

**S2 Fig. SNP and tag distribution across the chicken genome in 824 individual samples digested by *EcoR* I- *Mse* I.** In total, 292 K SNPs were identified among all individuals. The genome characteristics and genome-wide distribution of restricted digest fragments are represented circularly. The exterior circle displays the lengths of the chromosomes. The four interior circles show the distribution of fragments (green), GC islands (orange), repeat regions (black), and SNPs (red) from outside to inside.

