

Figure S3 Forward (F) and reverse (R) strand coverage of 50 nt-reads virtual HTS profile along the whole chromosome II. Some regions are repetitive on the both strands (1, 2) and some others are only repetitive on one strand (3, 4). This figure illustrates that virtual HTS profile is able to discriminate repetitiveness according to the strand in the reference genome.

4 SI C. Jubin et al.