



Figure S33: The proportional overlap of outliers above the 95% quantile of 100% neutral data for each of the 16 demographic treatments after 100 generations. Axis orderings were determined through hierarchical clustering. Heatmaps are shown for single measures of F_{ST} , D_{XY} and $\Delta\pi$ in the first column, and combined measures in the second column. Heatmaps are coloured according to a common scale of 0 to 1. Treatments are labelled with founding bottleneck (Bot), DP population size (Pop2), and migration (Mig) values.