



Development of microsatellite markers for *Hancornia speciosa* Gomes (Apocynaceae)

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ABSTRACT. Herein, we describe 34 microsatellite loci developed using an enrichment genomic library for the tree species *Hancornia speciosa* Gomes (Apocynaceae). Thirty-five individuals were genotyped using 34 primers to analyze the polymorphisms at each locus. The number of alleles per locus ranged from 4 to 20. The average number of alleles was 8.11, and the expected heterozygosity ranged from 0.62 to 0.94. These microsatellite primers will be useful in population genetics studies for this species.

Key words: Cerrado; Enrichment genomic library; Mangaba; SSR; STR