

Library type	Library fragment size	Read size	# of reads (after filtering)
PE ^{1,2}	550 bp	301 bp	43,347,614
PE ^{1,3}	350 bp	101 bp	31,177,878
PE ²	500 bp	301 bp	34,566,750
MP ^{1,2}	4 kb	301 bp	6,999,466
MP ²	6 kb	301 bp	3,761,466
MP ²	10 kb	301 bp	3,401,790
PACBIO	/	3.3 kb ⁴	1,390,999 ⁵

¹Libraries produced and sequenced by [40].

² Sequenced in a MiSeq Illumina sequencer (Illumina, California, United States)

³ Sequenced in a HiSeq2000 Illumina sequencer (Illumina, California, United States)

⁴ mean subread length

⁵ no filtering