

Supplementary material for

Measuring genome-wide genetic variation to reassess subspecies classifications in *Dodonaea viscosa* (Sapindaceae)

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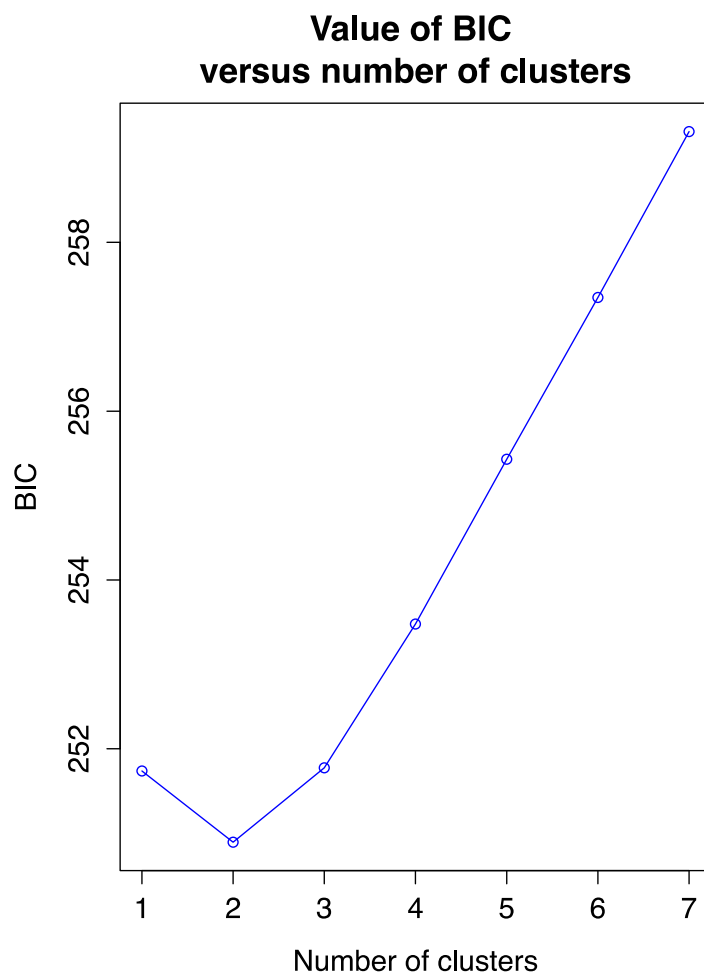


Fig. S1 Bayesian Information Criterion (BIC) values from a Discriminant analysis of principal components (DAPC) when $K = 1$ through to 7. The ‘elbow’ in the graph at $K = 2$ indicates 2 to be the most likely number of genetic clusters. This agrees with the STRUCTURE analysis.