Figure S2. Studied sequence around *mac-1*. Position of the four polymorphisms chosen for extensive genotyping is indicated by stars for SNPs and by triangles for PCR product length polymorphisms. The second polymorphism, *mac-1*, corresponds to the outlier locus identified by genome scan. The sequence has been segmented in three regions (region 1 of 880 bp; region 2 of 1500 bp; region 3 of 750 bp) for gene tree reconstruction.