

Supplementary Materials

Uncovering missed indels by leveraging unmapped reads

Mohammad Shabbir Hasan¹, Xiaowei Wu², and Liqing Zhang^{1*}

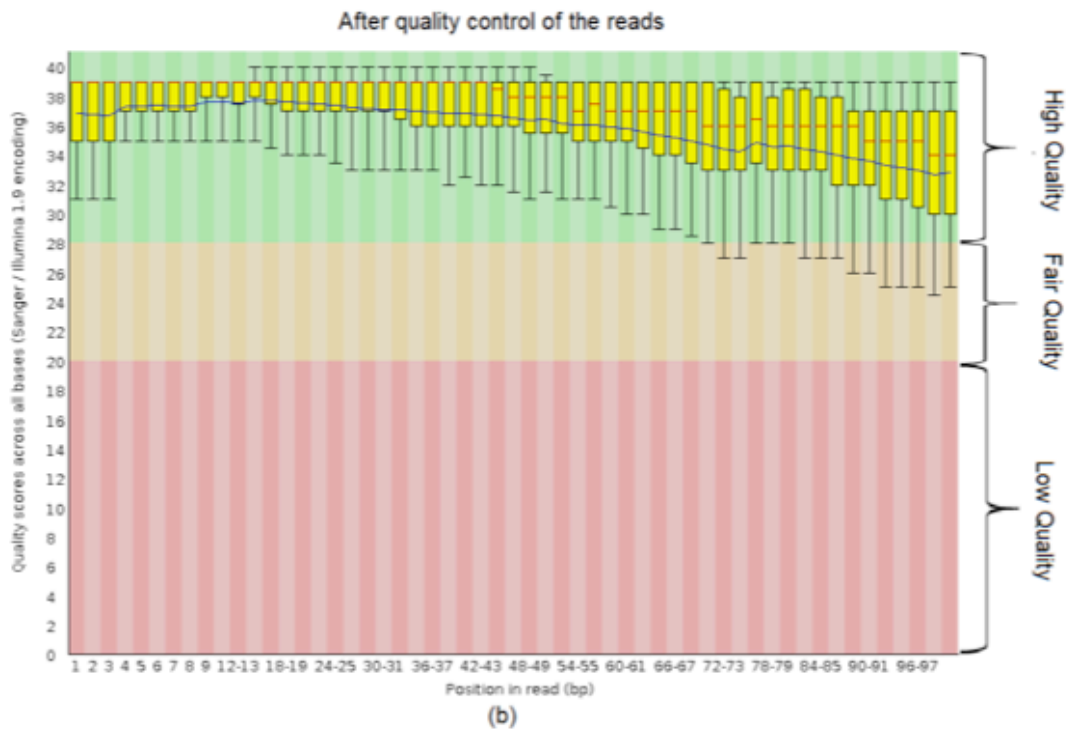
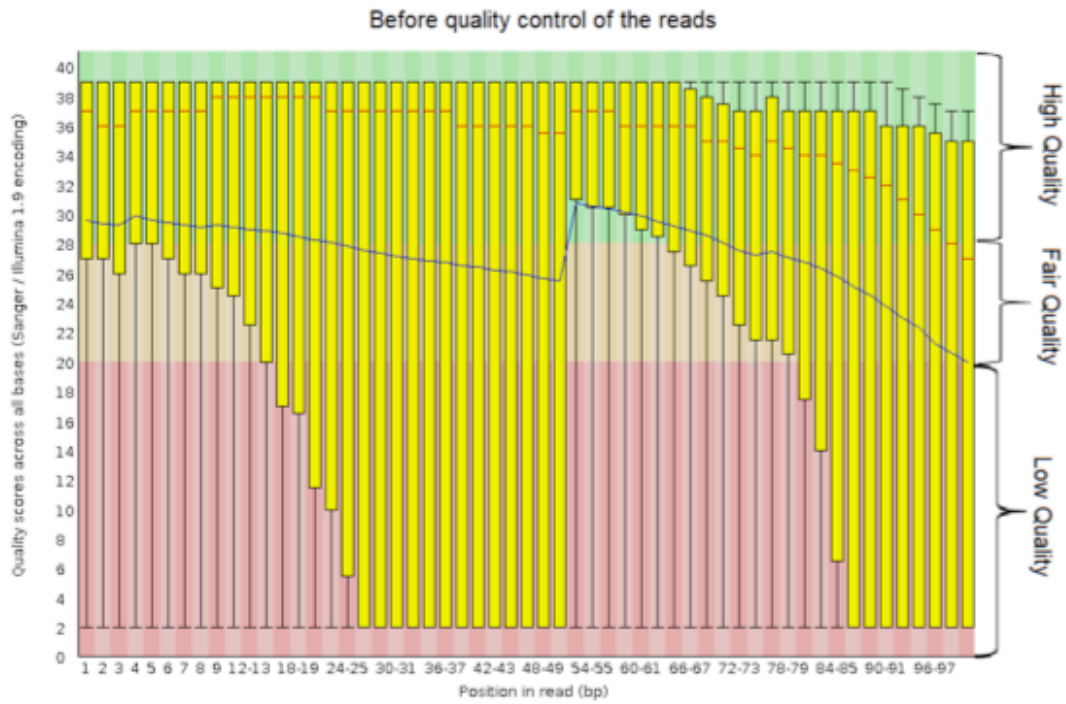
¹Department of Computer Science

