

Supplementary Table 1. Sequencing statistics

Sample Description	DNA/RNA library	% Reads Passing Quality Control	# of Raw Reads	# of Reads after Quality Filtering and Adapter Trimming	# of Reads Aligned to Human / Primate	# of Non-Human / Primate Reads	% Human / Primate Reads	# of Reads Aligned to NCBI NT database
No Template Control (NTC)	DNA	84.3	6,664,047	5,619,424	539,093	5,080,331	9.6	3,848,762
No Template Control (NTC)	RNA	59.1	13,243,517	7,829,946	2,174,615	5,655,331	27.8	3,065,872
External Positive Control (PC)*	DNA	62.9	15,488,998	9,735,493	9,154,273	581,220	94.0	469,147
External Positive Control (PC)*	RNA	64.8	6,841,850	4,433,299	1,550,481	2,882,818	35.0	1,616,732
Analytical Sample	DNA	67.8	23,638,587	16,037,745	15,999,078	38,667	99.8	24,156
Analytical Sample	RNA	80.8	9,161,626	7,400,632	5,635,904	1,764,728	76.2	889,944

*7 organism positive control mixture of CMV, HIV, *Streptococcus agalactiae*, *Klebsiella pneumoniae*, *Cryptococcus neoformans*, *Aspergillus niger*, and *Toxoplasma gondii*

Abbreviations: NTC, no template control; PC, positive control; NCBI, National Center for Biotechnology Information, NT, nucleotide