

The chromosomes of species, hybrids and cultivars of *Narcissus* L. (*Amaryllidaceae*)

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Summary. Chromosome counts are reported for 46 taxa, 8 F₁ hybrids and 230 cultivated varieties of *Narcissus* L. and the percentage pollen viability is given for most of them. B chromosomes were found in several species and cultivars. Chromosome numbers rise to the hexaploid level ($2n = 42, 43$) in species of the subgenus *Narcissus*, but the diploid ($2n = 14$) is the most commonly-occurring level of ploidy. In the subgenus *Hermione* diploid chromosome counts of $2n = 20$ and 22 were obtained ($x = 10, 11$), in confirmation of reports by previous workers. In cultivars solely derived from the subgenus *Narcissus* eu- and aneu-tetraploids ($2n = 28 \pm 2$) greatly exceed diploids in frequency. Cultivars with complex ancestry involving $x = 7$ and $x = 10$ or 11 plants display a wide range of chromosome numbers, and the probable derivation of the chromosome complements of these cultivars and of other real and apparent aneuploids, together with those of some of the more interesting euploids, are given in the light of current knowledge. The levels of ploidy extant in *Narcissus* cultivars are compared and contrasted with those of a range of other genera in which many ornamental cultivars have also been produced.

INTRODUCTION

The genus *Narcissus* L. is native in western Asia and in Europe as far north as Britain and Scandinavia (Fernandes 1951) and has been the subject of a number of chromosomal investigations (Moore 1982), notably the many works by A. Fernandes (summarised by Jefferson-Brown 1969). The majority of the species (subgenus *Narcissus*) have chromosome numbers based on $x = 7$ and most of these are diploids ($2n = 14$), but polyploids ascending through all the possible euploid intermediates to octoploid ($2n = 8x = 56$) have been reported in *N. bulbocodium* (Fernandes 1967). Occasional triploids ($2n = 21$) and tetraploids ($2n = 28$) occur in other species (Fedorov 1969, Moore 1982).

Not closely related cytologically to the above, the subgenus *Hermione* (Haw.) Spach comprises a group of species with basic numbers other than seven, being $x = 5$, giving $2n = 2x = 10$ and $2n = 6x = 30$ in *N. serotinus* and perhaps $2n = 4x = 20$ in *N. tazetta*, with a secondary alteration to $2n = 22$ in some races of the same species and also in related species such as *N. papyraceus* (= *N.* 'Paper White'), *N. pachybolbus* and others (Maugini 1953, Fernandes 1966, Weitz & Feinbrun 1972). The changes to $2n = 20$ and 22 are clearly very ancient, because these species behave strictly as diploids, forming only bivalents at meiosis (Fernandes 1950), and they will be referred to as diploids in this paper.

The genus is remarkable for the tremendous proliferation of cultivated varieties which has occurred, principally during the present century. These have been documented carefully by Dr T. Throckmorton of Des Moines, Iowa, U.S.A., and an abridged list has been published recently (Throckmorton 1980) which gives the name, breeder, year of introduction and coded entries summarising the size, flower morphology and colour, height and flowering

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time of each cultivar. The work is based on a computerised data bank covering over 10,000 cultivars held by the American Daffodil Society. This is more comprehensive and gives in addition to the above the parentage of each cultivar, its breeder and its chromosome number, if known. As chromosome counts are not given in Throckmorton (1980) the printout of the data bank obtained from the Iowa Methodist Medical Center, 1200 Pleasant Street, Des Moines, Iowa 50309, U.S.A.) was taken as the source of some of the chromosome data used in this article, and will be referred to below as A.D.S. (1981).

Although so many thousands of cultivars of *Narcissus* have been described, chromosome numbers are known for relatively few of them, amounting to about 300 (A.D.S. 1981, Fernandes & de Almeida 1971). The origins of these cultivars are very diverse and must include hybridisations, not only between diploids, but also between individuals of similar or differing levels of polyploidy (with the inevitable occurrence of aneuploid individuals among the progeny) and between individuals with differing basic chromosome numbers ($x = 7, 10$ or 11). Most of these hybridisations would have been made by breeders who had no idea of the chromosome complements of their plants, so the *Narcissus* cultivars constitute an ideal example of chromosome variation and evolution in a genus as influenced by the largely unknowing selection pressures exerted by Man.

In order to obtain more information on this variation the somatic chromosomes of over 200 cultivars were examined in the present study, the results of which will be presented below and analysed in conjunction with previous reports by other authors. While this work was under way the opportunity arose for us to examine a number of species (both in subgenus *Hermione* and in subgenus *Narcissus*) and F_1 hybrids between species of *Narcissus*, the chromosomes of some of which are unknown. These results are also included here.

The great majority of *Narcissus* cultivars are of hybrid origin and must therefore be propagated vegetatively in order that their identity can be retained. It follows that, assuming correct naming in every instance, any cultivar bearing the same name as one mentioned here which is used by anyone for further breeding will actually be the same clone as the one which we have studied and will consequently have the same chromosome constitution and behaviour. It was thus considered that it would be of considerable value to those intending to use particular cultivars in breeding programmes to report the pollen fertility of as many plants as possible, the pollen fertility being a good measure of the probable degree of success to be expected when a particular plant is used as a pollen parent in a trial cross. It will also give a similar measure of a plant's likelihood of success as a seed parent, except in those cultivars with double flowers (division 4, see below), many of which have no anthers and consequently zero pollen fertility. These can quite possibly possess a completely normal and fertile gynoecium and be fully seed-fertile.

This article will therefore present data on pollen fertility and chromosome number, the latter comprising 46 taxa, 8 F_1 hybrids and 230 cultivars, in order to increase the published knowledge of the cytology of the genus.

MATERIAL AND METHODS

Bulbs were obtained from three of Britain's major *Narcissus* collections; the Royal Horticultural Society's Garden, Wisley, the Rosewarne Experimental Horticulture Station and the National Collection at the New University of

Ulster, Coleraine; F_1 hybrids, mostly between miniature species, were given by Mr J. Blanchard, and the living collection at RBG Kew was also sampled.

Plants were grown in pots placed in cold frames and root tips were collected for chromosome study during periods of maximum root growth, September to November and February to March. Root tips were pretreated for 24 hours at 4°C in saturated aqueous alpha-bromonaphthalene and fixed in 1:3 acetic alcohol. After hydrolysis in 1N HCl at 60°C for 7 min. they were stained in Feulgen. Squashes were made in 45% acetic acid, or if the Feulgen stain was too faint, in 1% aceto-orcein. Each slide was frozen with liquid CO₂ at -75°C and the coverslip removed with a sharp blade. Both slide and coverslip were dehydrated in absolute ethanol and recombined with a drop of Euparal (B.D.H.). After a period of 2 weeks at 45°C for the mountant to harden the slides were screened with a Wild M20 microscope and suitable cells were photographed with a Zeiss photomicroscope. Slides are retained in the collection of the Cytology department of the Jodrell Laboratory.

For pollen fertility estimates mature dehisced anthers were tapped out on a slide in 1% cotton blue in lactophenol, the large debris removed and a coverslip applied. Stainable, presumed viable, pollen grains were expressed as a percentage of the total number of grains scored, the minimum being 500 for any single analysis.

Representative voucher specimens of leaves and flowers have been deposited in the Kew Herbarium.

CLASSIFICATION OF *Narcissus* CULTIVARS

The 10,000 (approx.) cultivars of *Narcissus* have been grouped for convenience by the Royal Horticultural Society into twelve divisions (Throckmorton 1980). As these are referred to frequently in this article it is appropriate to summarise them here as follows:

Division 1, Trumpet daffodils. Solitary flowers with corona equal to or longer than the perianth segments.

Division 2, Long-cupped daffodils. Solitary flowers with corona one third to almost equal to the length of the perianth segments.

Division 3, Short-cupped daffodils. Solitary flowers with corona less than one third the length of the perianth segments.

Division 4, Double-flowered daffodils of any affinity among Divisions 1-3, 5-9 or 11-12.

Division 5, Triandrus daffodils. Characteristics of *N. triandrus* dominant in hybrid.

Division 6, Cyclamineus daffodils. Characteristics of *N. cyclamineus* dominant in hybrid.

Division 7, Jonquilla daffodils. Characteristics of *N. jonquilla* dominant in hybrid.

Division 8, Tazetta daffodils. Characteristics of *N. tazetta* dominant in hybrid.

Division 9, Poeticus daffodils. Characteristics of *N. poeticus* dominant in hybrid.

Division 10, Species, wild forms and wild hybrids, including double-flowered forms.

Division 11, Split-corona daffodils. Corona divided distally for at least one third of its length into segments.

Division 12, Miscellaneous daffodils not falling into Divisions 1-11.

All divisions are plants of garden origin except Division 10. Within each

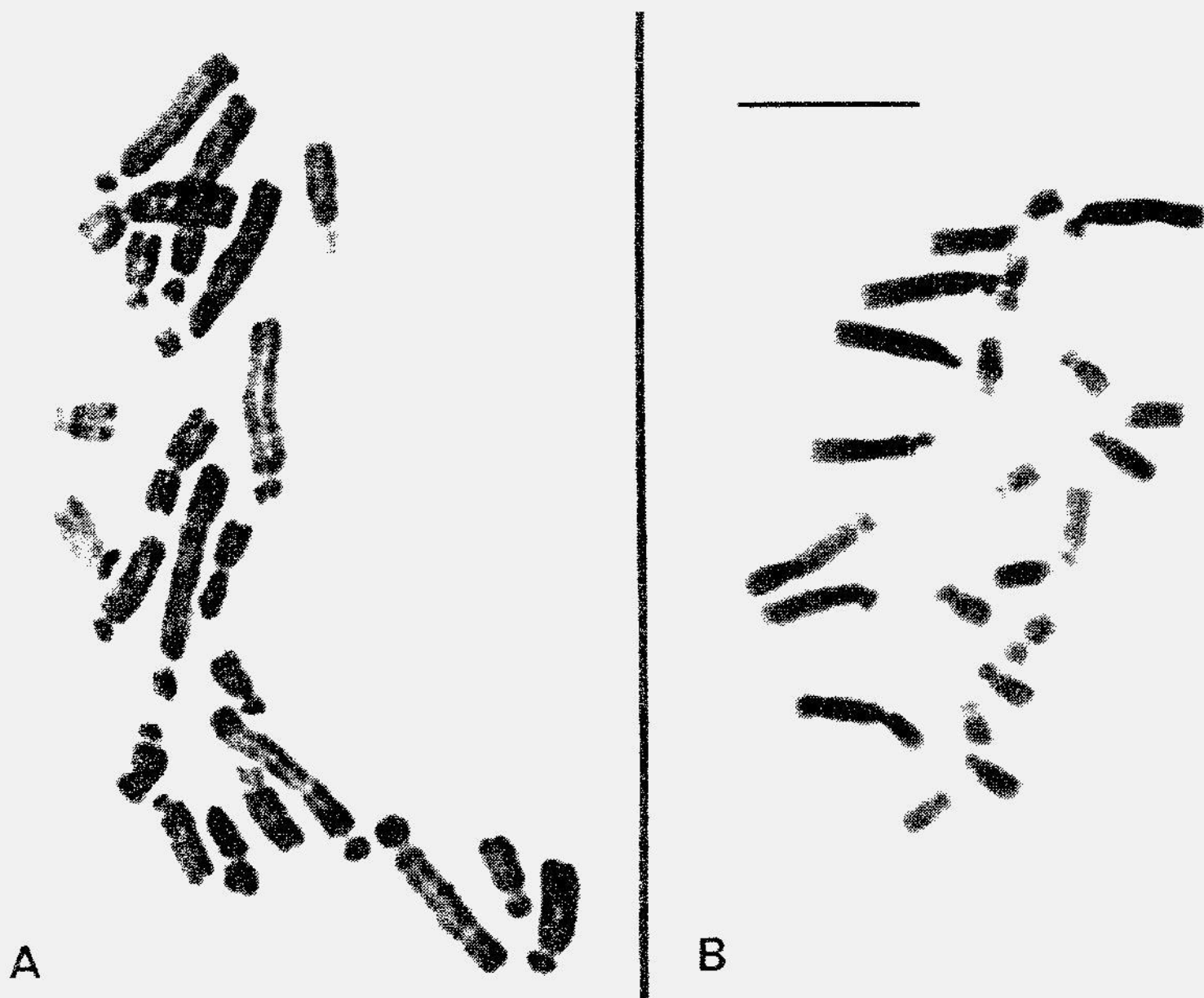


FIG. 1. Somatic chromosomes of *Narcissus* subgenus *Hermione*. **A** *N. aff. tazetta*, $2n = 22$; **B** *N. papyraceus*, $2n = 22$. Scale = $10\mu\text{m}$.

division flowers are colour-coded, firstly for the colour of the perianth and secondly for that of the corona, the latter being divided into two or three regions from the centre to the rim if two or more colour bands are present. Colour codes are W (white or nearly so), G (green), Y (yellow), P (pink), O (orange), and R (red). Under this system the cultivar 'Merlin', for example, is classified as 3W-YZR (Throckmorton 1980), which indicates that it has a solitary flower with a white perianth and a short yellow corona with a red rim. A second example 'Fermoy' (2W-YOO) has solitary flowers of which the corona has a yellow base but is predominantly orange and is not as long as the white perianth segments.

SPECIES AND F_1 HYBRIDS

Chromosome numbers

These are given in Table 1 and some of them are illustrated in Figs. 1–5. It is not intended here to describe in detail the karyotypes of every accession because this would largely duplicate the thorough work of Fernandes. Instead, description and comment will be restricted to features of particular interest.

