

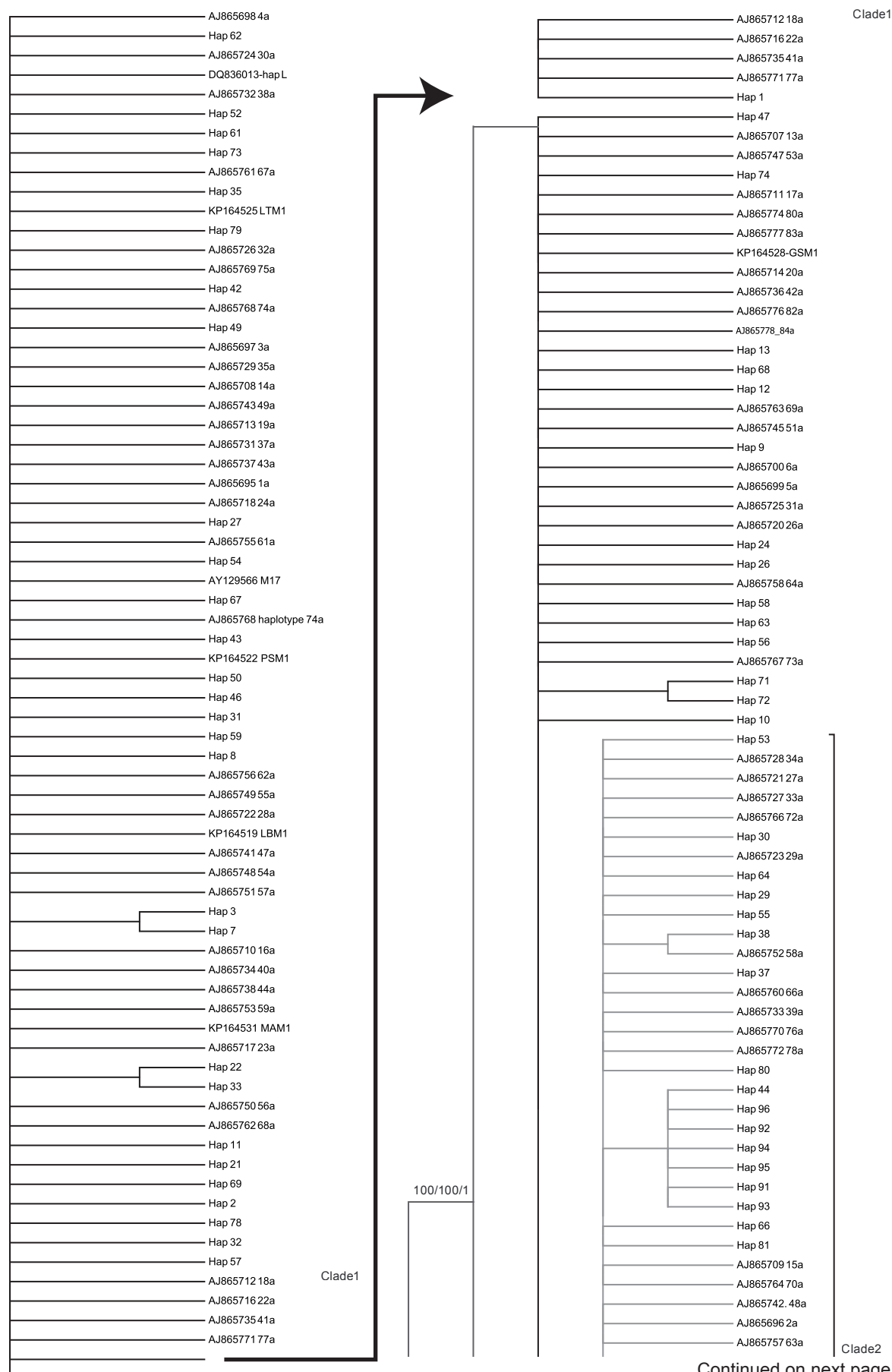
Supplementary Information

Genetic variability of the Lessepsian migrant mussel *Brachidontes pharaonis*

M Antit, N Amor, J Urra, AN Alagaili and S Farjallah

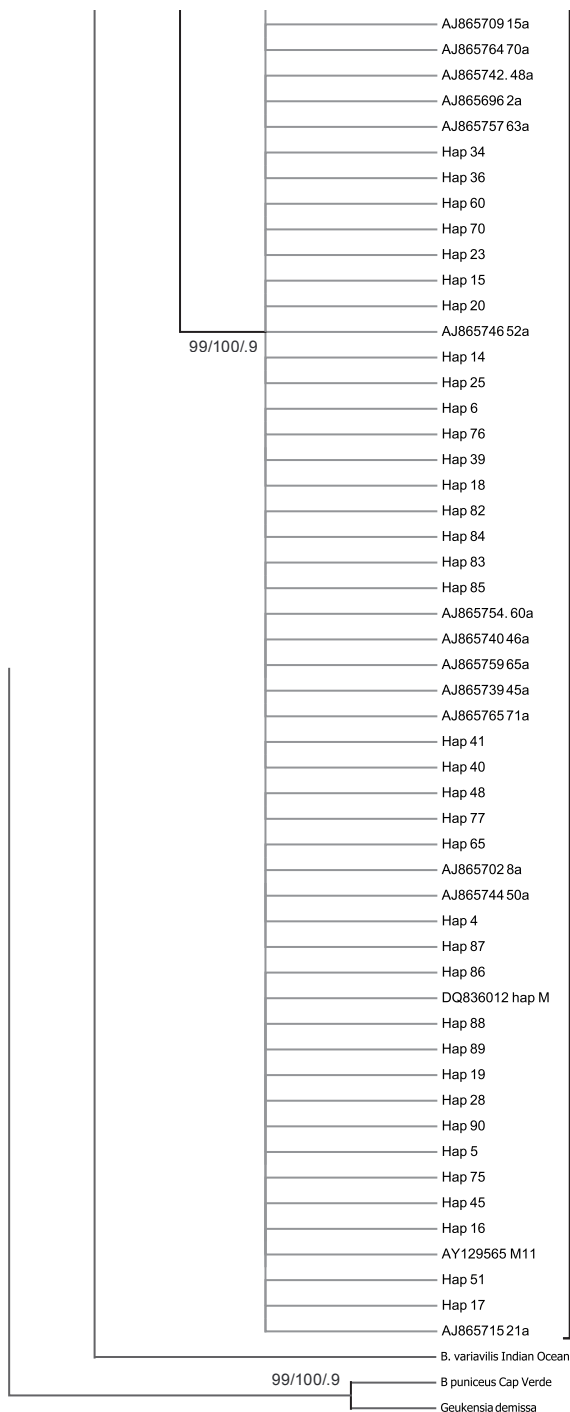
African Journal of Marine Science 2018, 40(2): 211–217.

<https://doi.org/10.2989/1814232X.2018.1476265>

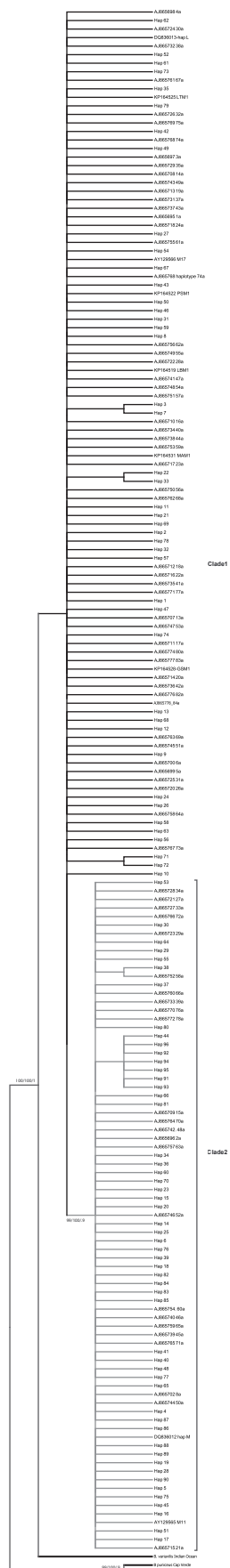


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Supplementary Figure S1: Consensus trees constructed with neighbour-joining (NJ), maximum-likelihood (ML), and Bayesian inference (BI) methods, showing phylogenetic relationships among *Brachidontes pharaonis* inferred from cytochrome oxidase subunit I (COI) sequences. Numbers indicated at branch nodes are bootstrap values (NJ and ML) and the posterior probability (BI)



Complete tree layout