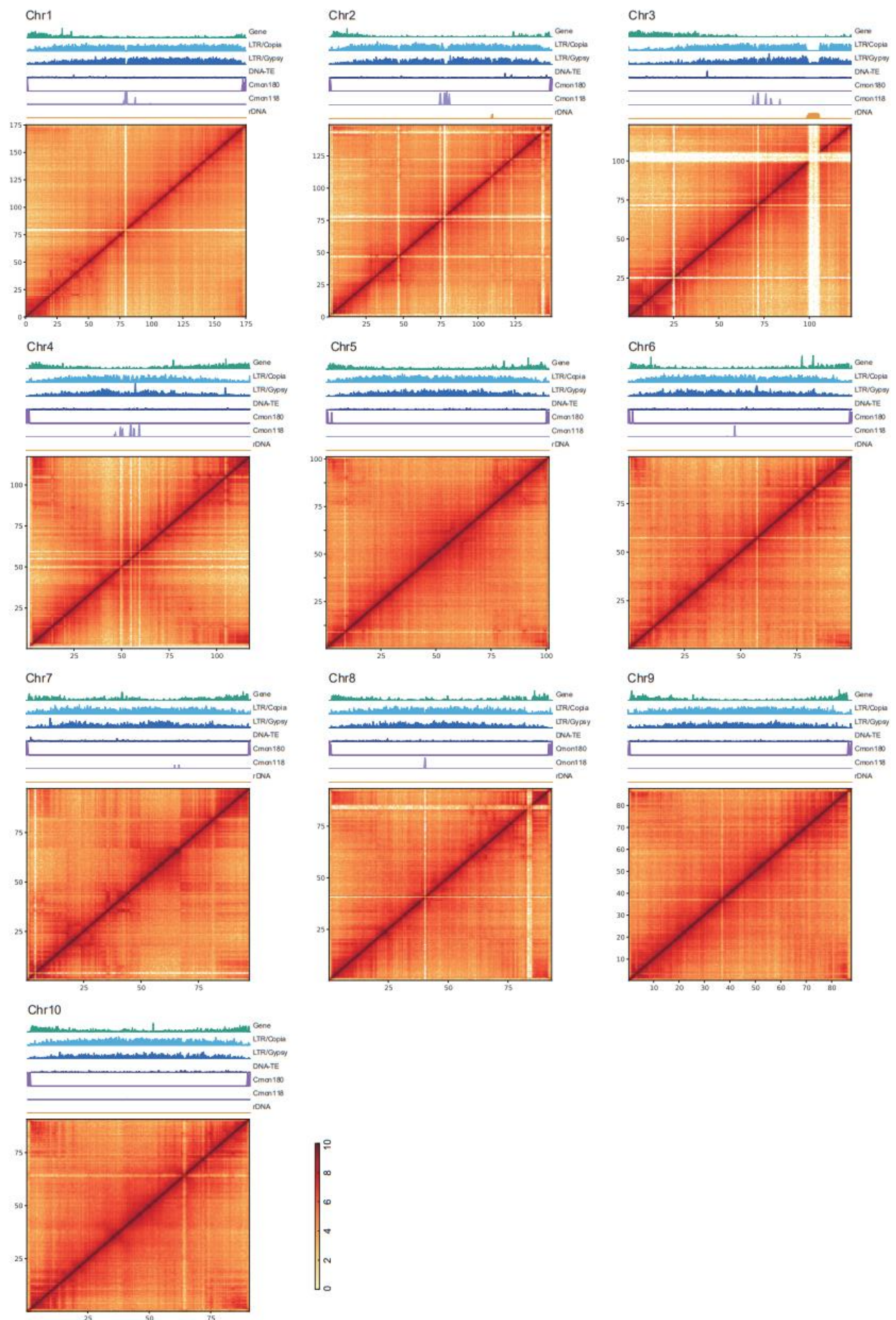




Supplementary Fig. 7 Detailed high-throughput chromosome conformation capture (Hi-C) chromatin interaction map of *C. monnieri*. For each image, the top tracks are the density of genes, long terminal repeat (LTR)/*Copia*, LTR/*Gypsy*, DNA transposons, *Cmon180*, *Cmon118*, and rDNA, respectively. The bottom panels show Hi-C chromatin interaction map (500 kb resolution) and the axes represent chromosome position in Mb. The color bar indicates the intensity of interaction. Chromosomes 1-10 of the *C. monnieri* genome are shown.