

Cluster Generation and Sequencing	MiSeq Reagent Kit v2				MiSeq Reagent Kit v2 Micro	MiSeq Reagent Kit v2 Nano		MiSeq Reagent Kit v3	
Read Length	1 × 36 bp	2 × 25 bp	2 × 150 bp	2 × 250 bp	2 × 150 bp	2 × 250 bp	2 × 150 bp	2 × 75 bp	2 × 300 bp
Output	540–610 Mb	750–850 Mb	4.5–5.1 Gb	7.5–8.5 Gb	1.2 Gb	500 Mb	300 Mb	3.3–3.8 Gb	13.2–15 Gb
Reads Passing Filter Per Flow Cell									
Single-end Reads (Clusters)	12-15 M				4 M	1 M		22-25 M	
Quality Scores* (average across the entire run)									
	> 90% bases higher than Q30 at 1 × 36 bp							> 85% bases higher than Q30 at 2 × 75 bp	
	> 90% bases higher than Q30 at 2 × 25 bp							> 70% bases higher than Q30 at 2 × 300 bp	
	> 80% bases higher than Q30 at 2 × 150 bp								
	> 75% bases higher than Q30 at 2 × 250 bp								
*Specifications based on Illumina PhiX control library at supported cluster densities. Performance may vary based on library type and quality, insert size, loading concentration, and other experimental factors.									