Accepted Manuscript

Molecular epidemiological survey and complete genomic phylogenetic analysis of H6 subtype avian influenza viruses in poultry in China from 2011 to 2016

Cheng Peng, Hongtao Sun, Jinping Li, Guangyu Hou, Suchun Wang, Shuo Liu, Qingye Zhuang, Shanju Cheng, Jiming Chen, Wenming Jiang

PII: S1567-1348(18)30527-6

DOI: doi:10.1016/j.meegid.2018.07.023

Reference: MEEGID 3595

To appear in: Infection, Genetics and Evolution

Received date: 29 March 2018 Revised date: 11 July 2018 Accepted date: 19 July 2018

Please cite this article as: Cheng Peng, Hongtao Sun, Jinping Li, Guangyu Hou, Suchun Wang, Shuo Liu, Qingye Zhuang, Shanju Cheng, Jiming Chen, Wenming Jiang, Molecular epidemiological survey and complete genomic phylogenetic analysis of H6 subtype avian influenza viruses in poultry in China from 2011 to 2016. Meegid (2018), doi:10.1016/j.meegid.2018.07.023

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



ACCEPTED MANUSCRIPT

Molecular epidemiological survey and complete genomic phylogenetic analysis of H6 subtype avian influenza viruses in poultry in China from 2011–2016

Cheng Peng¹, Hongtao Sun¹, Jinping Li, Guangyu Hou, Suchun Wang, Shuo Liu, Qingye Zhuang, Shanju Cheng, Jiming Chen, Wenming Jiang*

China Animal Health and Epidemiology Center, Qingdao, China

¹ These authors contributed equally to this work

^{*}Corresponding author. E-mail address: jiangwenming@cahec.cn

Download English Version:

https://daneshyari.com/en/article/8646483

Download Persian Version:

https://daneshyari.com/article/8646483

<u>Daneshyari.com</u>