection of African swine fever virus (ASFV) in pigs and wild boars. <i>Scientific Reports</i> , 2017, 7, 42903.	3.3	42
2	BA711 ^{CD2} : a New Recombinant Live Attenuated African Swine Fever Virus with Cross-Protective Capabilities. <i>Journal of Virology</i> , 2017, 91, .	3.4	189
3	Virulent strain of African swine fever virus eclipses its attenuated derivative after challenge. <i>Archives of Virology</i> , 2017, 162, 3081-3088.	2.1	10
4	African swine fever (ASF) and ticks. No risk of tick-mediated ASF spread in Poland and Baltic states. <i>Journal of Veterinary Research (Poland)</i> , 2017, 61, 375-380.	1.0	37
5	Genetic Assessment of African Swine Fever Isolates Involved in Outbreaks in the Democratic Republic of Congo between 2005 and 2012 Reveals Co-Circulation of p72 Genotypes I, IX and XIV, Including 19 Variants. <i>Viruses</i> , 2017, 9, 31.	3.3	40
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7	The Epidemiology of African Swine Fever in “Nonendemic” Regions of Zambia (1989–2015): Implications for Disease Prevention and Control. <i>Viruses</i> , 2017, 9, 236.	3.3	33
8	Chimeric DNA/LNA-based biosensor for the rapid detection of African swine fever virus. <i>Talanta</i> , 2018, 184, 35-41.	5.5	25
9	Linear epitopes in African swine fever virus p72 recognized by monoclonal antibodies prepared against baculovirus-expressed antigen. <i>Journal of Veterinary Diagnostic Investigation</i> , 2018, 30, 406-412.	1.1	34
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11	Co-circulation of multiple genotypes of African swine fever viruses among domestic pigs in Zambia (2013-2015). <i>Transboundary and Emerging Diseases</i> , 2018, 65, 114-122.	3.0	36
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13	Multi-locus sequence typing of African swine fever viruses from endemic regions of Kenya and Eastern Uganda (2011–2013) reveals rapid B602L central variable region evolution. <i>Virus Genes</i> , 2018, 54, 111-123.	1.6	29
14	Genome Sequences of Five African Swine Fever Virus Genotype IX Isolates from Domestic Pigs in Uganda. <i>Microbiology Resource Announcements</i> , 2018, 7, .	0.6	26
15	African swine fever virus does not express viral microRNAs in experimentally infected pigs. <i>BMC Veterinary Research</i> , 2018, 14, 268.	1.9	10
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18	Phylodynamics and evolutionary epidemiology of African swine fever p72-CVR genes in Eurasia and Africa. <i>PLoS ONE</i> , 2018, 13, e0192565.	2.5	44

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31	A Simple Method for Sample Preparation to Facilitate Efficient Whole-Genome Sequencing of African Swine Fever Virus. <i>Viruses</i> , 2019, 11, 1129.	3.3	35
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#	ARTICLE	IF	CITATIONS
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