

Table S1. Summary of quality sequences recovered from human subjects.

	Swab	Scrape	Punch	Total
Subject 1R	163	176	186	525
Subject 1L	189	182	186	557
Subject 2R	159	186	180	525
Subject 2L	191	161	186	538
Subject 3R	164	166	193	523
Subject 3L	181	164	185	530
Subject 4R	193	187	185	565
Subject 4L	175	173	185	533
Subject 5R	190	150	187	527
Subject 5L	191	171	188	550