Of the members of the subgenus *Hermione*, *N. pachybolbus*, *N. papyraceus* and *N. aff. tazetta* were found to have $2n = 22$ chromosomes and *N. tazetta* to have $2n = 20$, confirming earlier counts by Maugini (1953), Fernandes (1966) and Weitz & Feinbrun (1972). The karyotypes of these species tend towards bimodality, with 8 very large acrocentrics and 14 smaller acro- or metacentrics in the $2n = 22$ forms (Figs. 1A, 1B). It was not possible to analyse the karyotype of the $2n = 20$ plant, but Fernandes (1966) illustrated a chromosome

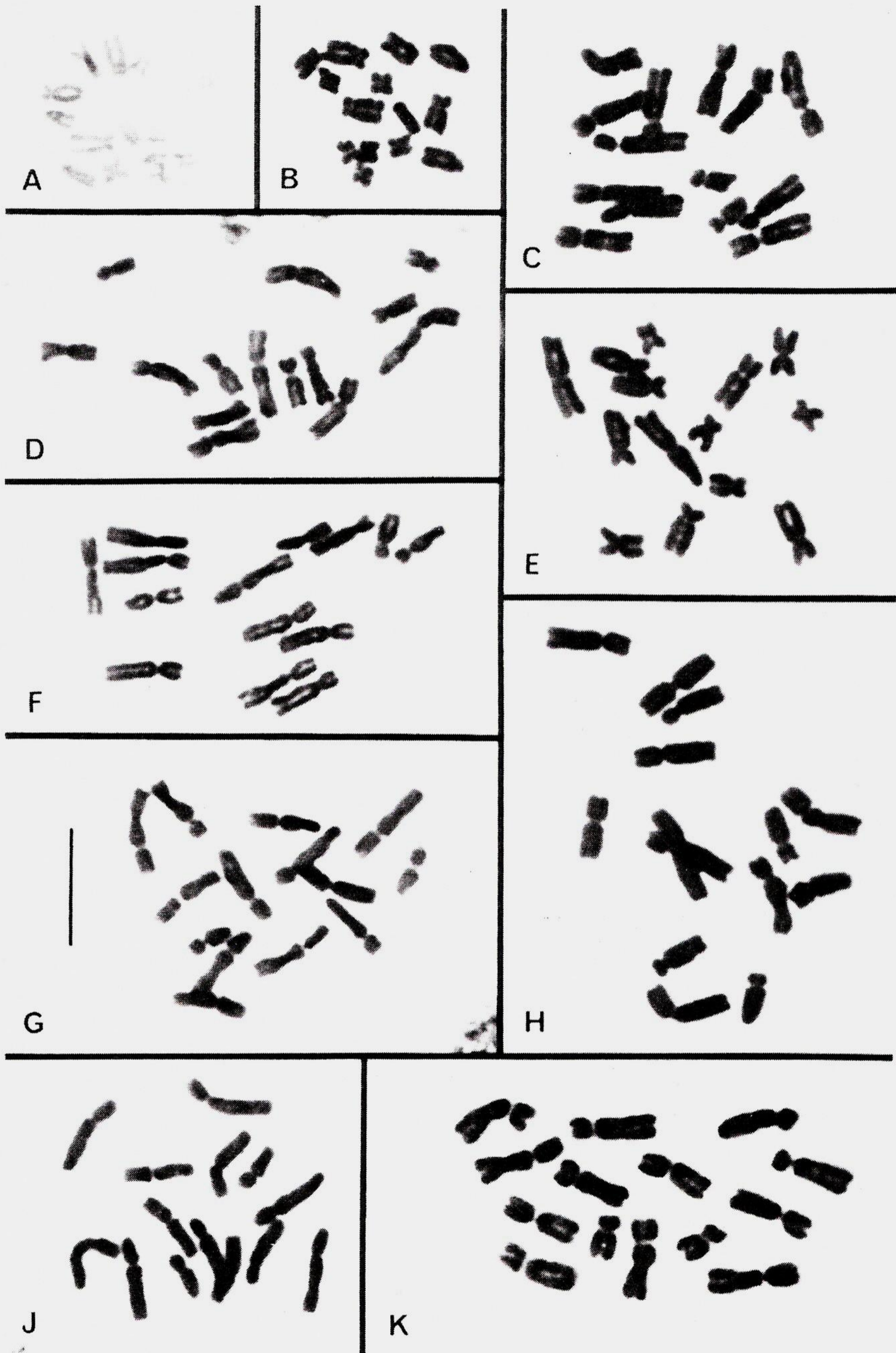


FIG. 2. Somatic chromosomes of diploid *Narcissus*, all $2n = 14$. **A** *N. hedraeanthus*; **B** *N. bulbocodium* var. *conspicuus*; **C** *N. longispathus*; **D** *N. juncifolius*; **E** *N. calcicola*; **F** *N. cyclamineus*; **G** *N. nanus*; **H** *N. jonquilla*; **J** *N. hispanicus*; **K** *N. poeticus*. Scale = $10\mu\text{m}$.

complement for *N. tazetta* ($2n = 20$) which was markedly different from the above, comprising 12 large acrocentrics and 8 smaller chromosomes.

The members of the subgenus *Hermione*, with $x = 10$ or 11 chromosomes are

TABLE 1. Chromosome numbers and pollen fertility in *Narcissus* species, varieties and interspecific F₁ hybrids.

Taxon	Kew accession number	Origin	2n	Pollen Fertility %
<i>N. bulbocodium</i> L. var. unknown	295-82.03045	Unknown	14	97
" " " "	295-82.03047	Unknown	26	99
" " " "	232-78.02456	Unknown	42	
" " var. <i>albidus</i> (Emb. & Maire) Maire	341-82.03524	Blanchard	14	
" " var. <i>citrinus</i> Baker	341-82.03525	Blanchard	34	86
" " " " "	400-82.04500	Coleraine	34	89
" " var. <i>conspicuus</i> Baker	341-82.03526	Blanchard	14	82
" " " " "	400-82.04501	Coleraine	14	87
" " var. <i>filifolius</i> Hort.	341-82.03527	Blanchard	26	99
<i>N. calcicola</i> Mendonça	295-82.03046	Unknown	14	
<i>N. cantabricus</i> DC. var. <i>cantabricus</i>	CYT 82.1498	Walker	14	
" " var. <i>foliosus</i> Maire	400-82.04502	Coleraine	28	
<i>N. compressa</i> Haw.	257-83.03161	Rosewarne	31	
<i>N. cyclamineus</i> DC.	400-82.04503	Coleraine	14 + B	86
<i>N. fernandesii</i> G. Pedro	326-79.02967	Mathew	21	
" " " "	341-82.03529	Blanchard	28	99
" " " "	400-82.04504	Coleraine	28	83
<i>N. gayi</i> (Henon.) Pugsl. (= <i>N. pseudonarcissus</i> L.)	400-82.04528	Coleraine	14	
<i>N. hedraeanthus</i> (Webb & Heldr.) Colmeiro	341-82.03530	Blanchard	14	61
<i>N. hispanicus</i> Gouan	400-82.04505	Coleraine	14	
<i>N. jonquilla</i> L. var. <i>jonquilla</i>	400-82.04506	Coleraine	14	97
" " var. <i>henriquesii</i> Samp.	341-82.03531	Blanchard	14	41
" " " " "	400-82.04507	Coleraine	14	
<i>N. juncifolius</i> Req. ex Lag.	400-82.04508	Coleraine	14	65
<i>N. lobularis</i> Schult.	400-82.04509	Coleraine	42	30
<i>N. longispathus</i> Pugsley	400-82.04510	Coleraine	14	76

Taxon	Kew accession number	Origin	2n	Pollen Fertility %
<i>N. minimus</i> Kunth	400-82.04511	Coleraine	14 + B	
<i>N. nanus</i> Steudel	400-82.04512	Coleraine	14	96
<i>N. ornatus</i> Haw. var. <i>ornatus</i> (= <i>N. poeticus</i> L.)	400-82.04519	Coleraine	14	
" " " " " "	257-83.03187	Rosewarne	14	
" " var. <i>maximus</i> Hort.	400-82.04520	Coleraine	14	96
" " " " " "	399-82.04463	Rosewarne	14	99
<i>N. pachybolbus</i> Durieu	400-82.04515	Coleraine	22	
<i>N. papyraceus</i> Ker-Gawl.	400-82.04516	Coleraine	22	97
<i>N. poeticus</i> L. var. <i>poeticus</i>	011-81.00043	<i>Sands</i>	14	95
" " var. <i>flore pleno</i> Hort.	400-82.04517	Coleraine	14	
" " " " " "	399-82.04459	Rosewarne	14	
" " subsp. <i>radiiflorus</i> (Salisb.) Baker	400-82.04523	Coleraine	14	
" " " " var. <i>poetarum</i> Burb. & Baker	400-82.04521	Coleraine	14	94
" " " " " " " " " "	257-83.03189	Rosewarne	14	
<i>N. pseudonarcissus</i> L. subsp. <i>abscissus</i> (Schultes & Schultes fil.) A. Fernandes	400-82.04527	Coleraine	14	
" " subsp. <i>obvallaris</i> (Salisb.) A. Fernandes	400-82.04530	Coleraine	21	
" " subsp. <i>pallidiflorus</i> (Pugsf.) A. Fernandes	232-78.02460	Unknown	14	
" " subsp. <i>pseudonarcissus</i>	232-78.02458	<i>A. & M.H. 7805</i>	43	
" " " " " "	400-82.04526	Coleraine	14	
<i>N. pumilus</i> Salisb. var. <i>plenus</i> Hort. (= 'Rip van Winkle')	400-82.04531	Coleraine	21	
<i>N. recurvus</i> Haw. (= <i>N. poeticus</i> L.)	400-82.04524	Coleraine	21	
" " " " " "	399-82.04498	Rosewarne	21	40
<i>N. romieuxii</i> Braun-Blanquet & Maire var. <i>romieuxii</i>	341-82.03532	<i>Blanchard</i>	28	
" " " " " "	400-82.04532	Coleraine	28	
" " var. <i>mesatlanticus</i> Hort.	341-82.03533	<i>Blanchard</i>	28	97

TABLE 1 *continued*. Chromosome numbers and pollen fertility in *Narcissus* species, varieties and interspecific F₁ hybrids.

Taxon	Kew accession number	Origin	2n	Pollen Fertility %
<i>N. romieuxii</i> Braun-Blanquet & Maire var. <i>mesatlanticus</i> Hort.	400-82.04533	Coleraine	28	77
„ „ var. <i>tananicus</i> Maire	341-82.03534	Blanchard	14	
<i>N. rupicola</i> Duf.	356-78.03601	Unknown	14	
<i>N. tazetta</i> L.	135-79.01296	Chesterman 103	20	
<i>N. aff. tazetta</i> L.	364-84.03754	Willis	22	
<i>N. telamonius</i> Link var. <i>plenus</i> Hort. (= <i>N. pseudonarcissus</i> L.)	257-83.03194	Rosewarne	14	
<i>N. triandrus</i> L. var. <i>albus</i> (Haw.) Baker	CYT.82.362	Blom	14	
„ „ „ „ „ „	CYT.83.935	Blanchard	14 + 0 - 3B	
„ „ var. <i>concolor</i> (Haw.) Baker	CYT.83.934	Blanchard	14	
„ „ var. <i>pulchellus</i> (Salisb.) Baker	CYT.83.936	Blanchard	14	
<i>N. willkommii</i> (Samp.) A. Fernandes	341-82.03535	Blanchard	14	79
„ „ „	400-82.04534	Coleraine	14	
F ₁ hybrids				
<i>N. atlanticus</i> F. C. Stern × <i>N. asturiensis</i> (Jord.) Pugsley 'Navarre'	341-82.03536	Blanchard	14 + B	3
<i>N. atlanticus</i> F. C. Stern × <i>N. cuatrecasasii</i> Casas, Lainz & Ruiz Rejon	341-82.03537	Blanchard	14	69
<i>N. jonquilla</i> L. × <i>N. watieri</i> Maire	341-82.03538	Blanchard	14	0
<i>N. poeticus</i> L. × <i>N. pseudonarcissus</i> L. (= <i>N. × incomparabilis</i> Miller)	232-78.02455	A. & M. H. 7854	14	
<i>N. pseudonarcissus</i> L. × <i>N. jonquilla</i> L. (= <i>N. × odorus</i> L. var. <i>plenus</i> Hort.)	400-82.04513	Coleraine	14	
„ „ „ „ (= <i>N. × odorus</i> L. var. <i>rugulosus</i> Hort.)	291-82.02815	Wisley	14	2
„ „ „ „ „ „	400-82.04514	Coleraine	21	
<i>N. rupicola</i> Duf. × <i>N. asturiensis</i> (Jord.) Pugsley 'Navarre'	341-82.03540	Blanchard	14 + B	0
<i>N. rupicola</i> Duf. × <i>N. gaditanus</i> Boiss. & Reut.	341-82.03541-3	Blanchard	14	0
<i>N. watieri</i> Maire × <i>N. calcicola</i> Mendonça	341-82.03544	Blanchard	14	
„ „ × <i>N. gaditanus</i> Boiss. & Reut.	341-82.03545-6	Blanchard	14	0.5
„ „ × <i>N. scaberulus</i> Henriq.	341-82.03547	Blanchard	14	

at variance with the remainder of the genus (subgenus *Narcissus*), of which all species are based on $x = 7$. Subgenus *Hermione* was said by Maugini (1953) to be related to the $x = 7$ group through the stabilisation or 'diploidisation' of triploids with $2n = 3x = 21$ chromosomes after aneuploid changes to $2n = 20$ or 22. This seems to be an extremely unlikely explanation, particularly in the light of the discovery of $2n = 10$ in *N. serotinus* (Fernandes 1975), which established a low basic number of $x = 5$ in the subgenus *Hermione* and consequently a tetraploid level in $2n = 20$ races of *N. tazetta*, which is of ancient origin since the species is functionally a diploid. Fernandes (1975) further suggested that the $x = 11$ forms are derived from the hybridisation of a $2n = 2x = 10$ plant with a $2n = 2x = 12$ ancestral type followed by chromosome doubling to give an allotetraploid with $2n = 22$.