²Department of Statistics

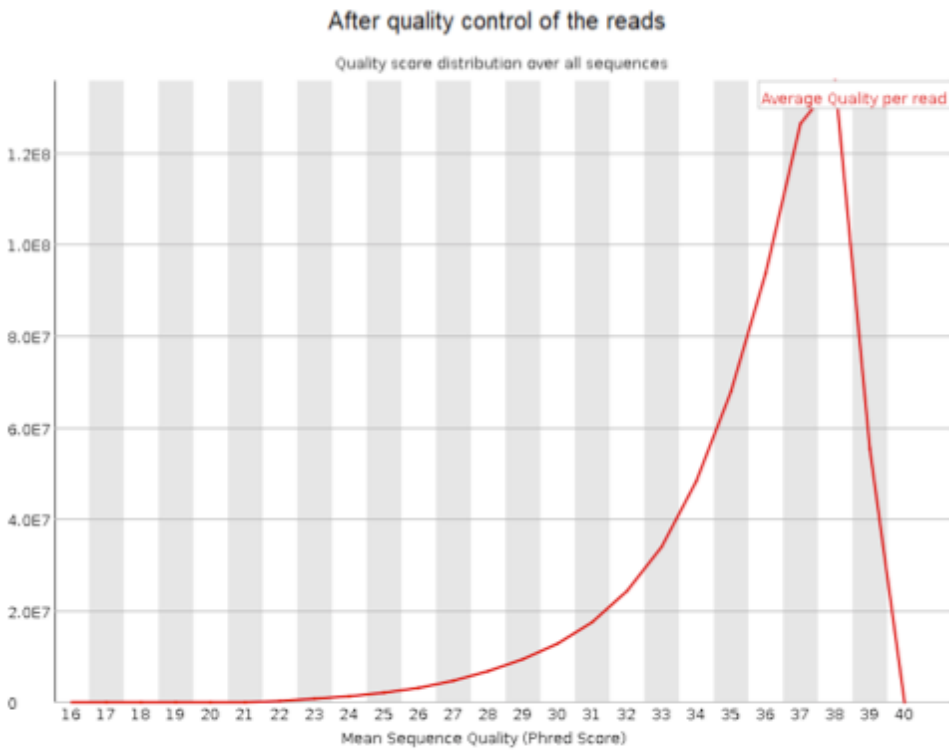
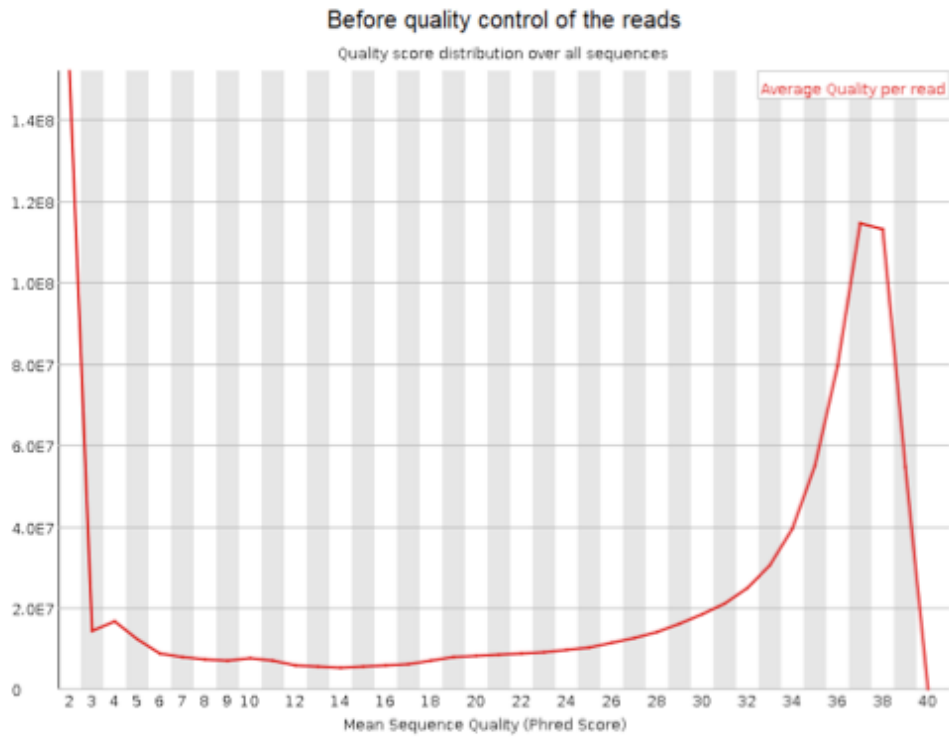
Virginia Tech, Blacksburg, VA 24061.

