**Suppl. Figure 3**: Mismatch distributions of Galápagos marine iguana mtDNA CR haplotypes. (a) clade C individuals from western islands; (b) clade B individuals from eastern and central islands; (c) clade C individuals from northern islands; (d) the entire marine iguana dataset. The line with closed circles is the observed distribution of pairwise differences and the smooth line is the expected distribution under the sudden expansion model \([90]\). The sum of squared deviation (SSD) between the observed and expected distributions, the raggedness statistic \((r)\) \([92]\), and their corresponding \(p\)-values are reported. The values of the expansion model parameters \(\tau\) (mutational time), \(\theta_0\) and \(\theta_1\) are also shown as well as the 95% confidence interval of the estimate of \(\tau\).