The remainder of the genus, subgenus *Narcissus*, has $x = 7$ chromosomes, suggested by Fernandes (1975) also to have arisen from an $x = 6$ ancestral stock through aneuploid change. The species are not chromosomally uniform, however, unlike some other plant groups such as the *Liliaceae-Aloineae* (Brandham 1971), and representative diploids with $2n = 14$ are illustrated in Fig. 2. There is a very wide variation in total chromosome size throughout these plants, with the smallest chromosomes in *N. bulbocodium* and *N. hedraeanthus* (Figs. 2A, 2B). The size increases through a range of intermediates to the longest, occurring in *N. longispathus* (Fig. 2C), *N. jonquilla* (Fig. 2H) and *N. poeticus*. *N. poeticus* has the most massive chromosomes of all, due largely to their increased thickness (Fig. 2K), the total chromosome length being little more than that of other species with large chromosomes.

Throughout most of the species in subgenus *Narcissus* the chromosomes display a range of chromosome length within a karyotype of about 2- or 2.5-fold between the longest and the shortest, although the chromosomes of *N. cyclamineus* vary less than this range (Fig. 2F) and those of *N. calcicola* a little more (Fig. 2E). The chromosomes are submetacentric in most cases, although all the short chromosomes of *N. bulbocodium* (Fig. 2B), *N. hedraeanthus* (Fig. 2A) and, unusually, one of the long pairs in each of *N. juncifolius* (Fig. 2D) and *N. calcicola* (Fig. 2E) are metacentric. In a few other species one or more of the shorter chromosome pairs are also metacentric (e.g. the second shortest pair in *N. cyclamineus*, Fig. 2F).

Polyploids

These occurred sporadically throughout the species studied, most polyploids being tetraploids with $2n = 28$. The greatest range of ploidy was in *N. bulbocodium*, in which all the ploidy levels between diploid ($2n = 14$) and hexaploid ($2n = 42$) were found with the exception of the triploid ($2n = 21$). This reflects the finding by Fernandes (1967) that this species has the widest range of ploidy of any in the genus, ranging from diploid to octoploid ($2n = 14-56$). In the present study triploids were recorded in several species, e.g. *N. fernandesii*, *N. pseudonarcissus* subsp. *obvallaris*, *N. pumilus* var. *plenus* and *N. recurvus* (Table 1), but it is not known to what extent these are representative of vegetatively-reproducing triploid populations similar to those known to occur in *Haworthia* (Brandham & Cutler 1981), or whether they were selected by plant collectors as being more vigorous than the rest of the individuals in the presumably diploid populations in which they occurred. In *N. bulbocodium* the former alternative seems to apply because Fernandes (1967) found two entirely triploid populations in an extensive survey of this species in Portugal. Hexa-

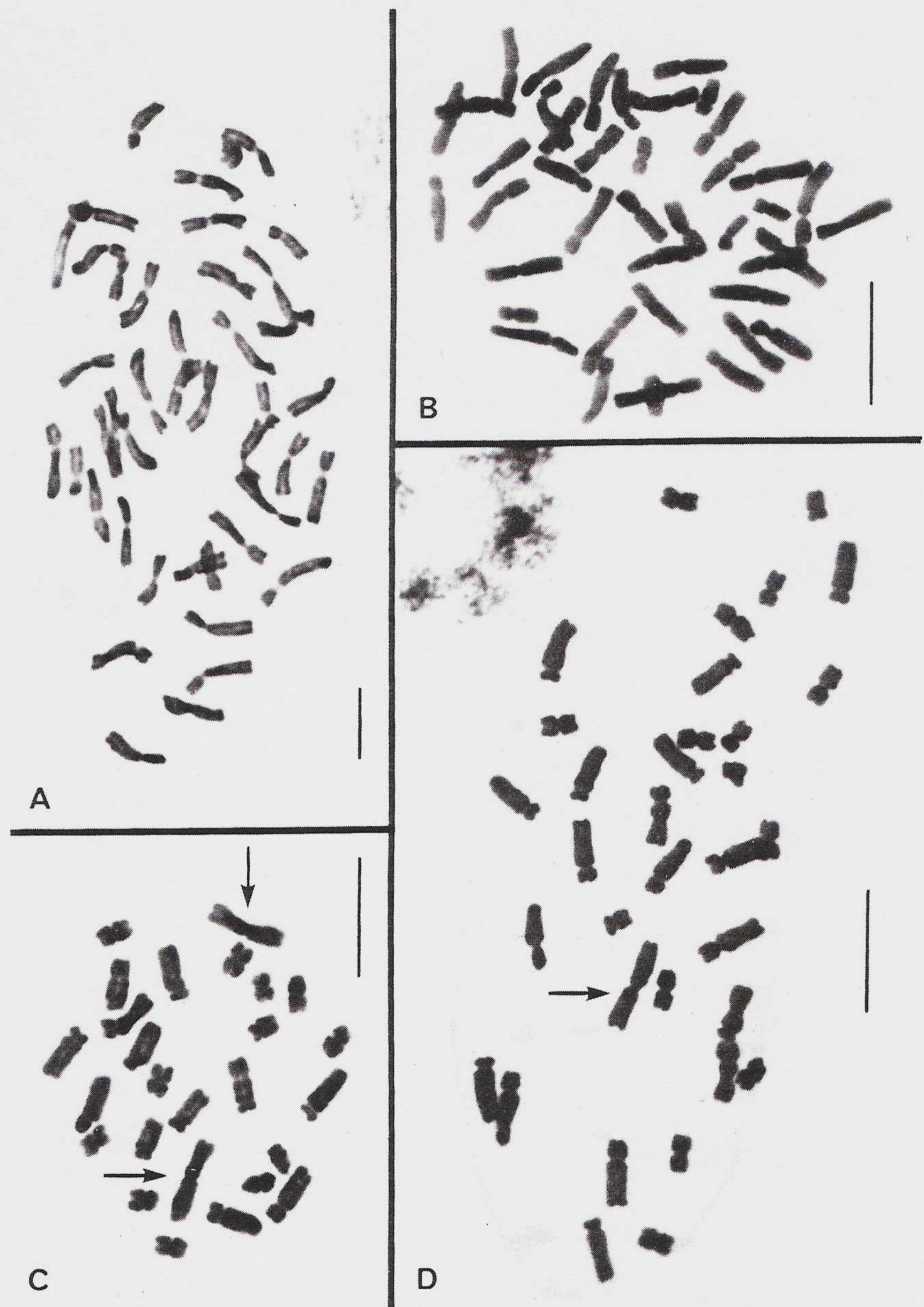


FIG. 3. Somatic chromosomes of polyploid *Narcissus*. **A** *N. pseudonarcissus*, $2n = 6x + 1 = 43$; **B** *N. lobularis*, $2n = 6x = 42$; **C** *N. bulbocodium*, $2n = 4x = 26$, derived from $2n = 28$ by homozygosity for a Robertsonian fusion producing 2 metacentrics (arrows) from 4 acrocentrics; **D** *N. bulbocodium* subsp. *bulbocodium* var. *citrinus*, a pentaploid with the chromosome number reduced from $2n = 35$ to $2n = 34$ by hybridity for the same Robertsonian fusion as in **C** (metacentric arrowed). Scales = $10\mu\text{m}$.

ploids or near-hexaploids were found in three species (Table 1), *N. bulbocodium*, *N. lobularis* (Fig. 3B) and *N. pseudonarcissus* subsp. *pseudonarcissus*.

Aneuploidy

This is uncommon in these wild-collected plants. A single instance occurred at the hexaploid level, with $2n = 6x + 1 = 43$ being found in *N. pseudonarcissus* subsp. *pseudonarcissus* (Fig. 3A). It is to be expected that aneuploidy should be more common in high polyploids. This is because the harmful effects, caused by the extra or missing chromosomes, on the aneuploid gametes that give rise to aneuploid individuals are extreme in haploid gametes but are progressively reduced in diploid and polyploid gametes because of the buffering effect of the presence of one or more extra intact sets of chromosomes. Thus in $x = 7$ plants aneu-haploid gametes with $n = x \pm 1 = 6$ or 8 chromosomes are less viable than aneu-diploid gametes with $n = 2x \pm 1 = 13$ or 15 chromosomes and these are less viable than aneu-triploid gametes with $n = 3x \pm 1 = 20$ or 22 chromosomes. The euploid and aneuploid gametes ($n = 21$ and 22 respectively) that gave rise to the $2n = 43$ aneu-hexaploid would have been almost equally viable, and populations of hexaploids such as the one from which our specimen came should therefore contain a significant number of aneuploids.

The apparent aneu-tetraploids reported in Table 1, $2n = 26$ ($?4x - 2$) in *N. bulbocodium* and $2n = 31$ ($?4x + 3$) in *N. compressa* and the apparent aneu-pentaploid $2n = 34$ ($?5x - 1$) in *N. bulbocodium* are the results of chromosomal events other than aneuploidy. Taking firstly the case of *N. compressa* with $2n = 31$ (Fig. 10A), this plant was described as a species and appears in Division 10 in Throckmorton (1980), so it has been included in Table 1. Nevertheless it is clearly allied morphologically to *N. tazetta* and is certainly of hybrid origin. It will be considered again below, together with the other hybrids and polyploids involving *N. tazetta*. Robertsonian fusion is the origin of the apparently aneuploid $2n = 26$ and 34 individuals of *N. bulbocodium*.

Robertsonian fusion

This is the process whereby two usually non-homologous acrocentric chromosomes each break at the centromere and the arms fuse in a new combination, the two short arms constituting a short metacentric and the long arms a long metacentric. Sexual reproduction following this change will result in the production of some viable gametes carrying the two metacentrics instead of the two original acrocentrics. Fusion of two such gametes produces a homozygote for the fusion in which four acrocentrics (two homologous pairs) are replaced by two long and two short metacentrics. Frequently Robertsonian fusion is accompanied by loss of the short arms, so the homozygote has a net loss of two chromosomes, four original acrocentrics being replaced by two larger metacentrics comprising the four long arms.

In tetraploids of *N. bulbocodium* homozygosity (or more accurately for tetraploids, duplex heterozygosity) for a single Robertsonian fusion, with loss of short arms, reduces the chromosome number from $2n = 28$ to $2n = 26$ including two metacentrics (Fig. 3C). The same event was reported over 50 years ago by Fernandes (1934) in the same species, and the present material is probably descended from plants brought into cultivation at that time. These $2n = 26$ plants are fully fertile and produce gametes with $n = 13$ chromosomes including one metacentric, and when they hybridise with a normal hexaploid of the same species ($2n = 42$) which produces gametes with $n = 21$ chromosomes (lacking the large metacentric) the resulting pentaploid plant contains

