

Supplemental Table [24](#)

Patient #	Usable Samples	Usable Sequences
03	B1, B2, D1, D2, R	8,819
04	B1, B2, D2	7,578
28	B1, B2, D1, D2, R	18,486
29	B1, B2	10,299
30	B1, B2, D1, D2, R	14,448
31	B1, B2, D1, D2, R	6,571
32	B1, B2, D1, D2, R1, R2	14,838
33	B1, B2, D1, D2, R	8,933
34	B1, B2, D1, D2, R1, R2	9,887
36	B1, B2, D1, D2, R	13,321
37	B1, B2, D1, D2, R1, R2	14,970
38	B1, B2, D1, D2, R	13,763
39	B1, B2, D1, D2, R1, R2	20,698
41	B1, B2, D1, D2, R	24,894
42	B1, B2, D1, D2, R	7,366
Total		194,871

Patients	15
Samples	74
Sequences Generated	213,375
Usable Sequences	194,871
OTUs at 97%	4,857

Table 42. Subject and sample descriptive statistics.

Usable sequences met quality standards: no Ns in sequence, exact match to 5' primer, exact match to barcode

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tag, no low quality sequence beyond 5' primer. Sequences that passed quality standards and that had a length 180 nt to 280 nt, after quality trimming, were retained.

