

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Sampling used for molecular analyses.

File Name: Supplementary Data 2

Description: The number of raw reads, number of reads passed filters (with the proper sample barcode and restriction enzyme cutsites present), number of reads mapped and unmapped on the reference genome, number of clusters and high-coverage clusters per sample, estimated heterozygosity and error rates, consensus clusters in each sample and the final number of loci in the assembly.