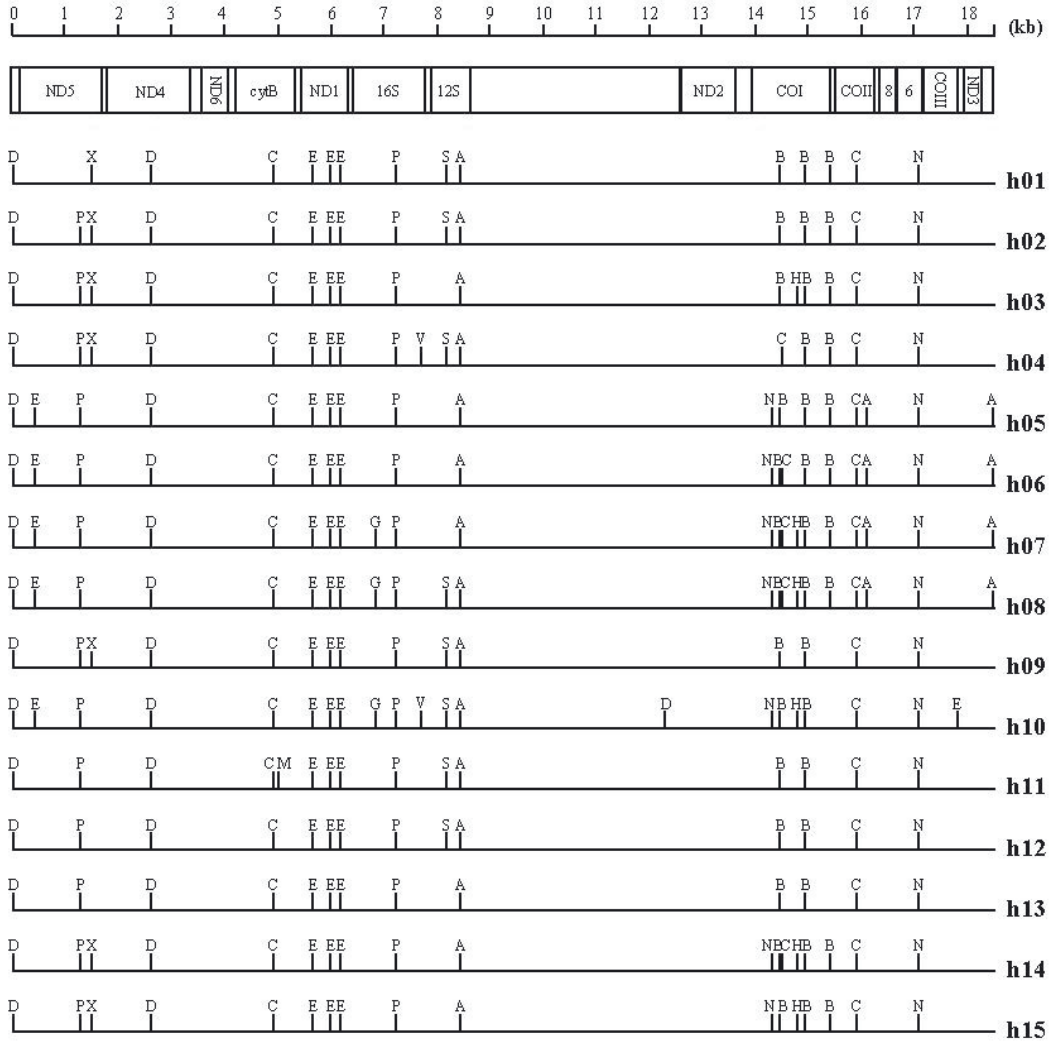


ELECTRONIC SUPPLEMENTARY MATERIAL



Supplementary Figure 1. Linearised restriction site maps and gene positions of the 15 haplotypes of *Plebeia remota* mitochondrial genomes. M: *Bam*H I, B: *Bcl* I, G: *Bgl* II, C: *Cla* I, E: *Eco*R I, V: *Eco*R V, A: *Hae* III, D: *Hind* III, N: *Nde* I, P: *Pst* I, S: *Sca* I, X: *Xba* I, and H: *Xho* I. ND5: NADH dehydrogenase subunit 5, ND4: NADH dehydrogenase subunit 4, ND6: NADH dehydrogenase subunit 6, cytB: cytochrome B, ND1: NADH dehydrogenase subunit 1, 16S: large subunit ribosomal RNA, 12S: small subunit ribosomal RNA, ND2: NADH dehydrogenase subunit 2, COI: cytochrome c oxidase subunit 1, COII: cytochrome c oxidase subunit 2, 8: ATP F_0 synthase subunit 8, 6: ATP F_0 synthase subunit 6, COIII: cytochrome c oxidase subunit 3, and ND3: NADH dehydrogenase subunit 3.