Supplementary Material

**Genomic Selection for Ascochyta Blight Resistance in Pea**

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A.

B.

Supplementary file 4: Determination of population structure. (A) The mean log likelihood of K (postulated number of subpopulations) for K = 1 to15, with bars representing ± standard deviation. (B) DeltaK for 1 to 15 subpopulations as calculated by the method of Evanno et al. (2005), indicating that population structure is best represented by 2 or 10 subpopulations.