

Corrigendum to: Suboptimal Biological Sampling as a Probable Cause of False-Negative COVID-19 Diagnostic Test Results

Natalie N. Kinloch,^{1,2} Gordon Ritchie,^{3,4} Chanson J. Brumme,^{2,5} Winnie Dong,² Weiyan Dong,² Tanya Lawson,³ R. Brad Jones,⁶ Julio S.G. Montaner,^{2,5} Victor Leung,^{3,4} Marc G. Romney,^{3,4} Aleksandra Stefanovic,^{3,4} Nancy Matic,^{3,4} Christopher F. Lowe,^{3,4} and Zabrina L. Brumme^{1,2}

¹Faculty of Health Sciences, Simon Fraser University, Burnaby, Canada, ²British Columbia Centre for Excellence in HIV/AIDS, Vancouver, Canada, ³Division of Medical Microbiology and Virology, St. Paul's Hospital, Vancouver, Canada, ⁴Department of Pathology and Laboratory Medicine, University of British Columbia, Vancouver, Canada, ⁵Department of Medicine, University of British Columbia, Vancouver, Canada, ⁶Infectious Diseases Division, Department of Medicine, Weill Cornell Medical College, New York, USA

In “Suboptimal Biological Sampling as a Probable Cause of False-Negative COVID-19 Diagnostic Test Results” by Kinloch et al [J Infect Dis 2020; doi: 10.1093/infdis/jiaa370], there was a typographical error in one of the primer sequences in the Methods

section. The sequence for the RPP30-Shear Forward Primer was reported as CCAATTTGCTGCTCCTTGGG, but the correct sequence for this primer is CCATTTGCTGCTCCTTGGG (less one A at position 3). The authors regret the error.