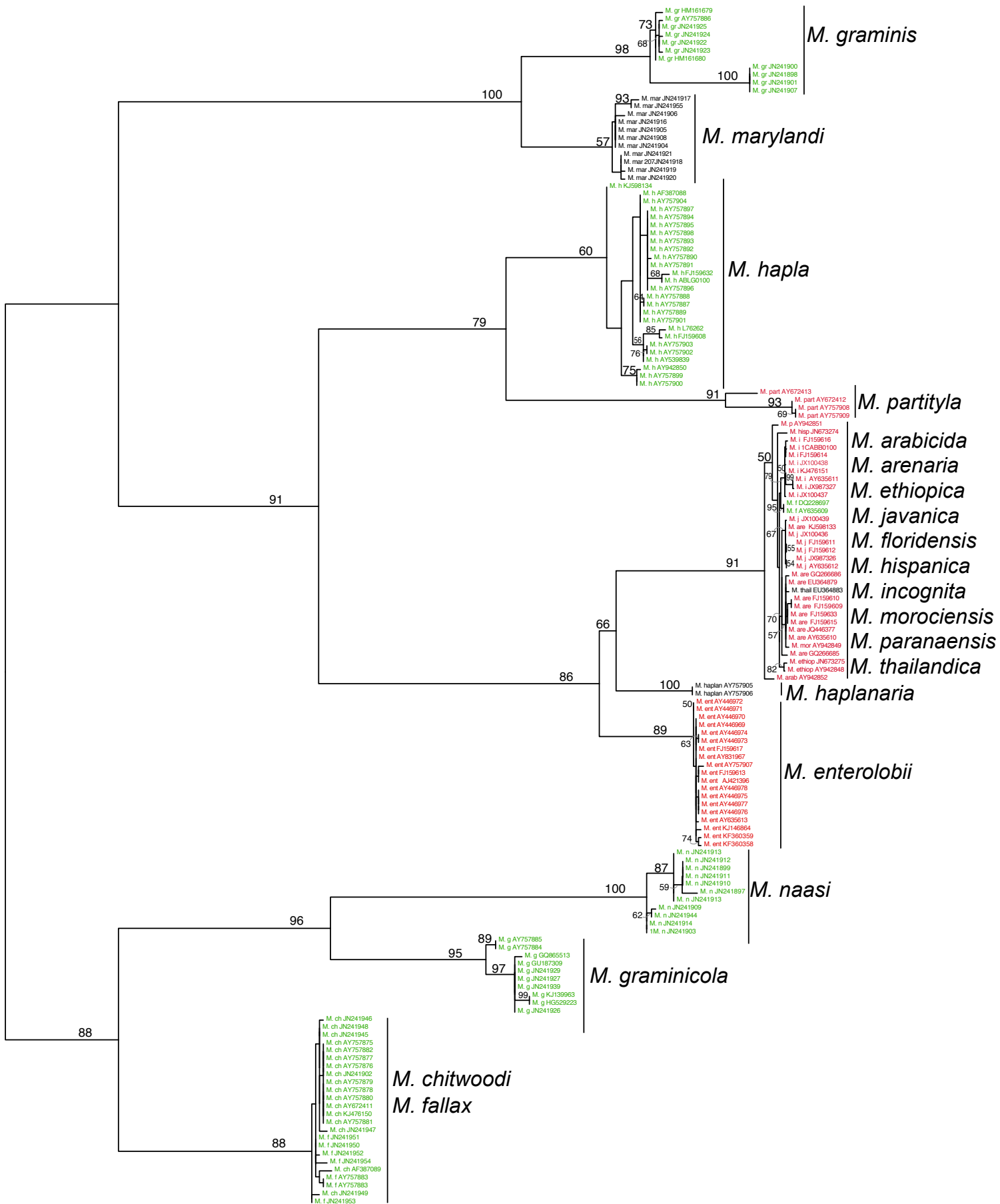


**S3 Figure. Maximum Likelihood (ML) phylogenetic tree of *Meloidogyne* spp. based on the *cox2-rrnS* mitochondrial region. Numbers above branches represent ML bootstrap values >50%. Species names in red, green or black indicate species with mitotic parthenogenesis, meiotic parthenogenesis or unknown reproductive strategy, respectively.**



0.1 substitutions/site