

ERRATUM

Molecular Evolution Analysis and Geographic Investigation of Severe Acute Respiratory Syndrome Coronavirus-Like Virus in Palm Civets at an Animal Market and on Farms

Biao Kan, Ming Wang, Huaiqi Jing, Huifang Xu, Xiugao Jiang, Meiyang Yan, Weili Liang, Han Zheng, Kanglin Wan, Qiyong Liu, Buyun Cui, Yanmei Xu, Enmin Zhang, Hongxia Wang, Jingrong Ye, Guichang Li, Machao Li, Zhigang Cui, Xiaobao Qi, Kai Chen, Lin Du, Kai Gao, Yu-teng Zhao, Xiao-zhong Zou, Yue-Ju Feng, Yu-Fan Gao, Rong Hai, Dongzhen Yu, Yi Guan, and Jianguo Xu

State Key Laboratory for Infectious Disease Prevention and Control (China CDC) and National Institute for Communicable Disease Control and Prevention, Chinese Center for Disease Control and Prevention, P.O. Box 5, Changping, Beijing 102206, People's Republic of China; Guangzhou Center for Disease Control and Prevention, 23 Third Zhongshan Road, Guangzhou, Guangdong 510080, People's Republic of China; and Department of Microbiology, The University of Hong Kong, Hong Kong Special Administrative Region, People's Republic of China

Volume 79, no. 18, p. 11892–11900, 2005. Page 11894, Table 1: The GenBank accession numbers of the indicated strains should read as follows: C029G, AY687372; B033G, AY687362; B040G, AY687364; C017G, AY687367; B039G, AY687363; A013G, AY687355; C019G, AY687369; C014G, AY687366; C028G, AY687371; GD03T0013, AY525636; B012G, AY687359; and SZ3, AY304486.

Page 11895, Fig. 3: The amino acid residues at position 227 of viruses SZ3 and SZ16 should read “K” instead of “N,” and the residues at position 743 of viruses SZ3 and SZ16 should read “A” instead of “T.”

Page 11899, column 1, line 11: “at residues 147, 228, 240, and 479” should read “at residues 227, 243, 462, and 479.” Lines 14 and 15: “(positions 609, 462, 360, 472, 480, 487, 613, 665, 743, 765, and 1163” should read “(positions 139, 609, 360, 472, 480, 487, 613, 665, 743, 765, and 1163.” Line 17: “positions 344, 778, 227, and 244” should read “positions 77, 344, 778, and 244.”