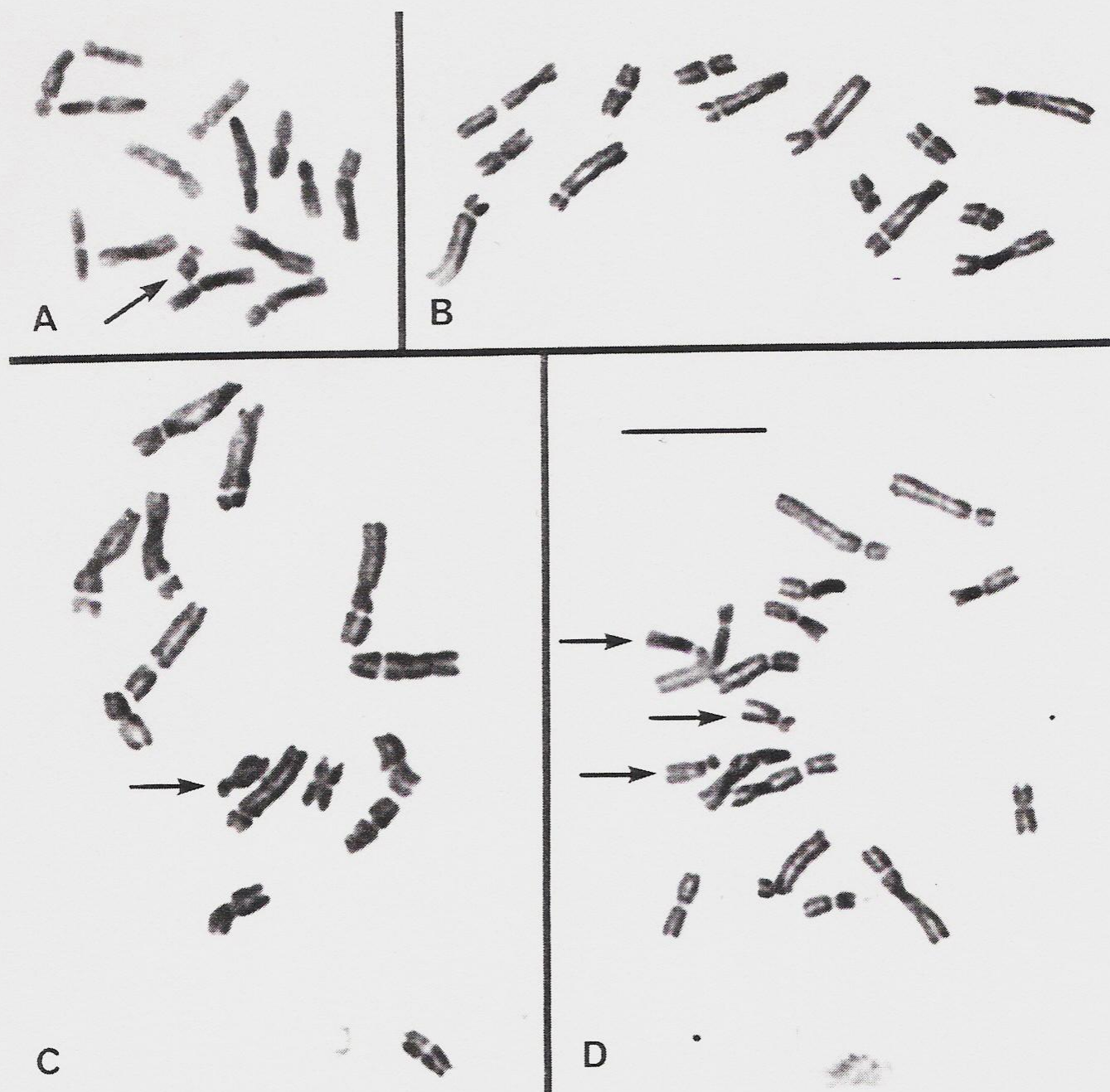


FIG. 4. Somatic chromosomes of diploid *Narcissus* ($2n = 14$) containing B chromosomes (arrows). **A** *N. minimus* (1B); **B, C, D** *N. triandrus* var. *albus* with (B), 0B, (C) 1B, (D) 3B. The Bs are constant within a plant but vary within the population. Scale = $10\mu\text{m}$.

$2n = 13 + 21 = 34$ chromosomes including just one large metacentric. One such plant was located in the present study (Fig. 3D), the first time that this event has been recorded in pentaploid *N. bulbocodium*. It is probable that this individual arose through a chance hybridisation in cultivation between Fernandes' original Robertsonian fusion tetraploid homozygote and a normal hexaploid of unknown provenance.

B chromosomes

These are additional to the normal or A chromosome complement of a species. They can be heterochromatic or euchromatic and are usually but not always smaller than the A chromosomes. They have few functional genes and often pair poorly during meiosis. They are consequently either absent or duplicated in the gametes of plants which carry them and thus vary in numbers from individual to individual within a population. Sometimes they will even vary in numbers from cell to cell within an individual because of mitotic irregularities, although this is very much less common than inter-individual variation.

In *Narcissus* they have been reported in several species and the numbers vary

from one to as many as five in *N. bulbocodium* (Wylie 1952, Jones & Rees 1982), with low numbers being the most common.

In the present study they were seen in three species: *N. cyclamineus* ($2n = 14 + 1B$), *N. minimus* ($2n = 14 + 1B$) and *N. triandrus* var. *albus* ($2n = 14 + 0 - 3B$). All of these are apparently euchromatic, staining identically to the A chromosomes. The single Bs found in *N. cyclamineus* and *N. minimus* resemble each other closely. They are acrocentric and are considerably shorter than the shortest A chromosomes (Fig. 4A). In *N. triandrus* var. *albus* the Bs are much larger acrocentrics, being longer than the shortest pair of A chromosomes, and varied in numbers from plant to plant. In a sample of eight bulbs that were examined, six had no Bs (Fig. 4B), one had a single B (Fig. 4C) and one had three (Fig. 4D). A short acrocentric B was also observed in Blanchard's F_1 hybrids between *N. atlanticus* and *N. asturiensis* 'Navarre' (Fig. 5A) and between *N. rupicola* and *N. asturiensis* 'Navarre' (Table 1), indicating that 'Navarre' also contains $2n = 14 + 1B$ chromosomes.

*F*₁ hybrids

Most of the F_1 hybrids studied in this investigation were crosses between very dissimilar miniature species which were carried out by J. Blanchard. The hybrid *N. atlanticus* × *N. asturiensis* 'Navarre' is typical of the majority of them and shows a very high frequency of morphologically unique chromosomes (Fig. 5A), and it is not possible to determine the homologies of these chromosomes simply from an examination of their length and centromere position, so it would be meaningless to attempt to construct a karyotype by pairing apparently similar chromosomes in this case.

Chromosome variation and heteromorphy is even more marked in another F_1 hybrid, *N. rupicola* × *N. gaditanus*, of which three accessions were studied, all with $2n = 14$. Taking as an example just the easily-recognisable large metacentrics which occur in these plants, no.03541 has two similar ones (Fig. 5C), 03542 has two dissimilar ones with one much longer than the other (Fig. 5D) and 03543 has three similar ones (Fig. 5E). It is evident that one or both of the parents of this hybrid contain heteromorphic homologous chromosomes which have segregated in various ways during gamete formation.

Another diploid F_1 hybrid between very dissimilar species, *N. pseudonarcissus* × *N. jonquilla* (= *N. odoratus* 'rugulosus') is also quite heteromorphic in its chromosome constitution, possessing several chromosomes having no closely-matching homologues. Indeed, the presence of one large and one small metacentric in this plant (Fig. 5B) in a complement of chromosomes which is otherwise largely submetacentric-acrocentric suggests the presence of a Robertsonian fusion (see above), and it is possible that such a fusion played a part in the chromosomal differentiation of the two parental species from the original ancestral *Narcissus* stock. Nevertheless, it would be necessary to examine meiosis in this plant before Robertsonian fusion hybridity could be confirmed or discounted. If present, the fusion would lead to the pairing of the large and small metacentric with two of the acrocentrics to form a quadrivalent. It has not been possible to carry out this particular investigation because of shortage of material.

The extreme heteromorphy found in these examples of F_1 hybrids indicates clearly that a considerable amount of alteration has occurred in chromosome morphology (rather than number) during speciation in the $x = 7$ subgenus

Narcissus and confirms the obvious chromosomal differences between species that can be seen in Fig. 2.

Pollen fertility

This is high in most of the species that were analysed, as might be expected in what are mostly the clonal descendants of wild-collected non-hybrid plants. Table 1 shows that the great majority of them were 80–99% pollen-fertile, even, surprisingly, the pentaploid Robertsonian fusion heterozygote *N. bulbocodium* subsp. *bulbocodium* var. *citrinus* which would be expected to be less fertile. The relatively low fertility of the diploid *N. jonquilla* var. *henriquesii* (03531) is also unexpected since no heteromorphy for any major structural change could be detected in its chromosome complement. It is therefore possible that the fertility of this plant is reduced by changes at the genic level or at the chromosome level, the latter necessarily being symmetrical and consequently irresolvable. The moderate fertility of the triploid *N. recurvus* is typical of autotriploids in which chromosome segregation is irregular, producing many aneuploid gametes, most of which are non-viable (Brandham 1982).

The remaining plant with low fertility, the hexaploid *N. lobularis*, is only one example of a wide range of fertilities that can exist among such high polyploids. A bivalent-forming allohexaploid would be expected to be very fertile, for instance the well-known example of bread wheat which has an AABBDD genotype (Darlington 1963) and forms mostly eu-triploid ABD gametes. A completely autohexaploid plant with six identical chromosome sets (i.e. an AAAAAA genotype) would form a large number of hexavalents at meiosis which, with the consequential irregularities of chromosome separation, would result in many widely aneuploid non-viable gametes and a considerable drop in fertility. Other hexaploids would display various amounts of homology between the chromosome sets and would present a range of fertilities between the two above extremes, and *N. lobularis* appears to be one such intermediate type. Again, a meiotic investigation, if sufficient material was available, would shed some light on the level of chromosome homology existing in this plant.

The pollen fertilities of the majority of the F₁ hybrids studied here are very low or zero (Table 1), reflecting the extreme dissimilarity of the chromosome sets donated by the two distantly-related parents of each hybrid as noted above. Only *N. atlanticus* × *N. cuatrecasii* has a reasonably high pollen fertility, indicating the closer chromosomal relationship of this particular pair of species.

THE CULTIVARS

Chromosome counts were made of 230 *Narcissus* cultivars. These are plants of unknown or complex parentage (i.e. not simply F₁ hybrids between pairs of species) and were selected at random from the lists offered by the suppliers mentioned above in Material and Methods. They may thus be regarded as a representative random sample of the many thousands of cultivars extant in the genus, which is relevant to the discussions of the occurrence of aneuploidy and polyploidy below. The counts are given in Table 2, together with the RHS division/colour/morphology code of each cultivar and the pollen fertility of nearly every one.

Because these cultivars are all of complex origin they are to be expected to display a considerable degree of heteromorphy among homeologous pairs or

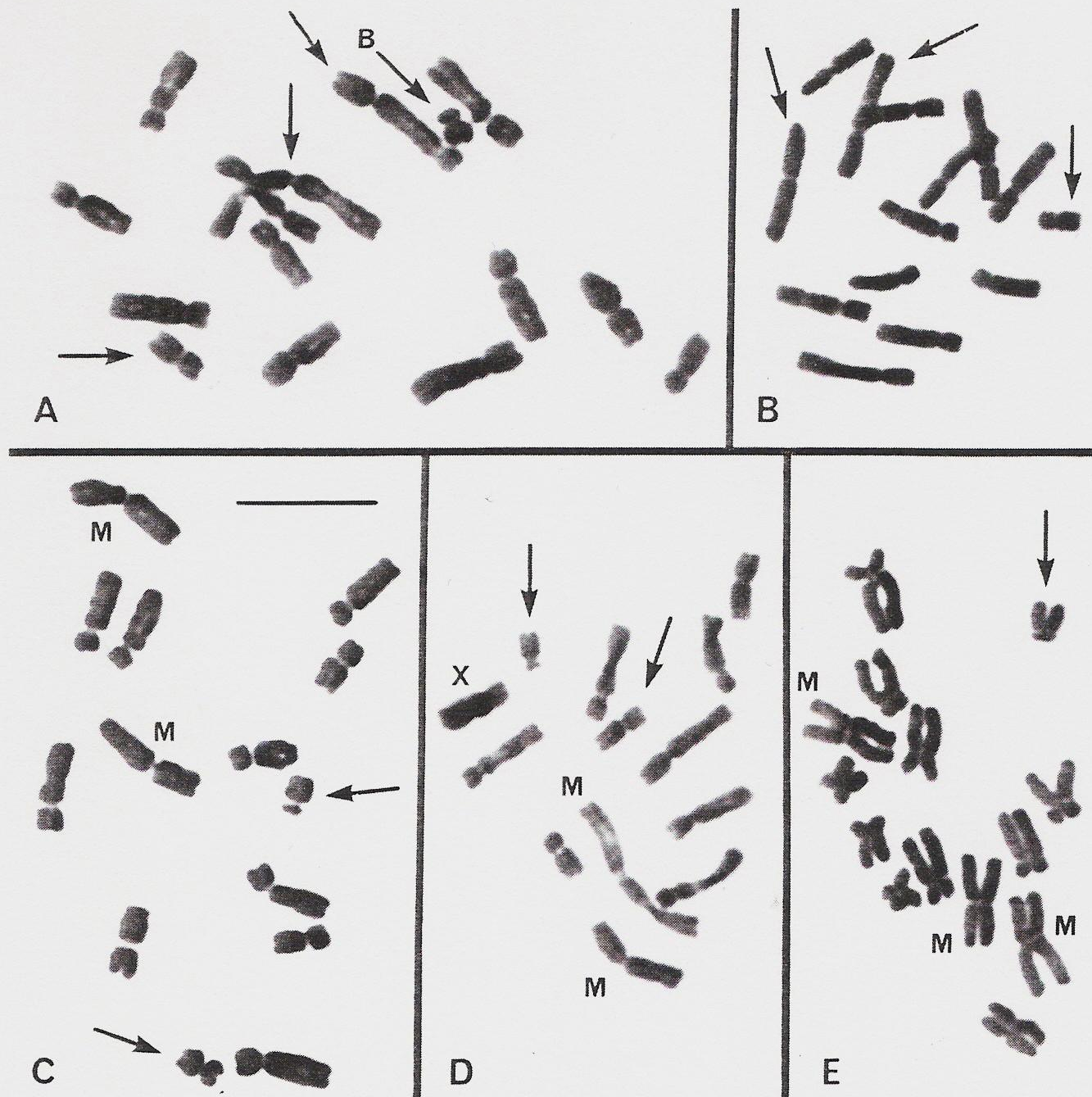


FIG. 5. Somatic chromosomes of sterile F_1 hybrid *Narcissus* diploids ($2n = 14$) showing heteromorphy of genomes (morphologically unique and unpaired chromosomes arrowed.) **A** *N. atlanticus* \times *N. asturiensis* 'Navarre' with $2n = 14 + 1B$; **B** *N. x odorus* var. *rugulosus*; **C, D, E** *N. rupicola* \times *N. gaditanus* showing variability in three sister hybrids, **C** with 2 similar metacentrics (M), **D** with 2 dissimilar metacentrics (M), **E** with 3 similar metacentrics (M). In **D** two chromosomes are completely overlapping at X. Scale = $10\mu\text{m}$.

groups of chromosomes, so much so that it is difficult to construct karyotypes showing chromosome homologies in these plants, since genetically similar homeologues might be very dissimilar morphologically. This part of the Results and Discussion section will therefore be devoted to the following matters only: the incidence of B chromosomes; the frequency of the various chromosome numbers; a summary, in the light of present knowledge, of the known parentage and the possible chromosomal derivation of a range of cultivars bearing the more uncommon chromosome numbers; pollen fertility; a comparison of ploidy levels in cultivation of *Narcissus* with those of some other popularly-cultivated decorative genera.

