

Fig. S1. Summary statistics of 2053 DArTag SNP markers used to assess the genetic diversity among the Ex-PVP inbred lines

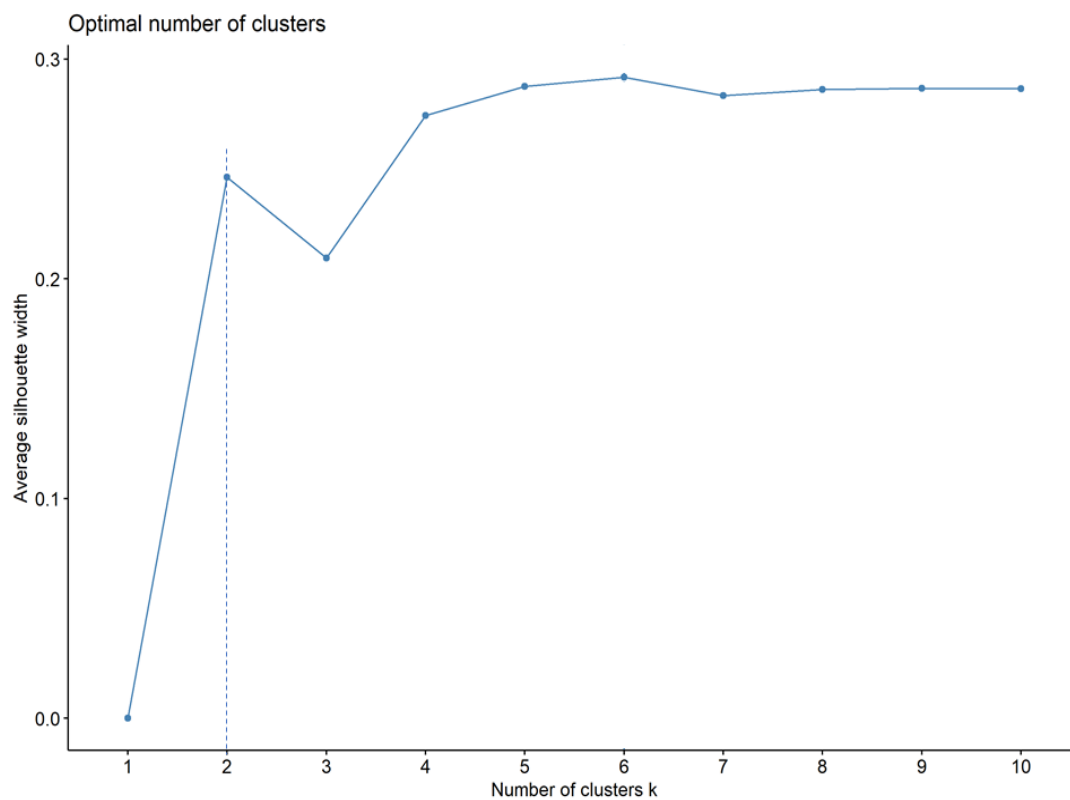


Fig. S2. Determination of the most appropriate number of clusters in structure analysis using cross-validation error (k) means