Figure S1. *Genotype accumulation curve.* The minimum number of loci necessary to discriminate between individuals in a population was calculated in *R* with the "poppr" package (Kamvar, Tabima & Grunwald, 2014). Multilocus genotypes (MLG) at eight microsatellite loci (112 ABFT larvae) were re-sampled 10.000 times to count the number of observed genotypes.