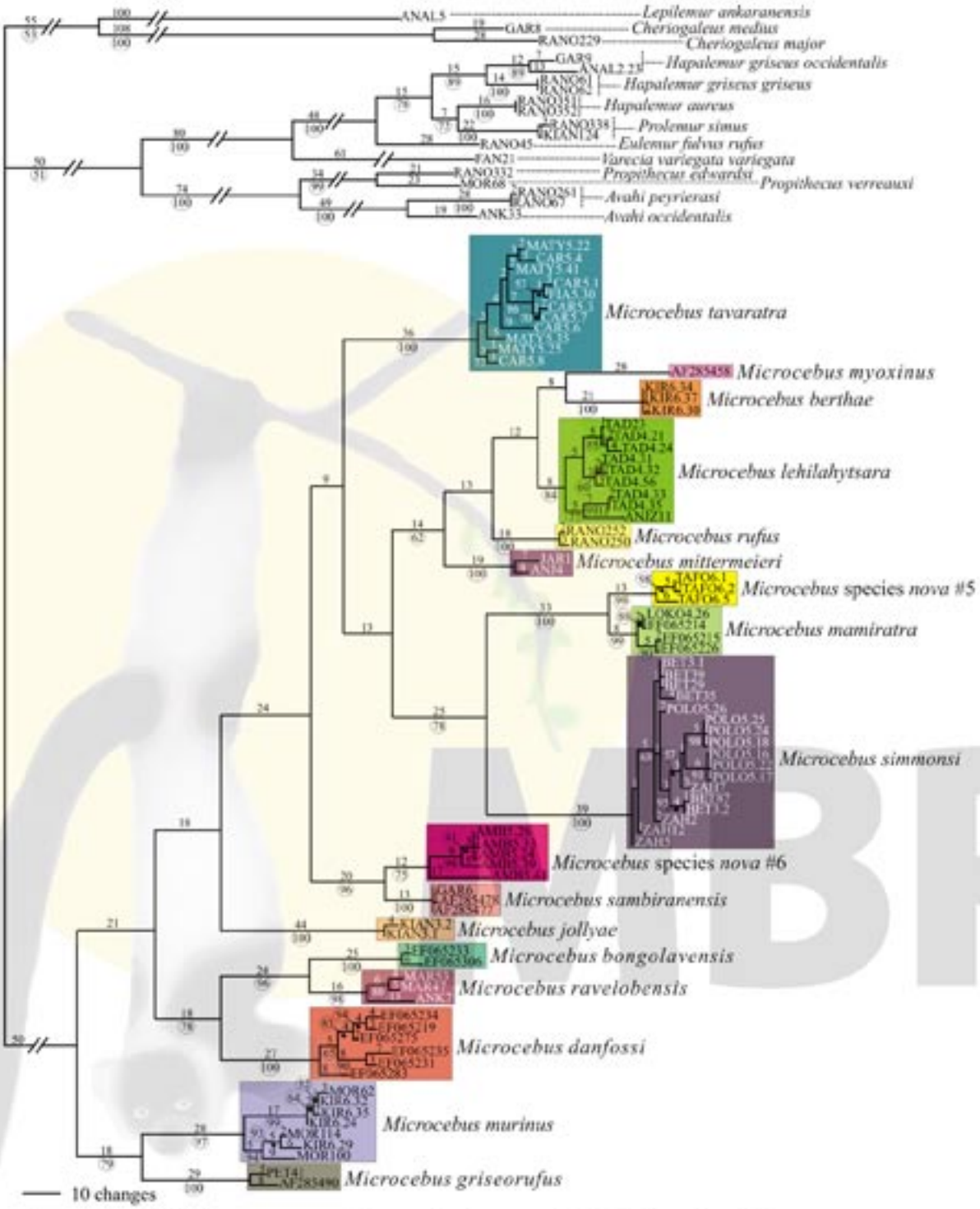

Appendix II(a). Maximum parsimony phylogram derived from the D-loop sequence data from 82 *Microcebus* individuals with 18 out-group taxa (one of 3886 most parsimonious trees). Values above branches indicate number of changes between nodes. Values within the circles along the branches indicate support of bootstrap pseudoreplicates. Length=2052; CI = 0.4235; RI = 0.8460; RC = 0.3583; HI = 0.5765.