

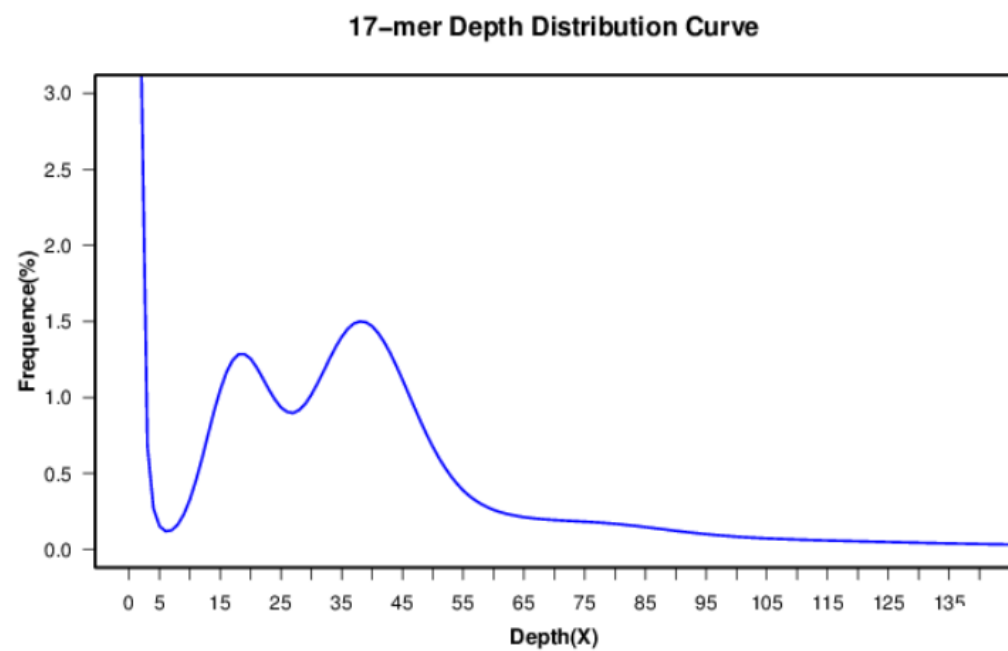
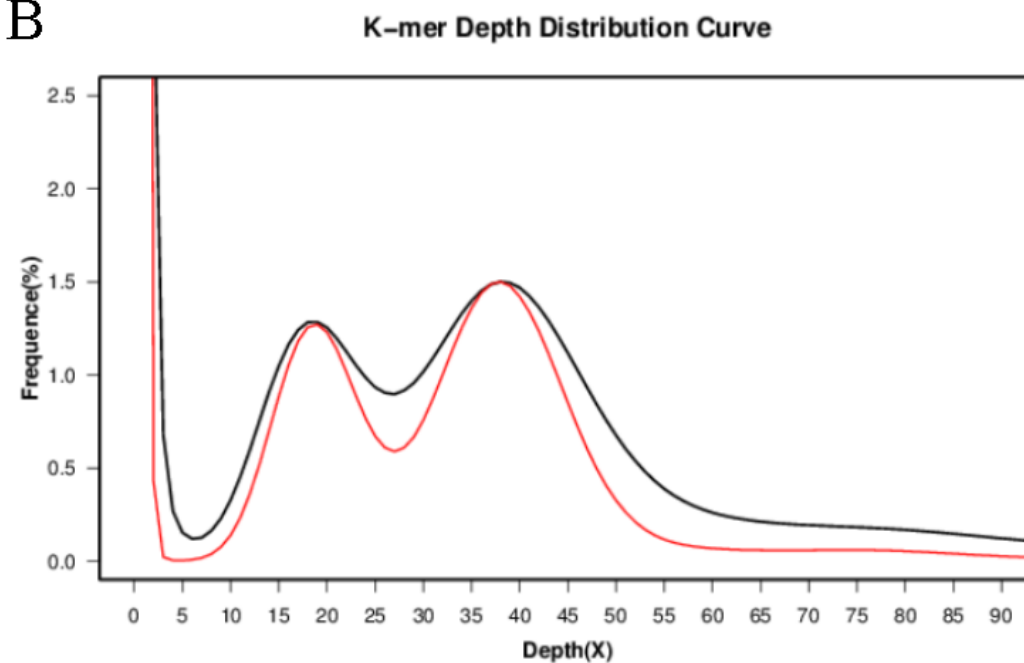
A**B**

Fig.S2 Estimation of genome size and heterozygosity. (A) 17-mer distribution curve. (B) Simulation curve of hybrid rate.