

Jumping through the hoops: the challenges of daffodil (Narcissus) classification

Article

Supplemental Material

Table 2

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Table 2. Summary statistics of the amplified 11 microsatellites based on 450–455 hoop-petticoat daffodil samples. H_o = observed heterozygosity; s.e. = standard error. N indicates number of samples across the table.

Locus	N	Total N of alleles	N of alleles per individual	Alleles per individual (mean $\pm$ s.e.)	H_o	Allelic diversity	Proportion of null alleles
NSB14	455	20	0-4	1.009 ($\pm$ 0.04)	0.262	0.772	0.305
NSB52	452	12	0-3	1.341 ($\pm$ 0.027)	0.301	0.767	0.004
NSB73	452	25	0-4	1.188 ($\pm$ 0.026)	0.212	0.799	0.049
NSB113	455	9	1-4	1.273 ($\pm$ 0.024)	0.244	0.624	0
NSB122	455	13	0-4	1.229 ($\pm$ 0.026)	0.229	0.798	0.031
NSB143	450	8	0-3	1.073 ($\pm$ 0.021)	0.138	0.674	0.067
NSB182	455	5	1-3	1.338 ($\pm$ 0.025)	0.312	0.446	0
NSB232	455	5	1-3	1.246 ($\pm$ 0.025)	0.196	0.353	0
NSB253	455	13	0-3	1.149 ($\pm$ 0.032)	0.301	0.832	0.158
NSB263	452	11	1-4	1.299 ($\pm$ 0.024)	0.274	0.736	0
NSB272	455	16	1-4	1.532 ($\pm$ 0.033)	0.424	0.778	0