Email: shabbir5@vt.edu, xwwu@vt.edu, lqzhang@vt.edu

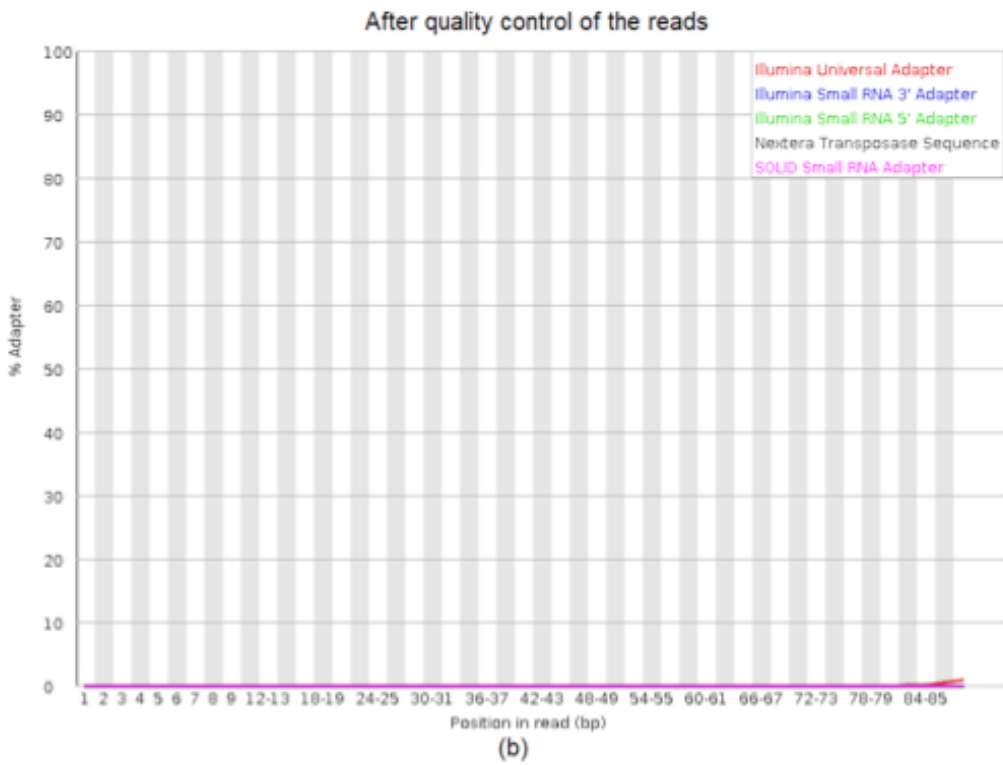
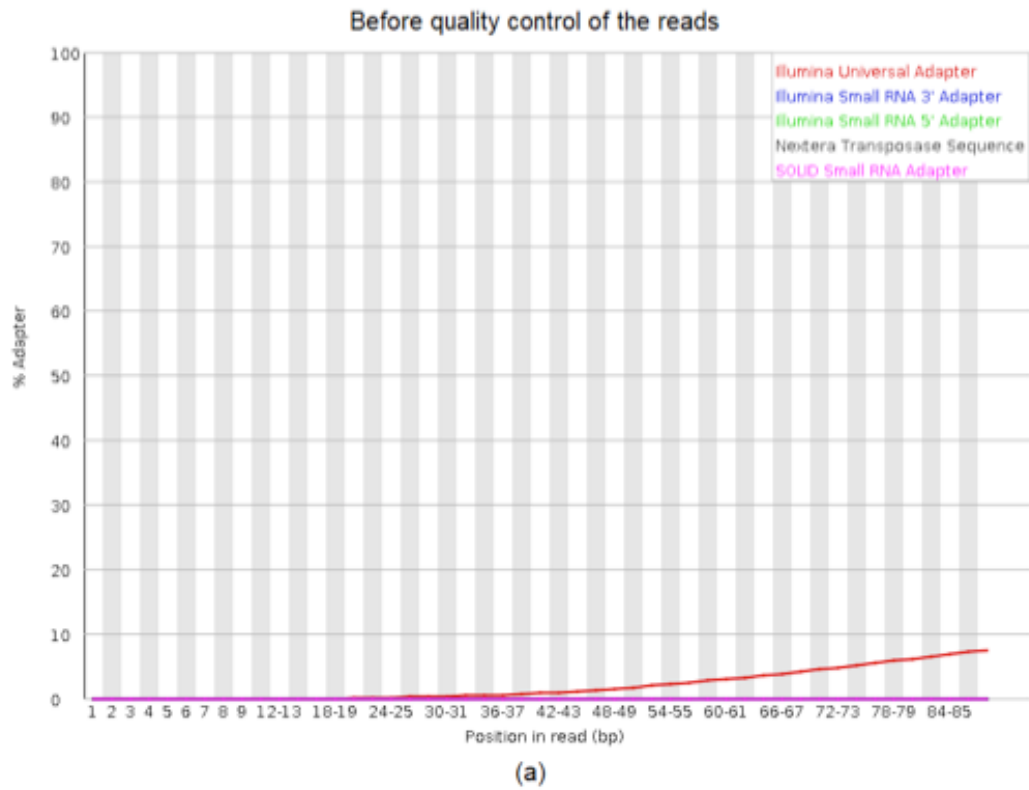
* Corresponding Author



