Figure S1: Parsimony-based haplotype networks for *Bufo balearicus*, and *B. viridis*.
Ellipsoids represent existing haplotypes, with locality numbers of the populations were they
occur; connection with a dash indicates single mutational steps; small empty dots symbolize
the number of hypothetical haplotypes not found; the rectangle refers to the most frequent
haplotype. For major clades, the bootstrap support from the Maximum-Likelihood phylogeny
(not shown) is given.