

## Additional file 2: Recovery of complete plasmid sequence directly from clinical specimens

ID	Plasmid copies/ $\mu$ l	Reads mapping to <i>C. trachomatis</i> plasmid	Mean read depth	Coverage of reference <i>C. trachomatis</i> plasmid
CT-33	4,350,790	37.91%	41945	100%
CT-34	45,417	2.31%	3727	100%
CT-35	16,948	0.20%	147	100%
CT-36	80,826	1.06%	797	100%
CT-37	54,154	0.80%	642	100%
CT-38	4,698	0.34%	301	100%
CT-39	4,056	0.12%	104	100%
CT-40	22,388	4.08%	5822	100%
CT-41	2,367	0.11%	91	100%
CT-42	5,093	0.30%	277	100%