Figure S2. Biogeographical scenario – ancestral ranges and biogeographical events – inferred by Lagrange M1 analysis represented on the median penalized likelihood (Med PL) dated phylogenetic tree. See Fig. 6 and main text for explanation. Pie charts represent ancestral area(s) based on the median PL estimation. Dispersal events are represented on palaeogeographic maps according to time slices. Thicknesses of arrows are proportional to the number of dispersals (see Table S4 in Appendix S2 for more details). Abbreviations: junk (in black): sum of ancestral area probabilities <0.1; star: hard incongruent nodes between methods; @: hard incongruent node within methods; grey lines: ranges between time slices and TS: time slice.