



Fig. S8. Estimated values of H_j (diversity) and δ_j (specialization) in chromosomes 21, 18, 13, 22, 20 and X (dataset C). Open circles are plotted in the mean of the 2000 bootstrap replicates for each parameter and the corresponding approximate 95% confidence intervals are plotted as continuous lines in each axis. This figure amplifies the brown box of Fig. 2.