



Figure S41: Effects of demographic treatments on measures of genetic divergence across all sampling generations for $\mu_\sigma = 0.1$ simulations. Point colour and shape denote treatment groups for founding bottlenecks (**A**), prolonged bottlenecks (**B**) and migration (**C**). Each point represents values of divergence averaged across all genes within individual treatments groups, averaged within treatment levels, and averaged over 20 iterations.