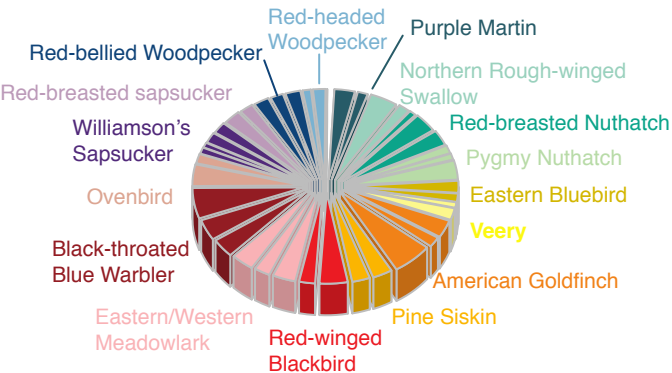
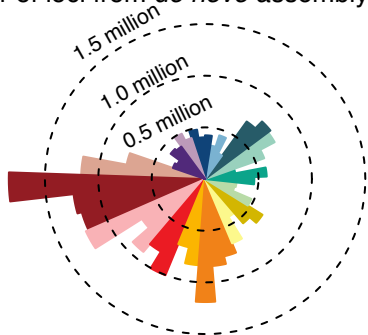


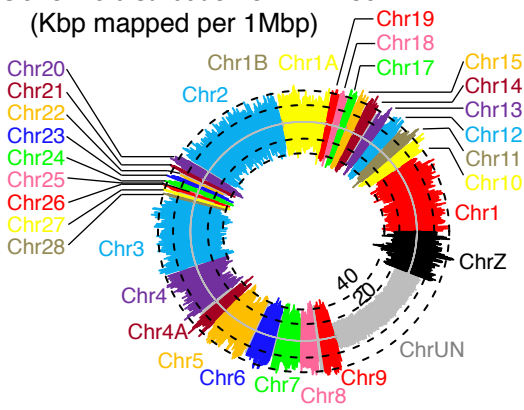
A. Raw illumina sequencing data



B. Number of loci from *de novo* assembly



C. Genomic distribution of RAD loci (Kbp mapped per 1Mbp)



D. Sequencing coverage and genomic coverage

