



Figure S1: (A) Chloroplast genome maps of *Q. fleuryi*; (B) Chloroplast genome maps of *Q. gilva*; (C) Chloroplast genome maps of *Q. glauca*; (D) Chloroplast genome maps of *Q. shennongii*; (E) Chloroplast genome maps of *Q. multinervis*; (F) Chloroplast genome maps of *Q. myrsinifolia*. Genes are colored based on different function. The inner circle (dashed gray area) indicates the proportional GC content of the corresponding genes. Regions of the LSC, SSR, IRa, and IRb are indicated.