

Table S1. Summary of RNA-Seq data for 6 groups of *T. asahii* biofilms

Groups -samples	Raw reads	raw bases (G)	Q20	Q30	Clean reads	Clean bases (G)
O-1	23,387,960	7.02	94.38%	90.39%	20,369,889	6.11
O-2	19,680,659	5.90	97.25%	92.25%	19,551,578	5.87
O-3	18,349,700	5.50	97.33%	92.42%	18,221,396	5.47
D-1	22,474,875	6.74	97.33%	92.62%	22,335,665	6.70
D-2	22,262,696	6.68	97.71%	93.52%	22,143,828	6.64
D-3	20,361,274	6.11	97.51%	93.03%	20,244,903	6.07
L20-1	20,013,410	6.00	97.67%	93.45%	19,921,988	5.98
L20-2	20,411,827	6.12	97.70%	93.43%	20,323,104	6.10
L20-3	21,406,905	6.42	97.72%	93.54%	21,309,440	6.39
L40-1	19,099,066	5.73	97.69%	93.51%	19,000,708	5.70
L40-2	20,003,785	6.00	97.64%	93.38%	19,918,940	5.98
L40-3	18,742,820	5.62	97.51%	92.99%	18,643,048	5.59
A-1	30,145,398	9.04	96.04%	92.52%	27,697,165	8.31
A-2	20,869,264	6.26	97.28%	92.27%	20,751,677	6.23
A-3	21,088,855	6.33	97.49%	92.73%	20,990,366	6.30
B-1	28,086,726	8.43	96.92%	93.48%	26,748,616	8.02
B-2	21,785,469	6.54	97.49%	92.57%	21,680,472	6.50
B-3	19,836,180	5.95	97.42%	92.60%	19,706,995	5.91