

Table S1. TaqMan MGB primers and probes.

Target	Sequence (5' to 3')
5.8S rRNA (<i>C. sinensis</i> JN681150.1; <i>C. grandis</i> JN681156.1)	Forward CGACTCTCGGCAACGGATA Reverse CGCATTTTCGCTACGTTCTTCA Probe CTCGGCTCTCGCATC
CiFT1 (Nishikawa et al. 2007)	Forward CTACCAACAAAATTTTCATCACTTGAATAG Reverse GATCTCTCTCCCTGCTAGACATATCA Probe TCTTACTACTTTTGTAGGCTGTGT
CiFT2 (Nishikawa et al. 2007)	Forward CTACCAACAAAATTTTCATCACTTGAATAG Reverse GGGTCTCTCTCCCTGCTAGACA Probe TCTTACTACTTTTGTAGGTTGTTTG
CiFT3 (Nishikawa et al. 2007)	Forward CAACAAAATTTTCATCACTTGAATAGTC Reverse AAACACTCAACAACACTTAGCACAAA Probe AGGCTGTGTGTGCGTGTA