

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

Article

Supplemental Material

Table 3

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Table 3. Genetic diversity estimators for each taxon. N_H = the number of different haplotypes N_C = the number of different clusters, A = the total number of different alleles across all loci; A_p = the number of private alleles across all loci; R = the genotypic richness. N without subscript indicates number of samples across the table.

Taxa	plastid DNA N (N_H)	Haplotypes (N)	SSRs N (N_C)	Clusters (N)	A	A_p	R
<i>Bulbocodii sensu Marques</i>							
<i>N. bulbocodium</i>							
	199 (13)	H28 (47), H29 (23), H30 (1), H31 (6), H33 (24), H34 (9), H35 (4), H37 (71), H38 (2), H39 (1), H40 (3), H41 (7), H42 (1)	198 (3)	1.1 (14) 3 (31) 4.1 (153)	116	34	1
<i>Meridionalis</i>							
<i>N. obesus</i>							
	25 (2)	H6 (2), H7 (23)	25 (2)	4.1 (1) 4.2 (24)	14	2	1
<i>N. cantabricus</i>							
	130 (6)	H1 (23), H2 (17), H29 (9), H3 (76), H32 (4), H34 (1)	129 (3)	1.1 (28) 1.2 (75) 1.3 (26)	63	3	0.95
<i>N. hedraeanthus</i> subsp. <i>hedraeanthus</i>							
	1 (1)	H8 (1)	1 (1)	2 (1)	12	0	N/A
<i>N. hedraeanthus</i> subsp. <i>luteolentus</i>							
	16 (1)	H1 (16)	14 (1)	2 (14)	12	2	1
<i>N. romieuxii</i>							
	75 (3)	H1 (28), H3 (33), H4 (14)	74 (2)	1.1 (29) 1.2 (45)	66	8	1
<i>Intersectional hybrids</i>							
<i>N. × cazorlanus</i>							
	2 (2)	H1 (1), h12 (1)	2 (2)	1.3 (1) 2 (1)	13	0	1
<i>N. × fosteri</i>							
	4 (3)	H34 (2), H36 (1), H28 (1)	4 (2)	1.2 (1) 4.1 (3)	30	0	1
<i>N. × litigiosus</i>							
	3 (1)	h12 (3)	3 (1)	1.2 (3)	20	0	1