

CORRECTION

Open Access



Correction to: Identifying high risk subgroups of MSM: a latent class analysis using two samples

M. Kumi Smith^{1*}, Gabriella Stein², Weibin Cheng³, William C. Miller⁴ and Joseph D. Tucker⁵

Correction to: *BMC Infect Dis* (2019) 19:213
<https://doi.org/10.1186/s12879-019-3700-5>

Following publication of the original article [1], the author reported his family name has been marked as the first name. His given name is M. Kumi and his family name is Smith.

The original article has been corrected as well.

The publisher apologizes for any inconvenience caused by this error.

Author details

¹Division of Epidemiology & Community Health, School of Public Health, University of Minnesota Twin Cities, 1300 South 2nd Street, Minneapolis, MN 55454, USA. ²Department of Biostatistics, University of North Carolina at Chapel Hill, 135 Dauer Drive, 3101 McGavran-Greenberg Hall, CB #7420, Chapel Hill, NC 27599, USA. ³Department of AIDS/STD Control and Prevention, Guangzhou Center for Disease Control and Prevention, 1 Jiaochang E Rd., Guangzhou Shi 510000, Guangdong Sheng, China. ⁴Division of Epidemiology, The Ohio State University, College of Public Health, 1841 Neal, Ave., 302 Cunz Hall, Columbus, OH 43210, USA. ⁵Division of Infectious Diseases, University of North Carolina at Chapel Hill, 130 Mason Farm Road, 2nd Floor, Chapel Hill, NC 27599, USA.

Received: 7 March 2019 Accepted: 7 March 2019

Published online: 26 March 2019

Reference

1. Smith MK, et al. Identifying high risk subgroups of MSM: a latent class analysis using two samples. *BMC Infect Dis*. 2019;19:213. <https://doi.org/10.1186/s12879-019-3700-5>.

* Correspondence: smi00831@umn.edu

¹Division of Epidemiology & Community Health, School of Public Health, University of Minnesota Twin Cities, 1300 South 2nd Street, Minneapolis, MN 55454, USA

Full list of author information is available at the end of the article

