Table S4: Mixed infection and lineage data from cohort one

ID	Fg/uL	Composite target coverage breadth %	Average coverage depth (x)	Species best match (main population (%)/min population (%))	SNP-based lineage (main (%)/min (%) if mixed)	Potential mixed infection
1	8.807	100	3052.72	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
2	1460.08	100	5898.61	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
3	137.357	99.99	2781.11	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
4	7.01	99.99	2052.28	Mycobacterium tuberculosis complex : 83.8 / Mycobacterium seoulense or Mycobacterium branderi : 15.7	Other than H37Rv	n/a
5	205.01	100	3556.78	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
6	623.22	99.99	2951.18	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
7	44.25	99.98	687.04	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
8	28.54	100	3181.43	Mycobacterium tuberculosis complex	Lineage 2 (M. tuberculosis)	n/a
9	42.36	99.99	3631.57	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
10	45.24	99.86	2866.22	Mycobacterium tuberculosis complex	Lineage 1 (M. tuberculosis)	n/a
11	51.172	99.94	2438.46	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
12	10.17	100	3774.00	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
13	18.71	99.98	570.04	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	Yes
14	30.71	99.96	1253.46	Mycobacterium tuberculosis complex	Lineage 2 (M. tuberculosis)	n/a
15	42.22	99.98	2866.14	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
16	9.83	99.99	1343.27	Mycobacterium tuberculosis complex : 92.3 / Mycobacterium tuberculosis complex : 7.7	Lineage 4.3 (M. tuberculosis)	n/a
17	57.67	99.79	413.57	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
18	0.03	99.98	894.44	Mycobacterium tuberculosis complex : 93.4 / Mycobacterium parmense : 6.6	Lineage 4.3 (M. tuberculosis)	Yes
19	2.39	99.99	1859.3	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
20	6.088	99.98	2586.46	Mycobacterium tuberculosis complex	Lineage 2 (M. tuberculosis)	n/a
21	36.08	99.94	578.95	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
22	11.03	99.98	1205.79	Mycobacterium tuberculosis complex	Lineage 4.3 (M. tuberculosis)	n/a
23	243.37	99.99	5221.00	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
24	0.1	99.84	188.34	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
25	0.009	99.90	307.05	Mycobacterium tuberculosis complex	Other than H37Rv : 100.0% Lineage 4.3 (M. tuberculosis) : 4.88%	Yes
26	1.112	99.98	2449.33	Mycobacterium tuberculosis complex	Other than H37Rv	n/a
27	5127.15	100	5236.54	Mycobacterium tuberculosis complex	Lineage 1 (M. tuberculosis)	n/a
28	2834	99.99	2070.04	Mycobacterium tuberculosis complex	Other than H37Rv	n/a