



**Fig. S2.** Genotype accumulation curve for OR forest *P. ramorum* isolates. The vertical axis denotes the number of observed MLGs, from 0 to the observed number of MLG in the forest populations, for a number of loci, indicated on the horizontal axis, randomly sampled without replacement. Each boxplot contains 1,000 random samples representing different possible combinations of  $n$  loci. The horizontal red dashed line represents 90% of MLG resolution.