

Origins and evolutionary genomics of the 2009 swine-o

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Citation Report

#	ARTICLE	IF	CITATIONS
4	Effective Containment of Pandemic Influenza (H1N1 2009) Virus Outbreak in an International Residence Hall of a University by Recommended Infection Control Guidance. <i>Infection and Chemotherapy</i> , 2009, 42, 99.	1.0	1
6	Genomic Signatures of Influenza A Pandemic (H1N1) 2009 Virus. <i>Emerging Infectious Diseases</i> , 2009, 15, 1897-1903.	2.0	69
7	2009 Swine-Origin Influenza A (H1N1) Resembles Previous Influenza Isolates. <i>PLoS ONE</i> , 2009, 4, e6402.	1.1	31
8	Evolutionary Trends of A(H1N1) Influenza Virus Hemagglutinin Since 1918. <i>PLoS ONE</i> , 2009, 4, e7789.	1.1	71
9	New methods for analyzing serological data with applications to influenza surveillance. <i>Nature Precedings</i> , 2009, , .	0.1	0
10	Insights into the Evolutionary History of an Emerging Livestock Pathogen: Porcine Circovirus 2. <i>Journal of Virology</i> , 2009, 83, 12813-12821.	1.5	208
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12	Poor Clinical Sensitivity of Rapid Antigen Test for Influenza A Pandemic (H1N1) 2009 Virus. <i>Emerging Infectious Diseases</i> , 2009, 15, 1662-1664.	2.0	167
13	Seasonal Influenza Virus Species in Patient Swab Samples Analyzed for the Presence of the Pandemic (H1N1) 2009 Influenza Virus. <i>Journal of Clinical Microbiology</i> , 2009, 47, 4187-4188.	1.8	1
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22	The Role of Genomics in Tracking the Evolution of Influenza A Virus. <i>PLoS Pathogens</i> , 2009, 5, e1000566.	2.1	81

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