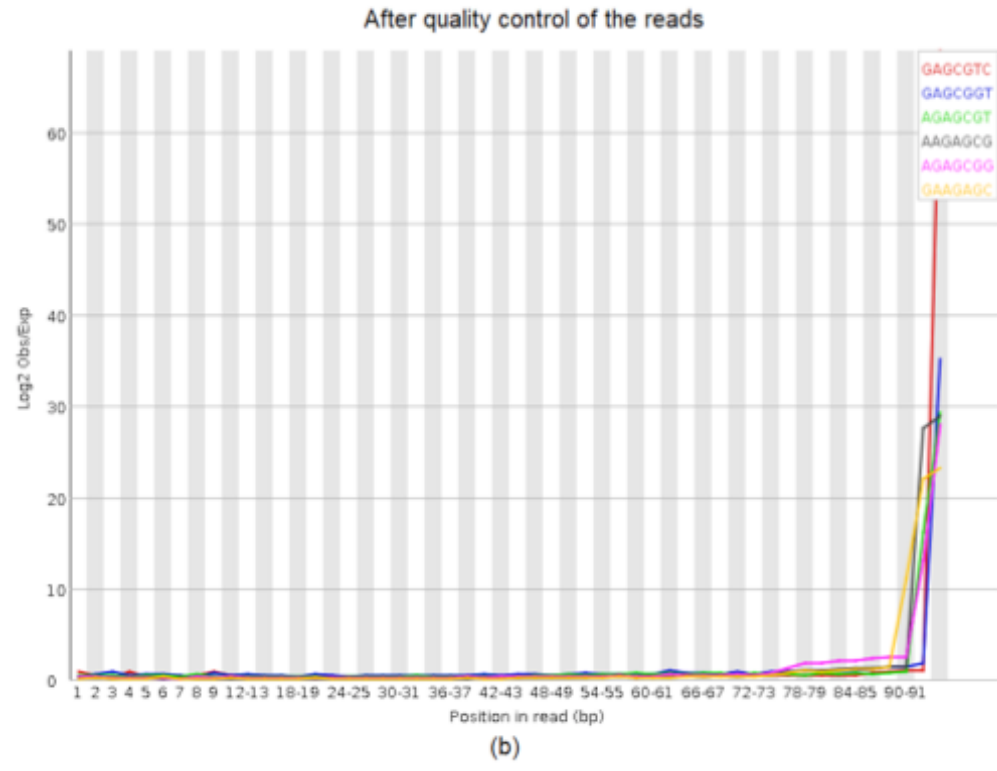
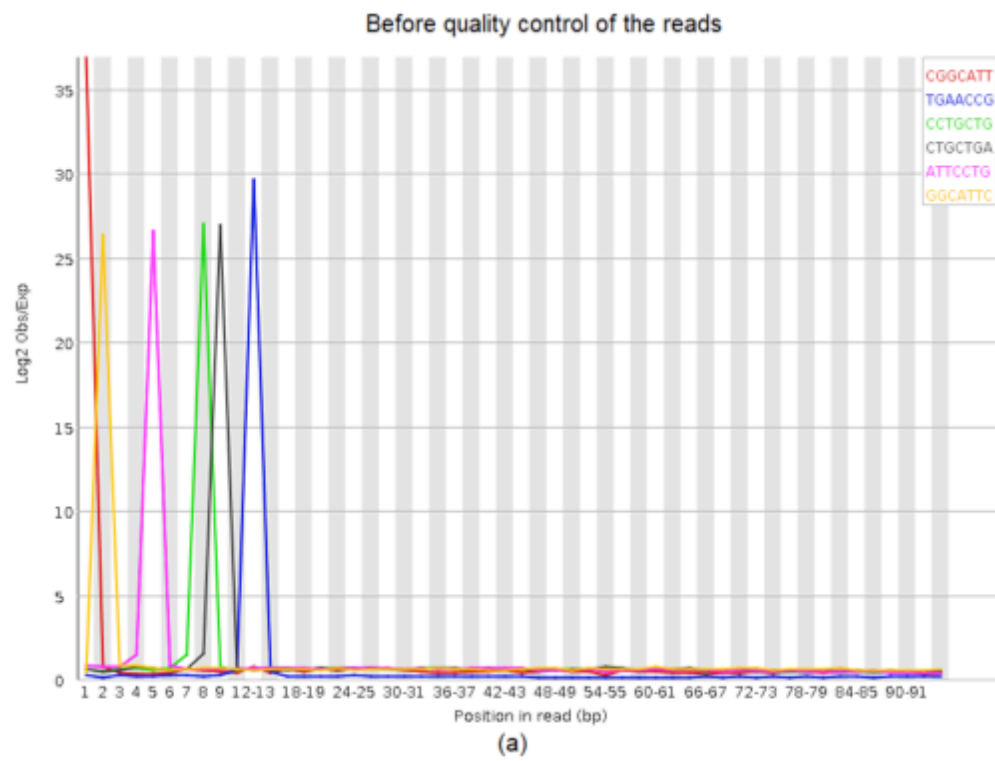
Supplementary Figure 1: Per base sequence quality (a) before and (b) after the quality control of the reads.



