



Supplementary Figure 4: Wright's F_{IS} values p -values for panmixia (A) and F_{ST} values with p -values for subdivision (B) of nine selected loci across all samples. Dots indicate mean values, dashes show lower and upper limits calculated with the following formula: $L_i = F_{XY} - \text{StdErr} \times t$ and $L_s = F_{XY} + \text{StdErr} \times t$. Dashes for All loci are the 95% confidence intervals as obtained by jackknife resampling over populations for each locus and by 10,000 bootstraps over loci for the average.