

Supplemental Table 1. Number of positive swabs detected at each time point, parsed by swab type. A swab was considered to be positive if more than 10 reads mapped to more than 100 bases of the scaffolded reference genome with greater than 95% identity.

Date	Body swab	Head swab	Oral swab	Rectal swab
April 2016	2	3	3	3
July 2016			1	3
October 2016	18	18	6	13
January 2017	1			
September 2017			1	