

a)

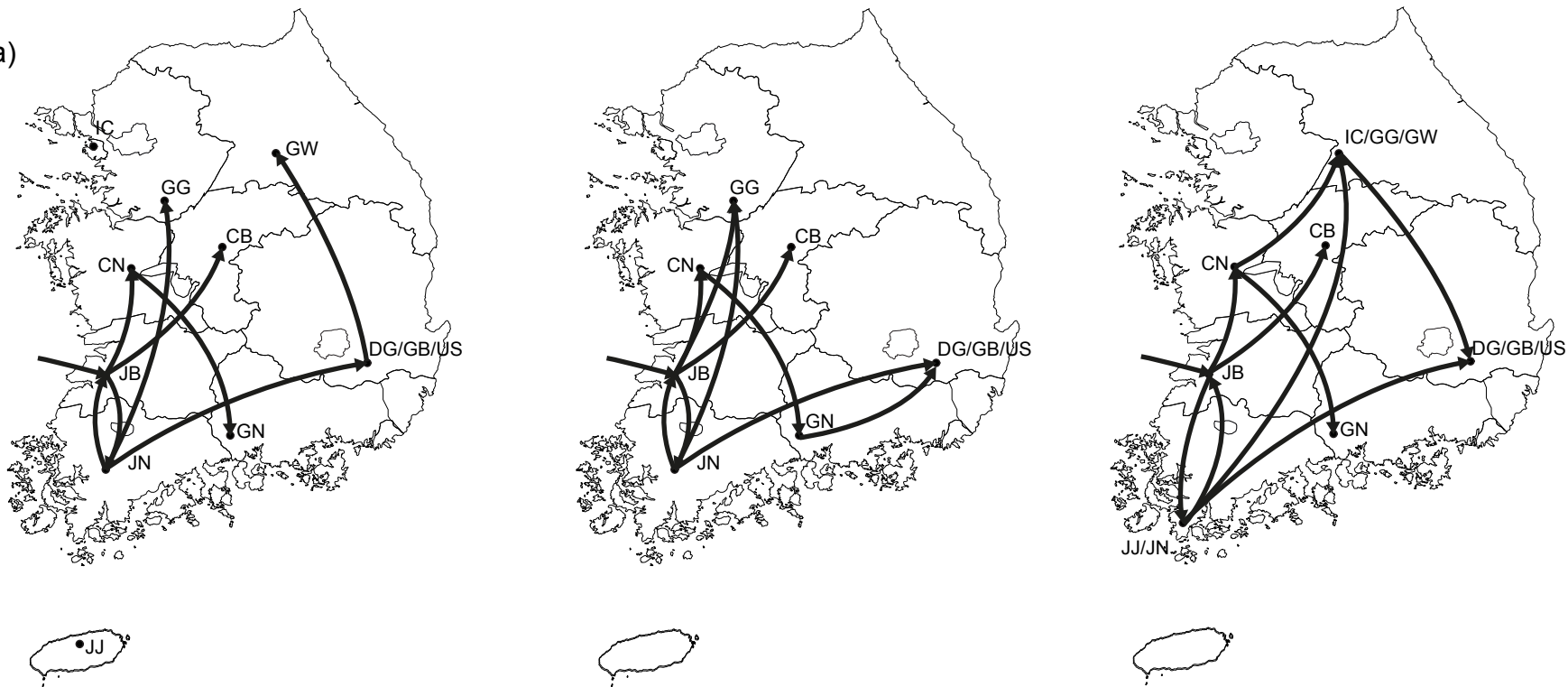


Fig. A.2. Effect of different sampling and grouping of provinces with only one sequence on phylogeographic inference. Arrows show inferred epidemiological links where Bayes Factor support for rate indicators >10, as determined using SPREAD. a) Results as shown in Figure 3. b) Sequences isolated in provinces from which only one sequence is available have been removed from the alignment (except DG/GB/US, which have been grouped). c) Sequences isolated in provinces from which only one sequence is available have been grouped with geographically neighbouring provinces for phylogeographic inference.