B chromosomes

These are rather difficult to identify with accuracy in these cultivars because they are mostly euchromatic and mitotically quite stable, not varying from root to root nor from bulb to bulb of a particular cultivar (all of which are

TABLE 2. Chromosome numbers, R.H.S. division and pollen fertility in 230 *Narcissus* cultivars of mixed origin.

Cultivar	Division/ colour	Kew accession number	Origin	2n	Pollen Fertility %
'Abel Tasman'	11 W-WYO	399-82.04378	Rosewarne	28	15
'Acropolis'	4 W-R	291-82.02813	Wisley	28	0
'Acropolis'	4 W-R	399-82.04379	Rosewarne	28	0
'Actaea'	9 W-GYR	291-82.02814	Wisley	28	94
'Alabaster'	4 W-W	399-82.04380	Rosewarne	14	
'Ann Abbott'	2 W-YPP	399-82.04381	Rosewarne	28	44
'Ardour'	3 Y-R	399-82.04382	Rosewarne	28	78
'Armada'	2 Y-O	291-82.02816	Wisley	28	84
'Aspasia'	8 W-Y	364-84.03755	Willis	24	
'Barrii Conspicuus'	3 Y-YYR	257-83.03154	Rosewarne	21	
'Bartley'	6 Y-Y	291-82.02818	Wisley	21	34
'Bath's Flame'	3 Y-YYR	257-83.03155	Rosewarne	21	
'Bawnboy'	1 Y-Y	291-82.02817	Wisley	28	
'Bealita'	3 Y-O	399-82.04384	Rosewarne	28	62
'Beersheba'	1 W-W	257-83.03156	Rosewarne	28	
'Benediction'	3 W-GWW	399-82.04385	Rosewarne	28	25
'Bernardino'	2 W-YYO	257-83.03157	Rosewarne	28	
'Big Wig'	4 W-Y	399-82.04386	Rosewarne	28	0
'Binkie'	2 Y-W	291-82.02819	Wisley	28	80
'Blaris'	2 W-P	291-82.02820	Wisley	28	83
'Brunswick'	2 W-Y	291-82.02821	Wisley	28	86
'Bryher'	3 W-GWW	399-82.04387	Rosewarne	28	56
'Camellia'	4 Y-Y	291-82.02822	Wisley	21	0
'Canasta'	11 W-Y	399-82.04388	Rosewarne	27	58
'Candida'	4 W-Y	399-82.04389	Rosewarne	27	0
'Cantatrice'	1 W-W	291-82.02823	Wisley	28 + B	50
'Cape Kennedy'	11 W-O	399-82.04390	Rosewarne	27	79
'Capparoe'	3 W-R	399-82.04391	Rosewarne	27	74
'Caramel'	9 W-O	399-82.04392	Rosewarne	28	94
'Caravelle'	1 Y-Y	291-82.02824	Wisley	27	93
'Carbineer'	2 Y-O	291-82.02825	Wisley	28	81
'Cassata'	11 W-W	399-82.04393	Rosewarne	28	83
'Ceylon'	2 Y-O	291-82.02826	Wisley	27	77
'Charity May'	6 Y-Y	291-82.02827	Wisley	21	0
'Charter'	2 Y-W	291-82.02828	Wisley	28	55
'Chinese White'	3 W-GWW	257-83.03158	Rosewarne	28	
'Cicely'	2 W-W	291-82.02829	Wisley	28	86
'C. J. Backhouse'	2 Y-YOO	257-83.03159	Rosewarne	14	
'Codlins and Cream'	4 W-Y	257-83.03160	Rosewarne	14	
'Colorange'	11 Y-O	399-82.04394	Rosewarne	28	47
'Content'	1 W-WWY	257-83.03162	Rosewarne	24	
'Cornet'	6 Y-Y	399-82.04396	Rosewarne	21	15
'Court Martial'	2 Y-R	291-82.02832	Wisley	28	73
'Crepello'	3 W-GWY	291-82.02830	Wisley	28	86
'Crescendo'	2 Y-YYO	399-82.04395	Rosewarne	28	81
'Crocus'	2 Y-Y	291-82.02831	Wisley	28	72
'Croesus'	2 Y-YOR	257-83.03163	Rosewarne	21	
'Cyclone'	6 Y-Y	399-82.04397	Rosewarne	28	63
'Dactyl'	9 W-YYR	399-82.04458	Rosewarne	14	88
'Dalboyne'	2 Y-O	399-82.04403	Rosewarne	28	58
'Dallas'	3 W-GWW	399-82.04404	Rosewarne	14	79
'Double Eagle'	4	399-82.04405	Rosewarne	27	0
'Double Event'	4 W-Y	399-82.04406	Rosewarne	29	0
'Doubledale'	4 W-Y	399-82.04407	Rosewarne	28	0
'Doublet'	4 W-W	399-82.04408	Rosewarne	14	0
'Dove Wings'	6 W-Y	291-82.02833	Wisley	21	3
'Downhill'	3 W-W	291-82.02834	Wisley	28	
'Dunkeld'	2 Y-O	257-83.03164	Rosewarne	28	
'Dunlewey'	2 W-W	291-82.02835	Wisley	28	30
'Early Splendour'	8 W-O	399-82.04479	Rosewarne	17	0
'Easter Moon'	2 W-GWW	291-82.02836	Wisley	28	25
'Eddy Canzony'	2 W-YYO	291-82.02837	Wisley	28	65

Cultivar	Division/ colour	Kew accession number	Origin	2n	Pollen Fertility %
'Edward Buxton'	3 Y-OOR	399-82.04409	Rosewarne	28	85
'Elvira'	8 W-Y	257-83.03165	Rosewarne	24	
'Elvira'	8 W-Y	364-84.03756	<i>Willis</i>	24	
'Emperor'	1 Y-Y	257-83.03166	Rosewarne	21	
'Empress'	1 W-Y	257-83.03167	Rosewarne	21 + B	
'Enterprise'	4 Y-O	399-82.04410	Rosewarne	28	0
'Entrancement'	1 Y-W	399-82.04411	Rosewarne	28	82
'Evening'	2 W-W	257-83.03168	Rosewarne	28	
'Extol'	4 Y-R	399-82.04412	Rosewarne	28	60
'Falaise'	4 W-O	257-83.03169	Rosewarne	26	
'February Gold'	6 Y-Y	291-82.02838	Wisley	21	39
'Fermoy'	2 W-YOO	399-82.04413	Rosewarne	28	6
'Firebrand'	3 W-R	257-83.03170	Rosewarne	14	
'Flamenco'	2 W-O	291-82.02839	Wisley	28	13
'Flower Drift'	4 W-O	291-82.02840	Wisley	28	87
'Fortwilliam'	1 Y-Y	399-82.04415	Rosewarne	28	80
'Furbelow'	4 Y-O	399-82.04416	Rosewarne	27	0
'Galway'	2 Y-Y	291-82.02841	Wisley	28	
'Garron'	1 Y-Y	399-82.04417	Rosewarne	28	84
'Gaucho'	2 Y-R	399-82.04418	Rosewarne	28 + B	78
'Gay Record'	4 W-O	399-82.04419	Rosewarne	28	0
'Gay Time'	4 W-R	399-82.04420	Rosewarne	28	94
'Gem of Ulster'	2 W-P	399-82.04421	Rosewarne	28	63
'Geranium'	8 W-O	399-82.04480	Rosewarne	17	3
'Geranium'	8 W-O	364-84.03757	<i>Willis</i>	17	
'Glorious'	8 W-R	257-83.03171	Rosewarne	24	
'Glorious'	8 W-R	364-84.03758	<i>Willis</i>	24	
'Glory of Lisse'	9 W-YYR	399-82.04460	Rosewarne	14	90
'Glowing Red'	4 W-R	399-82.04432	Rosewarne	28	0
'Golden Dawn'	8 Y-O	399-82.04481	Rosewarne	24	0
'Golden Dawn'	8 Y-O	364-84.03759	<i>Willis</i>	24	
'Golden Ducat'	4 Y-Y	291-82.02843	Wisley	28	0
'Golden Harvest'	1 Y-Y	291-82.02844	Wisley	28	96
'Golden Incense'	7 Y-Y	399-82.04430	Rosewarne	21	0
'Golden Lacquer'	6 Y-Y	399-82.04398	Rosewarne	28	2
'Golden Sceptre'	7 Y-Y	291-82.02845	Wisley	21	
'Golden Spur'	1 Y-Y	257-83.03172	Rosewarne	14	
'Golden Torch'	2 Y-Y	291-82.02846	Wisley	27	90
'Great Leap'	4 W-Y	399-82.04425	Rosewarne	21	0
'Gulliver'	3 Y-YYR	257-83.03173	Rosewarne	28	
'Hannibal'	11 Y-O	399-82.04426	Rosewarne	28	72
'Hawaii'	4 Y-R	399-82.04427	Rosewarne	27	0
'Helios'	2 Y-YOO	257-83.03174	Rosewarne	21	
'Henry Irving'	1 Y-Y	257-83.03175	Rosewarne	14	
'Hiawassee'	8 W-W	399-82.04482	Rosewarne	18	0
'Hiawassee'	8 W-W	364-84.03760	<i>Willis</i>	18	
'Highfield Beauty'	8 Y-GYO	399-82.04483	Rosewarne	31	0
'Home Fires'	2 Y-R	291-82.02847	Wisley	28	98
'Horace'	9 W-YOR	399-82.04461	Rosewarne	14	
'Horn of Plenty'	5 W-W	291-82.02848	Wisley	21	
'Hospodar'	2 Y-O	257-83.03176	Rosewarne	29	
'Ideal'	8 W-O	364-84.03761	<i>Willis</i>	17 + B	
'Inishkeen'	1 W-Y	291-82.02849	Wisley	28	89
'Irene Copeland'	4 W-Y	399-82.04428	Rosewarne	14	0
'Irish Luck'	1 Y-Y	399-82.04429	Rosewarne	28	90
'Irish Minstrel'	2 W-Y	291-82.02850	Wisley	28	88
'Jana'	6 Y-Y	399-82.04399	Rosewarne	21	39
'Jenny'	6 W-W	291-82.02851	Wisley	21	10
'Joseph Mcleod'	1 Y-Y	291-82.02852	Wisley	28	80
'Junco'	1 W-W	399-82.04437	Rosewarne	28	56
'Killara'	8 W-Y	399-82.04484	Rosewarne	45 + B	0
'Killigrew'	2 Y-R	257-83.03177	Rosewarne	28	
'Kilmorack'	2 Y-O	291-82.02853	Wisley	28	73
'Kilmore'	3 W-O	399-82.04438	Rosewarne	28	48
'Kilrea'	2 W-W	399-82.04439	Rosewarne	28 + B	3

TABLE 2 *continued*. Chromosome numbers, R.H.S. division and pollen fertility in 230 *Narcissus* cultivars of mixed origin.

