



Supplementary Figure 3

Figure S3. Alignments of a 2D, template, read and complement reads. Alignments were performed using the UCSC blat aligner (<https://genome.ucsc.edu/>). Color blocks represent alignments, while the gray scale of color blocks represent alignment identities. Red vertical lines within color blocks indicate mismatches. Two gray lines between color blocks indicate gaps. A 2D read has smaller gaps between alignment blocks and higher alignment identities when comparing to the template and complementary reads.