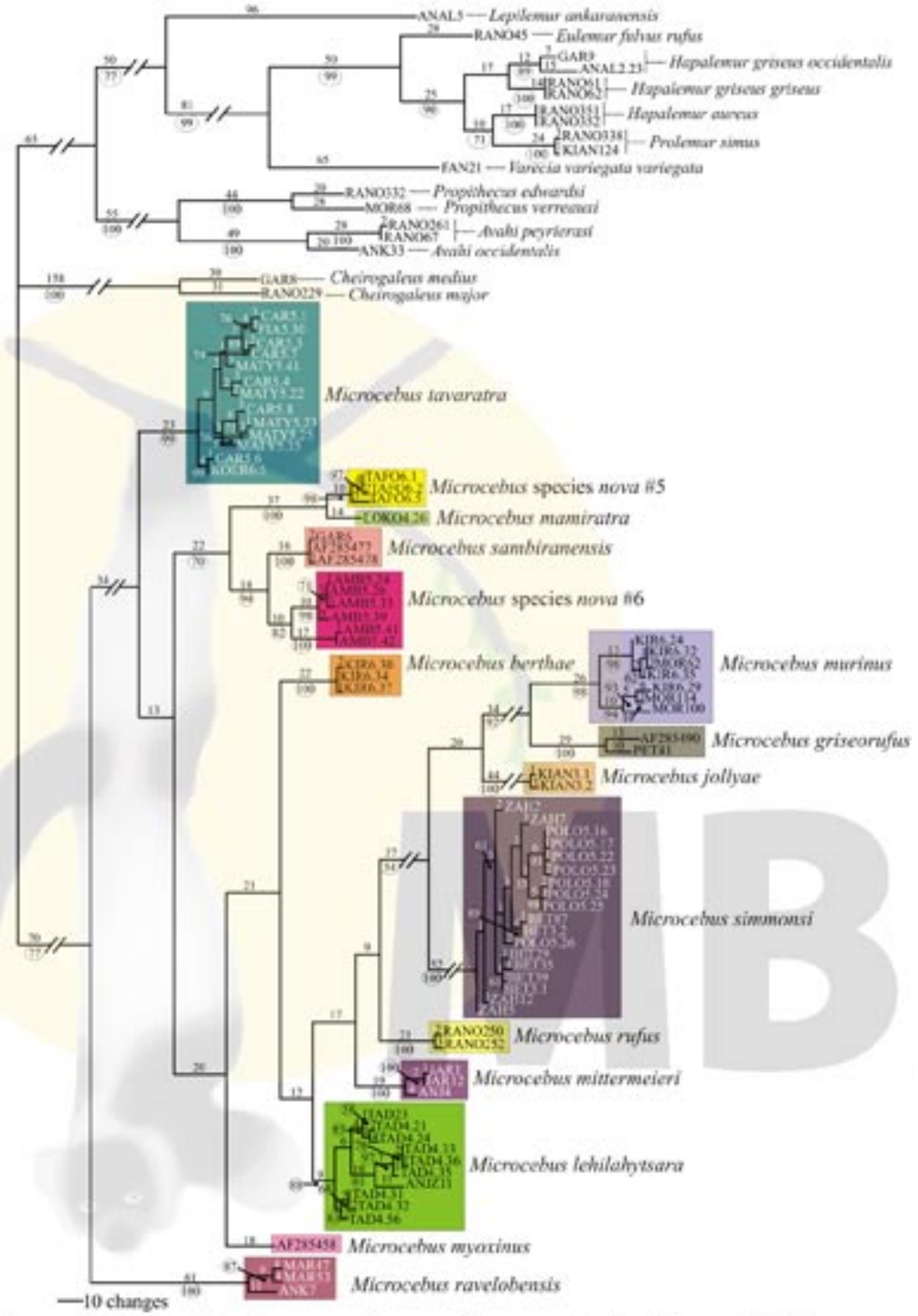

Appendix II(c). Maximum parsimony phylogram derived from the D-loop sequence data from 77 *Microcebus* haplotypes with 18 out-group taxa (one of 364 most parsimonious trees). Values above branches indicate number of changes between nodes. Values within the circles along the branches indicate support of bootstrap pseudoreplicates. Length=2138; CI = 0.4574; RI = 0.8578; RC = 0.3924; HI = 0.5426.