Figure 1. Each colour represents a genetic cluster in the bar chart with each line representing the assignment probability of an individual to each cluster. The population locations of each cluster are indicated by the colour-coded pie charts, overlayed upon a map of the study area. The study area is filled by habitat suitability scores (0-1, with 1 being most suitable in green). The assignment probabilities of individuals to each genetic cluster are represented as proportions of colour within each population. Total sample size = 186 with 8-20 individuals per population.