

Run Type	Configuration	Assay	Number of adapters multiplexed	Total number of reads	Number of reads pass filter	% reads passed filter	% PhiX	%Q30
MiSeq	2 x 250 PE	AMPSeq	21	3.22E+06	3.10E+06	96.27	31.22	93.4
MiSeq	1 x 50 SE	DNaseq	24	1.87E+07	1.71E+07	91.44	4.83	96.8
HiSeq	1 x 50 SE	RNAseq	24	2.16E+08	1.93E+08	89.35	4.79	93.7
HiSeq	1 x 50 SE	RNAseq	24	2.01E+08	1.89E+08	94.03	5.39	96.2
NextSeq	1 x 150 SE	RNAseq	15	1.30E+08	1.12E+08	86.15	0.77	89.3
NextSeq	1 x 75 SE	RNAseq	24	4.90E+08	4.07E+08	83.06	3.75	93.3
NextSeq	1 x 150 SE	RNAseq	24	1.00E+08	9.38E+07	93.80	14.63	86.7