

S2 Table. Primer sequences used to obtain analysed cpDNA fragments

| assay             | fragment           | primer sequence<br>(5'→ 3')                                                         |
|-------------------|--------------------|-------------------------------------------------------------------------------------|
| CAPS              | <i>trnC-trnD</i>   | F: CCAGTTCAAATCCGGGTGCC<br>R: GGGATTGTAGTTCAATTGGT                                  |
| SNaPshot          | <i>trnC-trnD</i>   | F: CCAGTTCAAATCCGGGTGCC<br>R: GGGATTGTAGTTCAATTGGT                                  |
| SNaPshot          | <i>psbC-trnS</i>   | F: GGTCGTGACCAAGAAACCAC<br>R: GTTTCGAATCCCTCTCTCTC                                  |
| SNaPshot          | <i>trnK1-trnK2</i> | F: GGGTTGCCCGGGACTCGAAC<br>R: CAACGGTAGAGTACTCGGCTTTTA                              |
| length difference | <i>trnC-trnD</i>   | FM13: CACGACGTTGTAAAACGACTGTTTTTCGTTTTCTAGACGTTTCG<br>R: TTGGATTTATGTTTTATTGACTCATT |
| length difference | <i>trnK1-trnK2</i> | FM13: CACGACGTTGTAAAACGACTTGACACAGCTTTCTCTATGT<br>R: GACCCCATAAATGAAATCGAAA         |