Supplementary Figure 2: F_{ST} values with p-values for subdivision of 10 loci across all samples. Dots indicate mean values, dashes show lower and upper limits calculated with the following formula: Li = F_{ST} − StdErr × t and Ls = F_{ST} + StdErr × 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.