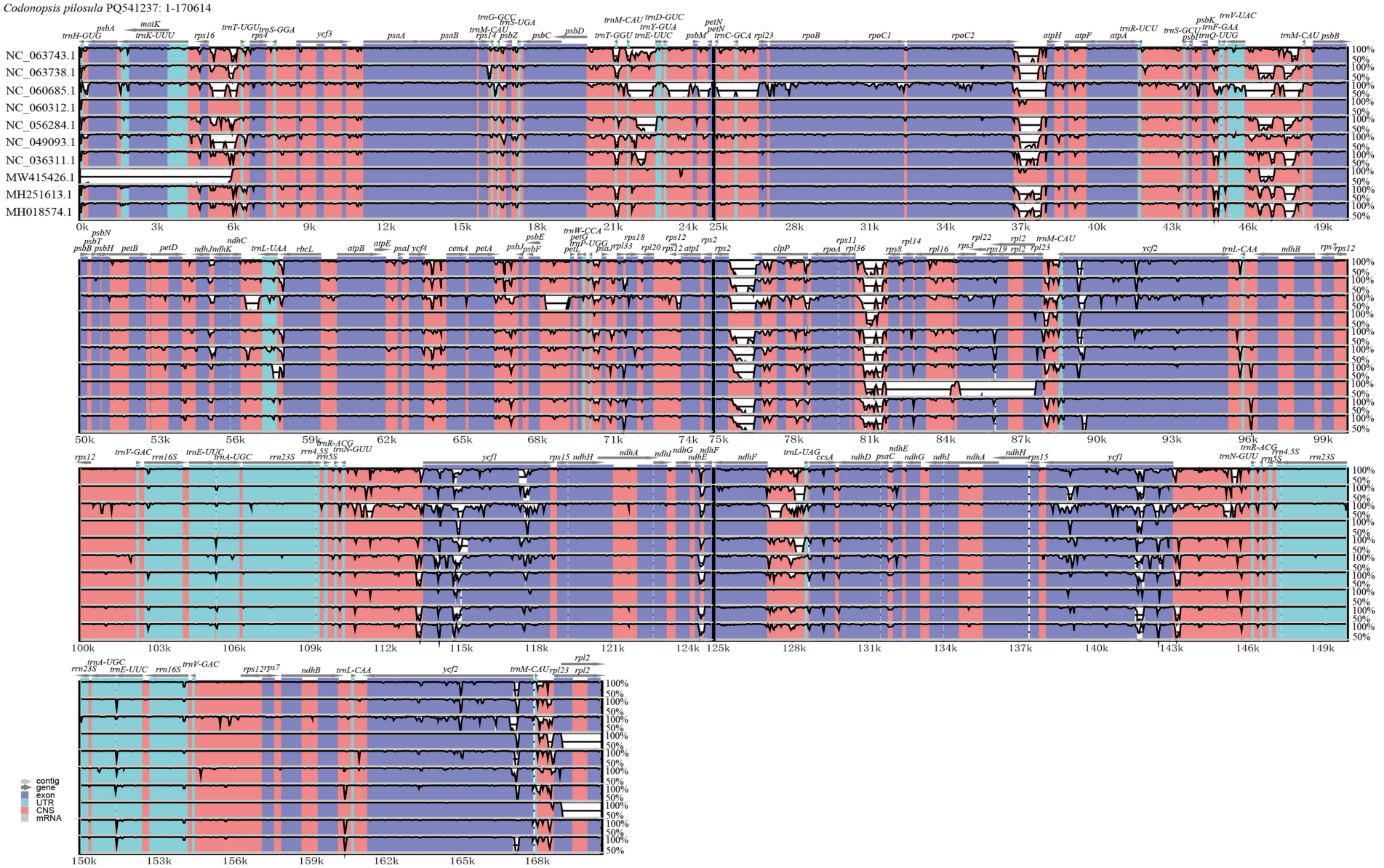


Fig. S2 Whole-chloroplast genome alignment of 11 *Codonopsis* plants. The newly assembled *C. pilosula* genome is used as the reference. The x-axis represents the coordinates of the reference genome, and the y-axis shows the percentage sequence identity (50-100%). Annotated genes are indicated by gray arrows.