Supplemental Table 1	Subject	Buccal mucosa microbiota: Number of sequences (after processing)	Tongue (dorsal side) microbiota: Number of sequences (after processing)	Nasal cavity microbiota: Number of sequences (after processing)	Age	Gender	Race	Current Smoker	Former Smoker	Second Hand Smoker Exposure	Alcoholic Beverages w/in past yr	Diabete	Lung Disease	Inhaler use	Anitbiotics use	Oral Steroid use	Inhaled Steroid Use	Nasal Rinse Use	Nasal Spray use	Current Upper respiratory infection	Current Lower respiratory infection	Current gum/dental infection
Subjects included in our intial analysis	A	1684	1597	269	38	male	Asian	No	No	No	<1/month	No	No	No	No	No	No	No	No	No	No	No
	B	1017	719	382	27	male	White	No	No	No	2-3/week	No	No	No	No	No	No	No	No	No	No	No
	C	912	599	1115	37	female	White	No	No	No	<1/month	No	No	No	No	No	Yes	Yes	Yes	No	No	No
	D	833	1195	1595	34	female	White	No	No	No	never	No	No	No	No	No	No	No	No	No	No	No
	E	1163	942	1431	54	female	White	Yes	n/a	Yes	<1/month	No	No	No	No	No	No	No	No	No	No	No
	F	1549	1255	939	36	female	White	No	No	No	<1/month	No	No	No	No	No	No	No	No	No	No	No
	G	715	962	778	30	female	White	Yes	n/a	No	2-4/month	No	No	No	No	No	No	No	No	No	No	No
	H	1023	519	781	54	male	Asian	No	No	No	never	No	No	No	No	No	No	No	Yes	No	No	No
	I	1199	1397	945	29	female	White	No	No	No	2-4/month	No	No	No	No	No	No	Yes	No	No	Yes	No
	J	1187	1041	911	31	male	White	No	No	Yes	4 or more/week	No	No	No	No	No	No	No	No	No	No	No
	K	1305	1301	not sequenced due to insufficient amplification	36	male	Black	No	No	Yes	4 or more/week	No	No	No	No	No	No	Yes	No	No	No	No
	L	788	1093	1	59	female	White	No	No	No	never	No	No	No	No	No	No	No	No	Yes	No	No
		Sequences generated from sff submitted to SRA (PRJNA248297)																				
				sff not submitted																		