Cultivar	Division/ colour	Kew accession number	Origin	2n	Pollen Fertility %
'King Alfred'	1 Y-Y	257-83.03178	Rosewarne	28	
'King Cardinal'	2 W-R	399-82.04440	Rosewarne	28	83
'Kingcraft'	8 W-GOO	364-84.03762	Willis	24	
'Kingscourt'	1 Y-Y	291-82.02854	Wisley	28	57
'Knowehead'	2 W-W	291-82.02855	Wisley	28	88
'Lady Margaret Boscawen'	2 W-Y	257-83.03179	Rosewarne	29	
'La Riante'	3 W-YOO	257-83.03180	Rosewarne	28	
'Larkelly'	6 Y-O	399-82.04401	Rosewarne	36	88
'Larkwhistle'	6 Y-Y	291-82.02856	Wisley	21	1
'Lawali'	4 W-P	399-82.04441	Rosewarne	28	80
'Lemon Doric'	2 Y-W	399-82.04442	Rosewarne	27	79
'Liberty Bells'	5 Y-Y	291-82.02857	Wisley	21	5
'Limelight'	1 Y-Y	291-82.02858	Wisley	28	74
'Lizard Light'	3 Y-O	399-82.04443	Rosewarne	27	60
'Love Song'	2 W-O	399-82.04444	Rosewarne	28	60
'Lucifer'	2 W-GRR	257-83.03181	Rosewarne	14	
'Lunar Spell'	2 Y-W	399-82.04445	Rosewarne	28	65
'March Sunshine'	6 Y-Y	291-82.02859	Wisley	21	28
'Matador'	8 Y-R	399-82.04485	Rosewarne	32 + B	66
'Maximus Superbus'	1 Y-Y	257-83.03183	Rosewarne	21	
'Merlin'	3 W-YYR	291-82.02860	Wisley	28	55
'Milan'	9 W-GYR	399-82.04462	Rosewarne	14	96
'Mistral'	11 W-Y	399-82.04446	Rosewarne	28	72
'Mitylene'	2 W-Y	257-83.03184	Rosewarne	28	
'Monterrico'	4 W-WOO	399-82.04447	Rosewarne	28	0
'Mount Hood'	1 W-W	291-82.02861	Wisley	29	77
'Mr Jinks'	3 W-O	399-82.04449	Rosewarne	29	2
'Mrs Langtry'	2 W-W	257-83.03185	Rosewarne	14	
'Mystic'	3 W-GWO	257-83.03186	Rosewarne	14	
'Nampa'	1 Y-W	399-82.04450	Rosewarne	28	73
'Nelly'	3 W-GYY	257-83.03182	Rosewarne	28	
'Nemo'	3 W-YYR	291-82.02862	Wisley	28	
'Ninth Lancer'	2 Y-O	291-82.02863	Wisley	28	78
'Niveth'	5 W-W	399-82.04490	Rosewarne	21	3
'Olivet'	2 W-W	399-82.04451	Rosewarne	28	64
'Orangery'	11 W-POY	399-82.04452	Rosewarne	28	41
'Ormeau'	2 Y-Y	291-82.02864	Wisley	28	
'Papillon Blanc'	11 W-W	399-82.04453	Rosewarne	29	14
'Papua'	4 Y-Y	399-82.04454	Rosewarne	27	0
'Passionale'	2 W-P	291-82.02865	Wisley	28	60
'Patagonia'	2 Y-R	291-82.02866	Wisley	28	52
'Penpol'	7 Y-Y	399-82.04431	Rosewarne	21	2
'Penvose'	2 W-Y	291-82.02867	Wisley	28	36
'Pepper'	2 Y-R	291-82.02868	Wisley	14	53
'Pick Up'	11 Y-O	399-82.04455	Rosewarne	28	46
'Pink Chiffon'	4 W-WPP	399-82.04456	Rosewarne	27	6
'Pinza'	2 Y-YOR	291-82.02869	Wisley	28	91
'Planet'	4 Y-Y	399-82.04457	Rosewarne	28	78
'Polar Ice'	3 W-W	291-82.02870	Wisley	29	13
'Pontresina'	2 W-Y	291-82.02872	Wisley	28	80
'Portchapel'	7 Y-Y	399-82.04432	Rosewarne	21	21
'Primrose Beauty'	4 Y-YOO	399-82.04465	Rosewarne	24	0
'Princeps'	1 W-Y	257-83.03188	Rosewarne	14	
'Quick Step'	7 W-P	399-82.04433	Rosewarne	28	66
'Rathkenny'	1 W-Y	399-82.04466	Rosewarne	28 + B	77
'Red Devon'	2 Y-O	291-82.02873	Wisley	28	82
'Red Goblet'	2 Y-R	291-82.02874	Wisley	28	87
'Rembrandt'	1 Y-Y	291-82.02875	Wisley	28	63
'Rijnveld's Early Sensation'	1 Y-Y	399-82.04469	Rosewarne	33	88
'Riotous'	4 Y-Y	399-82.04467	Rosewarne	28	24
'Rippling Waters'	5 W-W	291-82.02876	Wisley	21	19
'Roberta Watrous'	7 Y-GYP	399-82.04468	Rosewarne	28	63

Cultivar	Division/ colour	Kew accession number	Origin	2n	Pollen Fertility %
'Romeo'	8 Y-O	399-82.04486	Rosewarne	17	0
'Rose of Tralee'	2 W-P	257-83.03190	Rosewarne	27	
'Rushlight'	2 Y-W	291-82.02877	Wisley	28	76
'Rustom Pasha'	2 Y-R	257-83.03191	Rosewarne	28	
'Salmon Trout'	2 W-P	291-82.02878	Wisley	27	54
'Salome'	2 W-PPY	291-82.02879	Wisley	28	
'Sateen'	2 W-YPP	291-82.02880	Wisley	28	40
'Scarlet Gem'	8 Y-R	364-84.03763	Willis	17	
'Sebastopol'	2 W-Y	399-82.04470	Rosewarne	29	45
'Shah'	7 Y-Y	399-82.04434	Rosewarne	28	54
'Sidney Torch'	2 Y-YOO	399-82.04471	Rosewarne	28	89
'Silent Morn'	3 W-YYO	399-82.04472	Rosewarne	28	30
'Silver Chimes'	8 W-W	291-82.02881	Wisley	28 + B	
'Silver Chimes'	8 W-W	364-84.03764	Willis	28 + B	
'Sir Watkin'	2 Y-Y	257-83.03192	Rosewarne	21	
'Sir Winston Churchill'	4 W-O	399-82.04473	Rosewarne	17	0
'Snowhill'	2 W-GWW	399-82.04474	Rosewarne	28	53
'Sonata'	9 W-GYR	399-82.04464	Rosewarne	14	87
'Sparkling Eye'	8 W-GOO	364-84.03765	Willis	17	
'Spellbinder'	1 Y-W	291-82.02882	Wisley	29	66
'Spitzbergen'	1 W-Y	399-82.04476	Rosewarne	28	27
'St Agnes'	8 W-Y	364-84.03766	Willis	17	
'St Keyne'	8 Y-O	399-82.04488	Rosewarne	24	3
'St Keyne'	8 Y-O	364-84.03767	Willis	24	
'Straight'	1 W-Y	291-82.02883	Wisley	28	31
'Sunburst'	4 Y-Y	399-82.04477	Rosewarne	28	21
'Sunrise'	3 W-Y	257-83.03193	Rosewarne	14	
'Sweetness'	7 Y-Y	291-82.02884	Wisley	21	0
'Takoradi'	4 W-W	399-82.04478	Rosewarne	28	0
'Tête-à-Tête'	6 Y-O	296-82.03048	Rosewarne	24 + B	2
'Tibet'	2 W-W	291-82.02885	Wisley	28	51
'Tittle Tattle'	7 Y-Y	399-82.04436	Rosewarne	21	8
'Toreador'	3 W-R	291-82.02886	Wisley	28	50
'Trenithon'	1 Y-Y	399-82.04489	Rosewarne	28	78
'Tresamble'	5 W-W	291-82.02887	Wisley	21	16
'Verger'	3 W-R	291-82.02889	Wisley	28	95
'Verona'	3 W-W	291-82.02890	Wisley	28	23
'Victoria'	1 W-Y	257-83.03195	Rosewarne	21 + B	
'White Emperor'	1 W-W	257-83.03197	Rosewarne	28	
'White Lady'	3 W-Y	257-83.03198	Rosewarne	21	
'White Lion'	4 W-WYY	291-82.02892	Wisley	28	30
'White Marvel'	4 W-W	291-82.02893	Wisley	21	0
'White Nile'	2 W-W	257-83.03199	Rosewarne	29	
'White Sail'	4 W-W	399-82.04494	Rosewarne	14	0
'White Spire'	2 W-W	399-82.04495	Rosewarne	28	86
'Will Scarlett'	2 W-R	257-83.03196	Rosewarne	14	
'Windblown'	4 W-WYY	399-82.04496	Rosewarne	29	0
'Yankee Clipper'	2 Y-Y	291-82.02894	Wisley	28	83
'Yellow Cheerfulness'	4 Y-Y	399-82.04497	Rosewarne	24	0

hybrid and necessarily propagated clonally). It is thus not easy to decide whether a B chromosome is present or not in a plant with, for example, $2n = 3x + 1 = 22$ chromosomes, even if the extra chromosome is small and morphologically different from the remainder. It could simply be a normal member of one of the parents' complements and the extra chromosome is actually one of the other larger A chromosomes present in excess, as is common in polyploids. The cases cited below are nevertheless plants in which we are reasonably confident that B chromosomes exist.

No more than a single B chromosome was found in any cultivar, with occurrences in 'Ideal' ($2n = 17 + B$), 'Empress' and 'Victoria' ($2n = 21 + B$),

'Tête-à-Tête' ($2n = 24 + B$), 'Cantatrice', 'Gaucho', 'Kilrea', 'Rathkenny' and 'Silver Chimes' ($2n = 28 + B$), 'Matador' ($2n = 32 + B$) and 'Killara' ($2n = 45 + B$). These B chromosomes are all smaller than the smallest A chromosome and vary in morphology. In 'Silver Chimes' and 'Killara' (Fig. 10B, 10C) the B is a minute metacentric (the B chromosome in 'Silver Chimes' was clearly illustrated in her Fig. 5A by Wylie, 1952, but was not recognised, since she reported $2n = 29$ chromosomes in this cultivar rather than $28 + B$). The B's in these two cultivars are apparently identical (see below) and heterochromatic, staining less intensely than the A chromosomes. In all other cases the B is euchromatic. In 'Ideal' and 'Matador' (Fig. 9B) it is slightly larger than that in 'Killara' (Fig. 10C), but it is still small and metacentric. In 'Tête-à-Tête' it is a submetacentric of similar size (Fig. 7C) and in the remaining cultivars they are larger acrocentrics, almost as large as the smallest A chromosome.

Frequency of the various chromosome numbers

The chromosome numbers in our sample of 230 cultivars (ignoring the presence of B chromosomes in eleven of them) are summarised in Table 3. Also in Table 3 our results are combined with other published accounts of the chromosomes of *Narcissus* cultivars (de Mol 1923, Nagao 1929, Philp 1934, Fernandes 1950, Wylie 1952, Fernandes & de Almeida 1971, A.D.S. 1981) to give a total of 466 records which rises to 526 when those for the wild species and natural F_1 hybrids in division 10 are added. All records are classified in Table 3 according to the RHS division of the plants. In the first part of the following account all members of division 8 and some members of divisions 4, 6, 10 and 11 (identified by asterisks in Table 3) will be omitted. These plants are subgenus *Hermione*, *N. tazetta* and its allies ($x = 10, 11$) and cultivars having plants of this group in their ancestry. The combination, in various multiples and proportions, of $x = 10$ or 11 chromosomes of subgenus *Hermione* with $x = 7$ from subgenus *Narcissus* results in a series of complex aneuploids and apparent aneuploids which will be discussed in detail later.

The remainder of Table 3 is concerned with cultivated forms of species (division 10) or more complicated hybrid cultivars which are nevertheless derived solely from ancestors having $x = 7$ chromosomes. It shows that the species forms and natural hybrids of division 10 are largely diploid, in common with the majority of the species from which they have arisen (Fedorov 1969, Moore 1982), and only a relatively few of them are triploid ($2n = 21$) or tetraploid ($2n = 28$). To a lesser degree this pattern is repeated in division 9, the Poeticus group, which also has a majority of diploid records, with sixteen diploids against only six tetraploids and no triploids. In divisions 7 (Jonquilla hybrids), 6 (Cyclamineus hybrids) and 5 (Triandrus hybrids) there are indications of an increase in the number of polyploids compared with that of diploids, with the triploid level being predominant (Fig. 6B) and the tetraploid level being roughly similar to that of the diploids with the exception of division 5 in which no tetraploid has been located so far. A single aneu-pentaploid has been recorded in division 6; 'Larkelly' ($2n = 5x + 1 = 36$), and its derivation will be discussed below.

Divisions 1 to 3 are principally derived from *N. pseudonarcissus* with some interbreeding with other species to reduce the length of the corona from very long in division 1 to quite short in division 3. Division 11 is of similar origin,

TABLE 3. Total of known chromosome numbers in *Narcissus* cultivars arranged in RHS divisions (466 records plus 60 in division 10). Present authors' findings on 230 cultivars in brackets. B chromosomes are omitted, also the authors' records of division 10 plants which are in Table 1.

R.H.S. Division	2n =	No. of records of each chromosome number																		
		14	17	18	19	20	21	22	24	26	27	28	29	30	31	32	33	36	45	50
1		(3) 6					(4) 5	2	(1) 1	1	(1) 3	(21) 58	(2) 4		1		(1) 1			
2		(5) 10			1		(3) 6			2	(6) 9	(54) 131	(4) 4	1						
3		(4) 9					(3) 7				(2) 3	(17) 30	(2) 2							
4		(5) 6	(1)* 1*				(3) 8		(2*) 3*	(1) 1	(6) 6	(17) 18	(2) 4							
5		2					(5) 7													
6		4					(9) 10		(1*) 1*		(2) 4							(1) 1		
7		2					(6) 12				(3) 3			1						
8*			(7*) 14*	(1*) 1*		1*			(6*) 6*		(1*) 1*		4*	(1*) 1*	(1*) 2*				(1*) 1*	
9		(5) 16									(2) 6									
10		44	3*				5				5		1*	1*						1*
11							1				(1) 1	(6) 12	(1) 1	1*						
12											6									

*Plants with subgenus *Hermione* partially or totally in ancestry.

TABLE 4. Frequency of chromosome numbers in *Narcissus* cultivars based solely on subgenus *Narcissus* ($x = 7$). The authors' records are taken separately and combined with earlier records. B chromosomes are omitted, also all records in division 10 (species and natural hybrids), division 8 and any other plant known to have *N. tazetta* or its allies in its ancestry (marked with asterisks in Table 3).

The authors' records			Authors' plus earlier records		
2n =	No. of records	%	2n =	No. of records	%
14	22	10.58	14	55	12.82
19	0	0	19	1	0.23
21	33	15.87	21	56	13.05
22	0	0	22	2	0.47
24	1	0.48	24	1	0.23
26	1	0.48	26	4	0.93
27	16	7.69	27	22	5.13
28	122	58.65	28	268	62.47
29	11	5.29	29	15	3.49
30	0	0	30	1	0.23
31	0	0	31	2	0.47
33	1	0.48	33	1	0.23
36	1	0.48	36	1	0.23
Total	208		Total	429	
26-31	150	72.11	26-31	312	72.73

except that the corona is partially or completely divided into segments. In these divisions (except division 11) there are some diploids (Fig. 6A) and triploids, but these are completely overshadowed by large numbers of cultivars at or near the tetraploid level (Figs. 6C, 6D, 6E), with $2n = 4x \pm 2 = 26 - 30$. There are also two records in division 1 of higher numbers, $2n = 31$ and 33 (Fig. 7A). Division 12 (Miscellaneous daffodils) contains only tetraploids, but numbers sampled are too low to be significant. Division 4 (Double-flowered cultivars) has a distribution of diploids, triploids, tetraploids and aneuploids similar to those of divisions 1-3.

