

Corrigendum

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Development of microsatellite markers for the annual andromonoecious herb *Commelina communis* f. *ciliata* (Commelinaceae)

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Table 1 on p. 135 should be replaced with the corrected Table 1 shown bellow.

Table 1. Characteristics of microsatellite loci developed for *C. communis* f. *ciliata* based on all 65 samples

Locus	Primer sequences (5'-3') ^a	Repeat motif	Fluorescent label	Size range (bp)	GenBank Accession No.
<i>Ccfc01</i>	F: M13F-CGATCATAGGAAAGTGCACCC R: CGCATTACCAATTGAGTTTGACC	(CGA) ₈	6-FAM	156–182	LC339837
<i>Ccfc02</i>	F: M13R-CGATGGATCGAAGGCGAG R: CTACGTGCCTGCACATTACC	(CGA) ₇	VIC	229–235	LC339838
<i>Ccfc05</i>	F: M13R-CGTTTGATTTGGACGTTGGTC R: GAGCAATGGAAGTTGGAGCC	(ATT) ₈	VIC	171–178	LC339839
<i>Ccfc08</i>	F: M13R-GCATGTTCAATAAGTCCGAAAGG R: TGTACATAGATGACTTGAGCATTGG	(AT) ₈	VIC	177–183	LC339840
<i>Ccfc09</i>	F: AP2-TGGAGCTTCTAAGTCTAAACCG R: TGTCGGGATAAGGGTTCGAG	(AT) ₈	NED	275–296	LC339841
<i>Ccfc19</i>	F: M13F-CGAATGTGGCAAGAAAGGGC R: TAGGTCCACTTCACCTGGC	(GAA) ₇	6-FAM	224–250	LC339842
<i>Ccfc25</i>	F: M13F-TGAGAAGTGATGCCCATATCC R: GCAAGCCCATTATATTGTTTCTCC	(GT) ₉	6-FAM	163–180	LC339843
<i>Ccfc28</i>	F: M13F-GCTTTGCTCCTTGCTAGATCC R: GGTCTTTGTTGGTGACCCCTTG	(AT) ₉	6-FAM	236–258	LC339844
<i>Ccfc29</i>	F: M13R-AACTGGGTCCCAGTGTGG R: GGGATTAATATGGTTGGTCCTTTCC	(ACT) ₇	VIC	222–235	LC339845
<i>Ccfc31</i>	F: M13F-TAGGCCATAAACTGCACGG R: ACACTACTGTAAGCCTCGCC	(TA) ₈	6-FAM	248–264	LC339846
<i>Ccfc32</i>	F: M13R-TCTGTGGACCTCATGTTTAAGG R: GCCTCTGCCCTTCAGCCC	(TA) ₈	VIC	287–307	LC339847
<i>Ccfc37</i>	F: M13F-ACCTCATCCCAGCCAACAG R: CTGGAGGTCCCTTCTAGCC	(AG) ₈	6-FAM	259–278	LC339848
<i>Ccfc38</i>	F: M13R-CAAATCTTCACATGCACTGATAGG R: GATCGACTTGGCGCTTTGG	(AT) ₉	VIC	261	LC339849
<i>Ccfc39</i>	F: AP2-ATTGTTTCTCTGGGTTATAGAGC R: CAACTCTGATACTGGTGAGCG	(AT) ₈	NED	170–184	LC339850
<i>Ccfc44</i>	F: M13R-TTGGACCCCTTGAGCTTGG R: ACCTACTGGTTTCATAACAGGTC	(AT) ₁₀	VIC	323	LC339851

^aOne of the three following primer sequences was added to each forward primer: M13F, 5'-CACGACGTTGTAAAACGAC-3'; M13R, 5'-TGTGGAATTTGTGAGCGG-3'; AP2, 5'-CTATAGGGCACGCGTGGT-3'.