

Correction

APPLIED BIOLOGICAL SCIENCES

Correction for “Genome sequence of the Asian Tiger mosquito, *Aedes albopictus*, reveals insights into its biology, genetics, and evolution,” by Xiao-Guang Chen, Xuanting Jiang, Jinbao Gu, Meng Xu, Yang Wu, Yuhua Deng, Chi Zhang, Mariangela Bonizzoni, Wannes Dermauw, John Vontas, Peter Armbruster, Xin Huang, Yulan Yang, Hao Zhang, Weiming He, Hongjuan Peng, Yongfeng Liu, Kun Wu, Jiahua Chen, Manolis Lirakis, Pantelis Topalis, Thomas Van Leeuwen, Andrew Brantley Hall, Xiaofang Jiang, Chevon Thorpe, Rachel Lockridge Mueller, Cheng Sun, Robert Michael Waterhouse, Guiyun Yan, Zhijian Jake Tu, Xiaodong Fang, and Anthony A. James, which appeared in issue 44, November 3, 2015, of *Proc Natl Acad Sci USA* (112:E5907–E5915; first published October 19, 2015; 10.1073/pnas.1516410112).

The authors note that the affiliation for Thomas Van Leeuwen should instead appear as “^cLaboratory of Agrozoology, Department of Crop Protection, Faculty of Bioscience Engineering, Ghent University, B-9000 Ghent, Belgium; and ¹Institute for Biodiversity and Ecosystem Dynamics, University of Amsterdam, 1090 GE Amsterdam, The Netherlands.” The corrected author and affiliation lines appear below. The online version has been corrected.

Xiao-Guang Chen^{a,1}, Xuanting Jiang^b, Jinbao Gu^a, Meng Xu^b, Yang Wu^a, Yuhua Deng^a, Chi Zhang^b, Mariangela Bonizzoni^{c,d}, Wannes Dermauw^e, John Vontas^{f,g}, Peter Armbruster^h, Xin Huang^h, Yulan Yang^b, Hao Zhang^a, Weiming He^b, Hongjuan Peng^a, Yongfeng Liu^b, Kun Wu^a, Jiahua Chen^b, Manolis Lirakisⁱ, Pantelis Topalis^f, Thomas Van Leeuwen^{e,j}, Andrew Brantley Hall^{k,l}, Xiaofang Jiang^{k,l}, Chevon Thorpe^m, Rachel Lockridge Muellerⁿ, Cheng Sunⁿ, Robert Michael Waterhouse^{o,p,q,r}, Guiyun Yan^{a,c}, Zhijian Jake Tu^{k,l}, Xiaodong Fang^{b,1}, and Anthony A. James^{s,1}

^aDepartment of Pathogen Biology, School of Public Health and Tropical Medicine, Southern Medical University, Guangzhou 510515, China; ^bBeijing Genomics Institute-Shenzhen, Shenzhen 518083, China; ^cProgram in Public Health, University of California, Irvine, CA 92697; ^dDepartment of Biology and Biotechnology, University of Pavia, 27100 Pavia, Italy; ^eLaboratory of Agrozoology, Department of Crop Protection, Faculty of Bioscience Engineering, Ghent University, B-9000 Ghent, Belgium; ^fInstitute of Molecular Biology and Biotechnology, Foundation for Research and Technology–Hellas, 73100 Heraklion, Greece; ^gFaculty of Crop Science, Pesticide Science Lab, Agricultural University of Athens, 11855 Athens, Greece; ^hDepartment of Biology, Georgetown University, Washington, DC 20057; ⁱDepartment of Biology, University of Crete, Heraklion, GR-74100, Crete, Greece; ^jInstitute for Biodiversity and Ecosystem Dynamics, University of Amsterdam, 1090 GE Amsterdam, The Netherlands; ^kInterdisciplinary PhD Program in Genetics, Bioinformatics, and Computational Biology, Virginia Tech University, Blacksburg, VA 24061; ^lDepartment of Biochemistry, Fralin Life Science Institute, Virginia Tech University, Blacksburg, VA 24061; ^mCellular and Molecular Physiology, Edward Via College of Osteopathic Medicine, Blacksburg, VA 24060; ⁿDepartment of Biology, Colorado State University, Fort Collins, CO 80523; ^oDepartment of Genetic Medicine and Development, University of Geneva Medical School, 1211 Geneva, Switzerland; ^pSwiss Institute of Bioinformatics, 1211 Geneva, Switzerland; ^qComputer Science and Artificial Intelligence Laboratory, Massachusetts Institute of Technology, Cambridge, MA 02139; ^rThe Broad Institute of MIT and Harvard, Cambridge, MA 02142; and ^sDepartments of Microbiology & Molecular Genetics and Molecular Biology & Biochemistry, University of California, Irvine, CA 92697

www.pnas.org/cgi/doi/10.1073/pnas.1524968113