The chromosome numbers of our plants taken alone and added to those of earlier workers' material are summarised in Table 4. In this table only hybrid cultivars based solely on $x = 7$ chromosomes are taken into account (i.e. division 10 is excluded, and also all plants with subgenus *Hermione* in their ancestry, comprising division 8 and a few other cultivars asterisked in Table 3).

The very low overall frequency of diploids (10.5-13%) and triploids (13-16%) is quite evident, also the very high frequency of tetraploids and near-tetraploids. The records at the tetraploid level are very similar, our findings being 58.65% and the total records being 62.47%, but these figures approach each other even more closely when records in the aneuploid/euploid/aneuploid range from $2n = 26$ to 31 are combined (Table 4), ours becoming 72.11% and the total records becoming 72.73%. Thus we found relatively slightly fewer eu-tetraploids ($2n = 28$) and relatively more aneu-tetraploids ($2n = 26, 27, 29$) than the previous workers did, indicating possibly that some aneuploids at this level were scored by early workers as eu-tetraploids, perhaps because they were working with poorly spread mitotic cells or with sections.

Notwithstanding the slight discrepancy over aneuploid frequency at the tetraploid level our data on the relative frequency of the various chromosome numbers in *Narcissus* cultivars match those of previous workers very closely. It therefore follows that our data are representative of the total data, and the latter, an overall sample of 429 cultivars based solely on $x = 7$ plants, may be taken confidently as being characteristic of all of the ten thousand cultivars, i.e. if every one of them was analysed chromosomally the relative frequencies of the various chromosome numbers should not differ markedly from those presented here.

Aneuploids have not been detected at the diploid level, showing either that meiosis is quite regular in diploids, producing only $n = 7$ gametes, or that any aneuploids produced are non-viable, or at best are too weak to be selected by breeders. At the tetraploid level they are more common, with many records (Tables 3 & 4) of plants having $4x \pm 1 = 27$ or 29 chromosomes (Figs. 6D, 6E), and a few having $4x \pm 2 = 26$ or 30 chromosomes. It is well-known that in autotetraploids having a high quadrivalent frequency at meiosis the production of aneuploid gametes is quite common. The aneuploid products of the fusion of these gametes with each other or with normal ones are viable, due to the presence of several sets of chromosomes which buffer the adverse effects of the chromosomal imbalance, and have been reported to survive even under the harshly competitive conditions of the wild environment (Brandham & Johnson 1977). These are clearly more likely to survive in the less competitive environment found in cultivation, and the aneuploidy could easily endow the individual with a morphological characteristic that appeals to the hybridist, provided that the aneuploids do not show too much loss of vigour.

At the triploid level aneuploidy is rare and the two records of $2n = 3x + 1 = 22$, 'Grandis' and 'Horsefieldii' were even suggested by Wylie (1952) actually to be $2n = 21 + B$ (i.e. similar to 'Empress' and 'Victoria' which are recorded above as triploids with one B chromosome). Because of its demonstrable irregularity, tetraploid meiosis produces many aneuploid gametes and it might thus be expected that aneuploidy would be more common in triploids if it is assumed that they normally result from diploid/tetraploid hybridisations. The almost complete absence of aneuploids at the triploid level therefore indicates that they are probably only rarely the products of diploid \times tetraploid crosses and that the majority arise from diploids alone in another way. This is the process of non-reduction, in which a few gamete-mother cells of a diploid bypass normal meiosis and produce diploid gametes directly, which fuse with normal haploid gametes to produce triploid offspring (Brandham 1982). This process does not permit the formation of aneu-triploids, since the non-reduced gametes from a diploid can be only precisely diploid and not aneuploid. Non-reduction can occur at any level of polyploidy and also in wide hybrids. In fact in the latter the non-reduced gametes are often the only viable ones that the plant can produce (Wylie 1952). Non-reduction will be shown below to be an important and frequent source of many of the *Narcissus* cultivars at the triploid and higher levels of polyploidy.

Above the tetraploid level there are a few aneuploid cultivars which approach or slightly exceed the pentaploid level of $2n = 35$ (to be considered individually below), but no higher polyploids have been found. Wylie (1952) found an aneu-hexaploid ($2n = 6x - 1 = 41$) occurring spontaneously in a population of a presumably tetraploid cultivar, originating probably from the fusion of a non-reduced gamete ($n = 28$) with an aneuploid one ($n = 13$). This



FIG. 6. Somatic chromosomes of *Narcissus* cultivars. **A** Diploid 'Firebrand' (division 3), $2n = 14$; **B** Triploid 'Jenny' (division 6), $2n = 21$; **C** Tetraploid 'Nemo' (division 3), $2n = 28$; **D** Aneu-tetraploid 'Salmon Trout' (division 2), $2n = 27$; **E** Aneu-tetraploid 'Papillon Blanc' (division 11), $2n = 29$. Scale = $10\mu\text{m}$.

she reported to be viable but weak and it is very clear that the tetraploid is the optimal level for the cultivated forms of *Narcissus*. The allohexaploid *N. × dubius* ($2n = 50$) has been placed by A.D.S. (1981) in division 10 (Table 3) and

is a naturally occurring species of hybrid origin rather than a cultivar (see below).

The derivation of apparent aneuploids and other uncommon chromosome numbers

The subgenus *Hermione* of *Narcissus* differs from subgenus *Narcissus* in having the basic chromosome numbers $x = 10$ or 11 rather than $x = 7$, but despite this difference members of the two subgenera are often sexually compatible, at least as far as the F_1 hybrid stage. In these inter-subgeneric crosses breeders have mostly used the $x = 10$ forms of *N. tazetta* and its allies ($2n = 20$), giving hybrids with $2n = 17$ when crossing them with diploids having $2n = 14$ and hybrids with $2n = 24$ when crossing them with tetraploids having $2n = 28$. A wide range of true and apparent aneuploids also occurs within this particular group of hybrids (division 8), and their origin and probable chromosome constitution will now be considered individually, together with plants from other divisions which either have *N. tazetta* affinities, or are aneuploid or have uncommon or otherwise interesting euploid chromosome numbers. Data are taken from Nagao (1929), Philp (1934), Wylie (1952), Jefferson-Brown (1969) and A.D.S. (1981), supplemented by our own observations (Table 3).

In predicting the chromosomal origin of a cultivar several different types of chromosomal behaviour at meiosis are taken into account as possible mechanisms for the origin of particular gamete types:- a, non-reduction, in which a plant of any chromosome number can produce a gamete with the same chromosome number, or occasionally double that number, as in the extreme case of the *N. tazetta* hybrid 'Geranium' ($2n = 17$) in which the only viable gametes have $n = 17$ or 34 chromosomes (Wylie 1952); b, the irregular nature of meiosis in autotriploids (in this case with $2n = 21$). Through the irregular 2:1 segregation of each of the potential maximum of seven trivalents at anaphase I the gametes of such triploids can contain any number of chromosomes between $n = 7$ and $n = 14$ with the numbers nearest to the extremes being the most viable (Brandham 1982); c, the irregular nature of meiosis in tetraploids ($2n = 28$). Although normally more regular than in triploids, meiosis in these plants can produce viable gametes with $n = 13$ or 15 (less commonly 12 or 16) in addition to the expected $n = 14$ chromosomes.

Commonly-occurring aneuploids between $4x - 2$ ($2n = 26$) and $4x + 1$ ($2n = 29$) arise mostly as a result of meiotic irregularities in tetraploids and only two aneuploids between $2n = 26$ and 29 will be considered in detail, together with the apparently eu-tetraploid 'Silver Chimes' ($2n = 28 + B$) which is actually a tribasic allotriploid of complex parentage (see below).

The analysis of the origin of selected cultivars follows, the account being arranged in R.H.S. division order and in order of increasing chromosome number within each division.

Division 1.

'Maximus Superbus' ($2n = 21$). Selected from *N. pseudonarcissus* subsp. *major* Baker (= *N. hispanicus* Gouan), a diploid with $2n = 14$. The result of the fusion of a non-reduced gamete with a normal one.

'Magnificence' ($2n = 21$). A seedling from triploid 'Maximus'. *Narcissus* is not known to be self-fertile, so this cultivar is probably the result of a triploid \times diploid cross or a triploid \times tetraploid cross. Either of these crosses can

produce some triploid progeny, with a diploid gamete being produced by the autotriploid parent in the former case and a haploid gamete in the latter case (see Brandham 1982). The crosses can be written thus: $2n = 14$ (from 21) + 7 (from 14) = 21 (the triploid \times diploid cross) or $2n = 7$ (from 21) + 14 (from 28) = 21 (the triploid \times tetraploid cross). Other crosses or probable crosses will be expressed in the same way below.

'Victoria' ($2n = 21 + B$). Unknown origin; probably arose from a diploid ($2n = 14 + B$) through the fusion of a non-reduced gamete ($n = 14 + B$) with a normal gamete ($n = 7$) from another diploid ($2n = 14$). This plant has given rise vegetatively to the split-corona (division 11) cultivar 'Buttonhole', which also has $2n = 21 + B$.

'Emperor' ($2n = 21$) and 'Empress' ($2n = 21 + B$). Both are very old triploids introduced in 1865 from a cross between *N. pseudonarcissus* var. *pseudonarcissus* and *N. pseudonarcissus* var. *bicolor* Willk. The latter was reported to be tetraploid by A. & R. Fernandes (1946), but was suggested to be triploid by Wylie (1952) because of its very low fertility. Thus these triploids arose in one of two ways: $2n = 7$ (from 14) + 14 (from 28) = 21, or $2n = 7$ (from 14) + 14 (from 21) = 21, the former according to A. & R. Fernandes and the latter according to Wylie. The B chromosome in 'Empress' could have been inherited from var. *bicolor* (see below under 'Grandis').

'Grandis' ($2n = 22$, or $21 + B$ according to Wylie (1952)). Selected from *N. pseudonarcissus* var. *bicolor*, also most probably $2n = 21 + B$, after pollination by a diploid. Thus $2n = 14 + B$ (from $21 + B$) + 7 (from 14) = $21 + B$.

'Horsefieldii' ($2n = 22$, or $21 + B$ according to Wylie (1952)). From triploid 'Maximus' \times probable triploid (+ B) *N. pseudonarcissus* var. *bicolor*. $2n = 7$ (from 21) + $14 + B$ (from $21 + B$) = $21 + B$.

'Content' ($2n = 24$). No affinity with subgenus *Hermione*, unlike several division 8 cultivars with $2n = 24$ (see below). From 'Lord Antrim' \times 'Beersheba'. The latter is known to have $2n = 28$ chromosomes, but the former is chromosomally unknown. As a cross between two tetraploids is unlikely to give progeny lacking four chromosomes ($2n = 24 = 4x - 4$) it is more probable that 'Lord Antrim' is a triploid, contributing $n = x + 3 = 10$ chromosomes to the hybrid as a result of irregular triploid meiosis. Thus in 'Content' $2n = 10$ (from 21) + 14 (from 28) = 24.

'Charles' First' ($2n = 26$). Unknown origin; possibly from tetraploid \times tetraploid or tetraploid \times triploid cross with the involvement of aneuploid gametes, i.e. $2n = 13$ (from 28) + 13 (from 28) = 26 or $2n = 13$ (from 28) + 13 (from 21) = 26. It could also have arisen from a diploid \times triploid cross through a combination of non-reduction and aneuploidy, thus $2n = 14$ (from 14) + 12 (from 21) = 26.

'Madame de Graaff' ($2n = 31$, or more probably $30 + B$). No affinity with subgenus *Hermione*, unlike other plants with approximately this number (see below). From 'Empress' ($2n = 21 + B$) \times *N. pseudonarcissus* var. *albescens*. The most probable origin assumes var. *albescens* also to be triploid. Thus with non-reduction in the female parent the plant is $2n = 21 + B$ (from $21 + B$) + 9 (from 21) = $30 + B$. Two less probable origins are as follows, assuming respectively that var. *albescens* is diploid or tetraploid; a, $2n = 16 + B$ (from $21 + B$) + 14 (from 14 by non-reduction) = $30 + B$, a very unlikely alterna-



FIG. 7. Somatic chromosomes of *Narcissus* cultivars. **A** Sub-pentaploid 'Rijnveld's Early Sensation' (division 1), $2n = 33$; **B** Allotriploid 'Yellow Cheerfulness' (division 4), $2n = 24$; **C** Allotriploid 'Tête-à-Tête' (division 6), $2n = 24 + B$ (arrow). Scale = $10\mu\text{m}$.

tive because a triploid does not normally produce a gamete meiotically with more than the diploid number of chromosomes (in this case 14); b, $2n = 14 + B$ (the largest possible gamete from $21 + B$) + 16 (from 28 through irregular meiosis) = $30 + B$.

