



Contents lists available at [ScienceDirect](#)

Vaccine

journal homepage: www.elsevier.com/locate/vaccine



Corrigendum

Corrigendum to “A computationally designed H5 antigen shows immunological breadth of coverage and protects against drifting avian strains” [Vaccine 37 (2019) 2369–2376]



Ted M. Ross^a, Joshua DiNapoli^b, Maryann Giel-Moloney^{b,*}, Chalise E. Bloom^a, Kateri Bertran^c, Charles Balzli^c, Tod Strugnell^b, Raymond P. Oomen^b, Mariana Sá e Silva^e, Teshome Mebatsion^e, Michel Bublot^d, David E. Swayne^c, Harry Kleanthous^b

^aUniversity of Georgia, Center for Vaccines and Immunology, Department of Infectious Diseases, Athens, GA 30602, USA

^bSanofi-Pasteur, 38 Sidney Street, Cambridge, MA 02139, USA

^cExotic and Emerging Avian Viral Diseases Research Unit, Southeast Poultry Research Laboratory, U.S. National Poultry Research Center, Agricultural Research Service, U.S. Department of Agriculture, Athens, GA 30602, USA

^dBoehringer Ingelheim, S.A.S., R&D, 69007 Lyon, France

^eBoehringer Ingelheim, R&D, Athens, GA 30601, USA

The authors regret that Ray Oomen, a co-author to the paper was mistakenly left out of the submitted manuscript accepted for publication on March 12, 2019. This was an honest oversight by the Sanofi Pasteur's co-authors.

The author information has been updated as above.

The authors would like to apologize for any inconvenience caused.

DOI of original article: <https://doi.org/10.1016/j.vaccine.2019.03.018>

* Corresponding author.

E-mail address: Maryann.Giel-Moloney@sanofi.com (M. Giel-Moloney).

<https://doi.org/10.1016/j.vaccine.2019.07.058>

0264-410X/© 2019 Elsevier Ltd. All rights reserved.