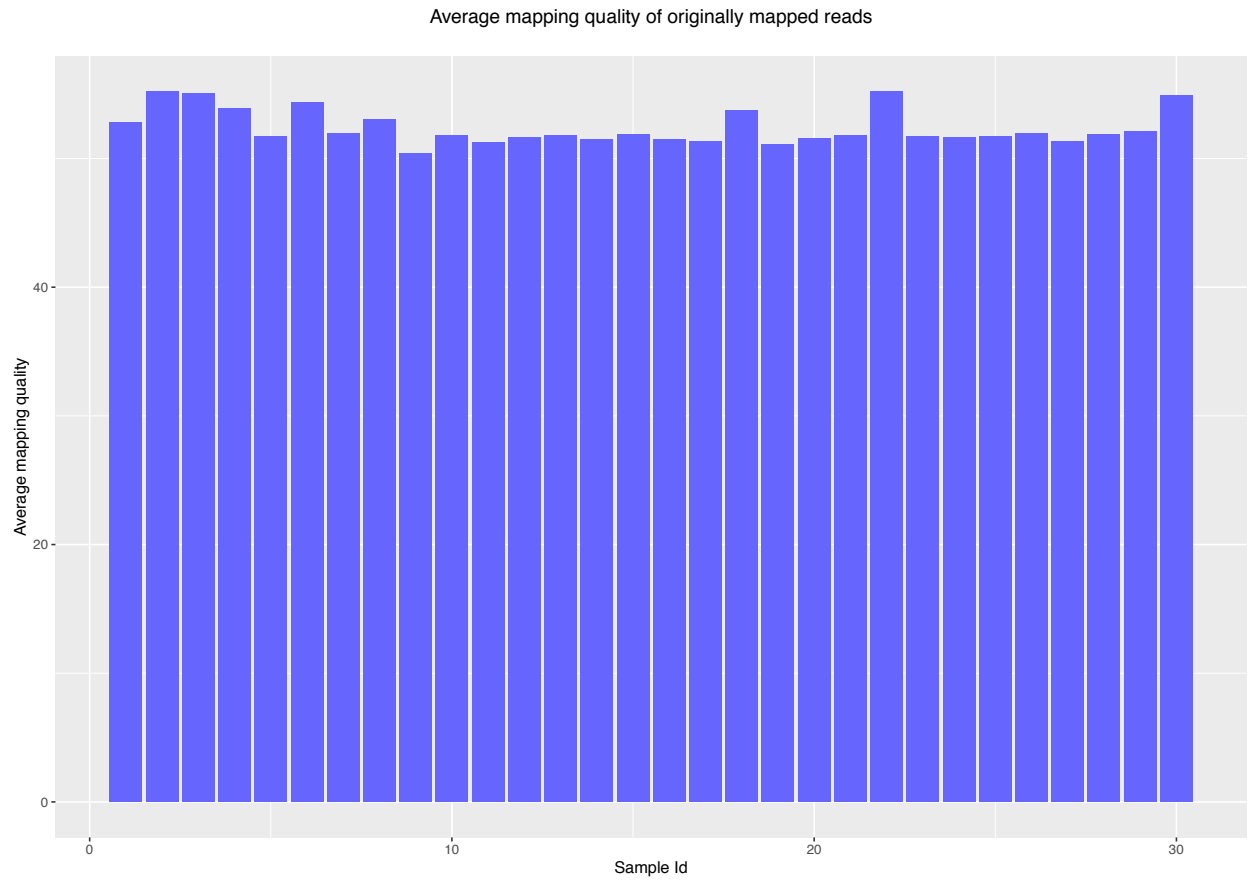
Supplementary Figure 2: Per sequence quality score (a) before and (b) after the quality control of the reads.



Supplementary Figure 3: Percentage of Adapter contents in the reads (a) before and (b) after the quality control.



Supplementary Figure 4: Observance of different k-mers in the reads (a) before and (b) after the quality control.



Supplementary Figure 5: Average mapping quality of the samples for the originally mapped reads.