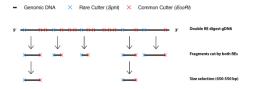**Supplementary Fig. S1.** Overview of ddRADseq protocol. (A) Genomic DNA for each individual is double restriction digested and size selected, (B) individual-specific adapters are ligated, (C) samples are pooled and sequenced in multiplex, (D) resulting reads are demultiplexed by individual, (E) *de novo* assembly of loci is performed, and presence/absence and coverage are calculated across loci, (F) loci from individuals are compared, and (G) h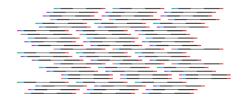aplotypes are called for variable loci.