

## Jaboticaba genotypes: analysis of fruits, seeds, and artisanal wine

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### Supplementary Document

**Table 1.** SSR primers, sequences (forward and reverse), annealing temperature (at), and reason for developing the sequences for species of the family, Myrtaceae, and amplified for *Plinia* sp.

Locus SSR	Primer sequence (5' - 3')		A t ( <sup>o</sup> C )	Motif
	Forward	Reverse		
mPgCIR 18	5' TAAGCTGCATGTGTGC 3'	5' ATGGCTTTGGATGAAA 3'	55	(GA) <sub>23</sub>
mPgCIR 21	5' TGCCCTTCTAAGTATAACA G 3'	5' AGCTACAAACCTTCTA AA 3'	55	(AG) <sub>15</sub> GG(AG) <sub>7</sub>
mPgCIR 97	5' GACCTCAGTAGTTCAGCA TGT 3'	5' TAGAGTGGACGGGAG GAG 3'	50	(GGAA) <sub>3</sub> /(GGTC) <sub>3</sub>
mPgCIR 108	5' GGACCTCACAGAAGTTCA C 3'	5' CGCTGTTTACTGTCTG TT 3'	50	(GA) <sub>13</sub> /(GGAG) <sub>3</sub>
mPgC IR 174	5' GCCACTGTCTAAGAGGA TTG 3'	5' ATTGTGGGAGATTGGA GAC 3'	56	(GA) <sub>20</sub> /(GA) <sub>9</sub> /(GG TC) <sub>3</sub>
mPgC IR 187	5' AAGCCTAATAATGCGGA AGT 3'	5' TCTTTCCAGGATGGAG TA 3'	55	(GGAA) <sub>3</sub>
mPgC IR 198	5' CTCGATCAGAAGAACAA CAC 3'	5' ACTGTTCTGATGGCTC TC 3'	55	(GA) <sub>22</sub> /(GGTTGA) 3
mPgC IR 209	5' CTAAAGCCACATCCAGCA 3'	5' CTAACATTTGCCTTCTA CAGC 3'	55	(GA) <sub>15</sub>
mPgC IR 225	5' ATGGCTCAATGAAGATCC TA 3'	5' GCTCATCCTTGTTAGGA AAG 3'	55	(GT) <sub>9</sub> /(AGAA) <sub>3</sub>
mPgCIR 347	5' CTCTGAAAGGGAGAGG ACT 3'	5' AGAATCTTCGCCTATT GCTT 3'	55	(GA) <sub>6</sub> (GA) <sub>13</sub>
mPgCIR 378	5' GTCACCTGCCTGTTGAT TTT 3'	5' GCTTTCATTCTTA ACCG 3'	55	(CT) <sub>10</sub> (CT) <sub>6</sub>
mPgCIR 446	5' GTGACCACTTTCTCAA GCTA 3'	5' ATCTCCATTCTTGCT CGT 3'	55	(TC) <sub>24</sub> (CA) <sub>9</sub>