'Rijnveld's Early Sensation' ($2n = 33$). A sub-pentaploid of unknown origin; no evidence of *Hermione* ancestry unlike other plants with $2n = 33$ (see below). As it is a $4x + 5$ plant it is unlikely to have arisen from the hybridisation of two tetraploids because the degree of aneuploidy is too high. It can therefore have arisen in one of three ways; a, triploid \times triploid with non-reduction in one parent and an aneuploid gamete from the other, i.e. $2n = 21$ (from 21) + 12 (from 21) = 33; b, triploid \times tetraploid, again with non-reduction and aneuploidy, i.e. $2n = 21$ (from 21) + 12 (from 28) = 33; c, the least likely origin is the involvement of a hexaploid ($2n = 42$), of which very few are known to exist, i.e. $2n = 14$ (from 21 or 28) + 19 (from 42) = 33 (Fig. 7A).

Division 2.

'Bethany' ($2n = 19$). From 'Binkie' ($2n = 28$) crossed with an unnamed seedling (which could only have been diploid) with the involvement of an aneuploid gamete from the former; thus $2n = 12$ (from 28) + 7 (from 14) = 19.

'Gervo' ($2n = 30$). Unknown origin, but no *Hermione* ancestry. Probably $2n = 15$ (from 28) + 15 (from 28) = 30.

Division 4.

'Sir Winston Churchill' ($2n = 17$). From *N. poeticus* × *N. tazetta* ($2n = 20$), i.e. $2n = 7$ (from 14) + 10 (from 20) = 17.

'Cheerfulness' ($2n = 24$). Double-flowered form of 'Elvira' (division 8, see below) originating vegetatively.

'Yellow Cheerfulness' ($2n = 24$). A yellow-flowered mutation derived vegetatively from 'Cheerfulness' (Fig. 7B).

'Primrose Beauty' ($2n = 24$). Unknown origin, but clearly related to subgenus *Hermione*, so presumably derived from *N. aff. tazetta* crossed with an unknown tetraploid of subgenus *Narcissus*, i.e. $2n = 10$ (from *taz.* 20) + 14 (from 28) = 24.

'Falaise' ($2n = 26$). From the double-flowered 'Mary Copeland' ($2n = 21$) × unknown (probably $2n = 28$) with the involvement of an aneuploid gamete from the former; i.e. $2n = 12$ (from 21) + 14 (from 28) = 26. Among plants of the *Aloineae* tribe (*Liliaceae*) crosses between triploids ($2n = 21$) and tetraploids ($2n = 28$) produce quite large numbers of $2n = 26$ aneuploids, among others (Brandham 1982). "Weedy little... 'Falaise'... cannot be said to be outstandingly robust" (Jefferson-Brown 1969: 115, 31), which is perhaps indicative of the fact that loss of two of the genetically more vital chromosomes from a tetraploid is quite a major degree of aneuploidy and can have extensive deleterious effects on the morphology and vigour of the plant.

Division 6.

'Tête-à-Tête' ($2n = 24 + B$). An allotriploid of complex parentage, including 'Soleil d'Or' (a diploid plant of subgenus *Hermione*) in its origin. The plants involved are: [(diploid *N. cyclamineus* × diploid 'Soleil d'Or') = 'Cyclataz'] × 'Op' (presumably also diploid). Chromosomally the origin is as follows: a. $2n = 7$ (from 14) + 10 (from 20) = 17 'Cyclataz'; b. $2n = 17$ (from 17 'Cyclataz' with non-reduction) + 7 (from 14) = 24 'Tête-à-Tête'. The B chromosome could have been inherited from any one of the three immediate ancestors (Fig. 7C).

'Larkelly' ($2n = 36$). This cultivar ($5x + 1$) is from an unnamed plant (presumably an aneu-tetraploid with $2n = 29$) crossed with *N. cyclamineus*, of which forms with more than 14 chromosomes are unknown (A. & R. Fernandes 1946). Its probable origin is through a non-reduced gamete from the aneu-tetraploid, hence $2n = 29$ (from 29) + 7 (from 14) = 36.

Division 7.

'Golden Perfection' ($2n = 31$). There are conflicting accounts of the origin of this cultivar, and without access to illustrations of its chromosomes we cannot say which is correct. Possibly there are two different cultivars with the same name in this case. Wylie (1952) proposed two types of *N. jonquilla-poeticus-tazetta* ancestry: a, tetraploid *N. poeticus* ($2n = 28$) crossed with diploid *N. tazetta* to give a $2n = 24$ hybrid which produced a non-reduced gamete to fuse with a

haploid *N. jonquilla* gamete ($n = 7$) to result in a $2n = 31$ allotetraploid with a genomic constitution Jo-Po-Po-Tz; b, diploid *N. poeticus* ($2n = 14$) crossed with diploid *N. tazetta* ($2n = 20$) to give a $2n = 17$ hybrid which produced a non-reduced ($n = 17$) gamete to fuse with a non-reduced ($n = 14$) gamete from *N. jonquilla* to result in a $2n = 31$ hybrid with the genomic constitution Jo-Jo-Po-Tz which is indistinguishable from the above.

The version of the origin of 'Golden Perfection' in A.D.S. (1981) is significantly different from Wylie's account, still with the involvement of *N. poeticus* and *N. jonquilla*, but with *N. pseudonarcissus* rather than *N. tazetta*. Details are as follows: The hybrid ($2n = 31$) is derived from the diploid ($2n = 14$) *N. × odorus* (from *N. pseudonarcissus* × *N. jonquilla*), which produced a non-reduced gamete to fuse with an aneuploid $n = 17$ gamete from the other parent 'Homespun' which has no *Hermione* ancestry. The chromosome constitution of 'Homespun' is unknown, but as the production of $n = 2x + 3 = 17$ gametes from tetraploids is very rare it is more likely that 'Homespun' is a pentaploid with $2n = 35$, from which gametes with $n = 17$ would be more common. 'Homespun' is from diploid *N. poeticus* subsp. *ornatus* × 'Golden Spur' (? $2n = 4x = 28$), a selection from *N. pseudonarcissus* subsp. *major*. Non-reduction in 'Golden Spur' would give a gamete with $n = 28$ chromosomes to fuse with a normal $n = 7$ gamete from *N. poeticus* subsp. *ornatus* to give $2n = 35$ in 'Homespun'. There is also a possibility that 'Golden Spur' is triploid ($2n = 21$), and that 'Homespun' arose from non-reduction in both of its immediate parents i.e. $2n = 14$ (from 14) + 21 (from 21) = 35.

Division 8.

'Cyclataz' ($2n = 17$). From *N. tazetta* 'Soleil d'Or' ($2n = 20$) × *N. cyclamineus* ($2n = 14$), i.e. $2n = 10$ (from 20) + 7 (from 14) = 17.

'Early Splendour', 'Romeo', 'Sparkling Eye' (all $2n = 17$), 'Ideal' ($2n = 17 + B$). All from unknown subgenus *Narcissus* diploids crossed with diploid *N. tazetta*, i.e. $2n = 7$ (from 14) + 10 (from 20) = 17 (Fig. 8A). The B chromosome in 'Ideal' could have originated from either parent.

'Alsace', 'Cragford', 'Geranium', 'Irene', 'Jaune à Merveille', 'L'Innocence', 'Laurens Koster', 'Scarlet Gem', 'St. Agnes' (all $2n = 17$). From *N. poeticus* ($2n = 14$) × *N. tazetta* ($2n = 20$), i.e. $2n = 7$ (from 14) + 10 (from 20) = 17. These hybrids are sterile except for the production of some non-reduced or doubled gametes (Wylie 1952).

'Hiawasse' ($2n = 18$). From diploid 'Cassandra' (a hybrid between *N. poeticus* subsp. *poeticus* and *N. poeticus* subsp. *radiiflorus* var. *poetarum*) crossed with 'Paper White' (= *N. papyraceus*, $2n = 22$), an ally of *N. tazetta*, i.e. $2n = 7$ (from 14) + 11 (from 22) = 18 (Fig. 8B).

'Soleil d'Or' ($2n = 20$). A diploid selection from *N. tazetta* ($2n = 20$).

'Aspasia', 'Kingcraft', 'St. Keyne' (all $2n = 24$). From unknown tetraploids of subgenus *Narcissus* ($2n = 28$) crossed with *N. tazetta* ($2n = 20$), i.e. $2n = 14$ (from 28) + 10 (from 20) = 24.

'Elvira', 'Glorious' (both $2n = 24$). From tetraploid *N. poeticus* crossed with diploid *N. tazetta*, i.e. $2n = 14$ (from 28) + 10 (from 20) = 24 (Fig. 8C). All of these *N. tazetta* hybrids with $2n = 24$ chromosomes are dibasic allotriploids

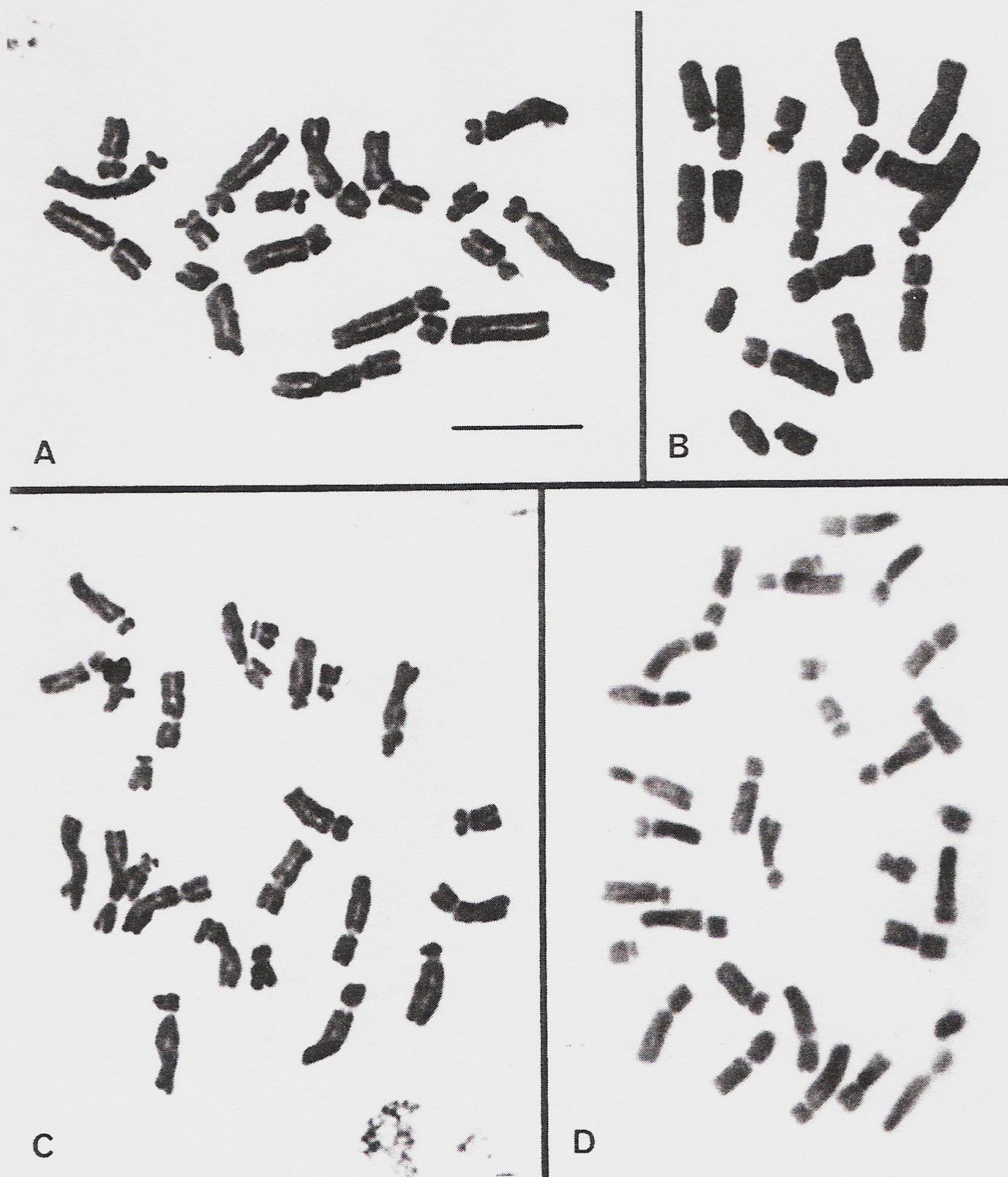


FIG. 8. Somatic chromosomes of *Narcissus* cultivars in division 8, the *N. tazetta* hybrids. **A** 'Sparkling Eye', $2n = 17$; **B** 'Hiawasee', $2n = 18$; **C** 'Elvira', $2n = 24$; **D** 'Highfield Beauty', $2n = 31$. Scale = $10\mu\text{m}$.

which would be expected to be totally sterile except for the occasional production of non-reduced gametes (see discussion of 'Killara', below).

'Chinensis', 'Grand Emperor of China', 'Grand Primo Citronière', 'Grand Soleil d'Or' (all $2n = 30$). All autotriploids selected from *N. tazetta* ($2n = 20$). 'Grand Soleil d'Or' is a triploid derivative of the diploid 'Soleil d'Or' (Wylie 1952).

'Highfield Beauty' ($2n = 31$). A cross between two plants of subgenus *Hermione* having $2n = 20$ and 22 respectively, with non-reduction in the former. i.e. $2n = 20$ (from 20) + 11 (from 22) = 31 (Fig. 8D).

'Scilly White' ($2n = 32$). Another cross between two plants of subgenus *Hermione* having $2n = 20$ and 22 but with non-reduction in the latter, i.e. $2n = 10$ (from 20) + 22 (from 22) = 32 .

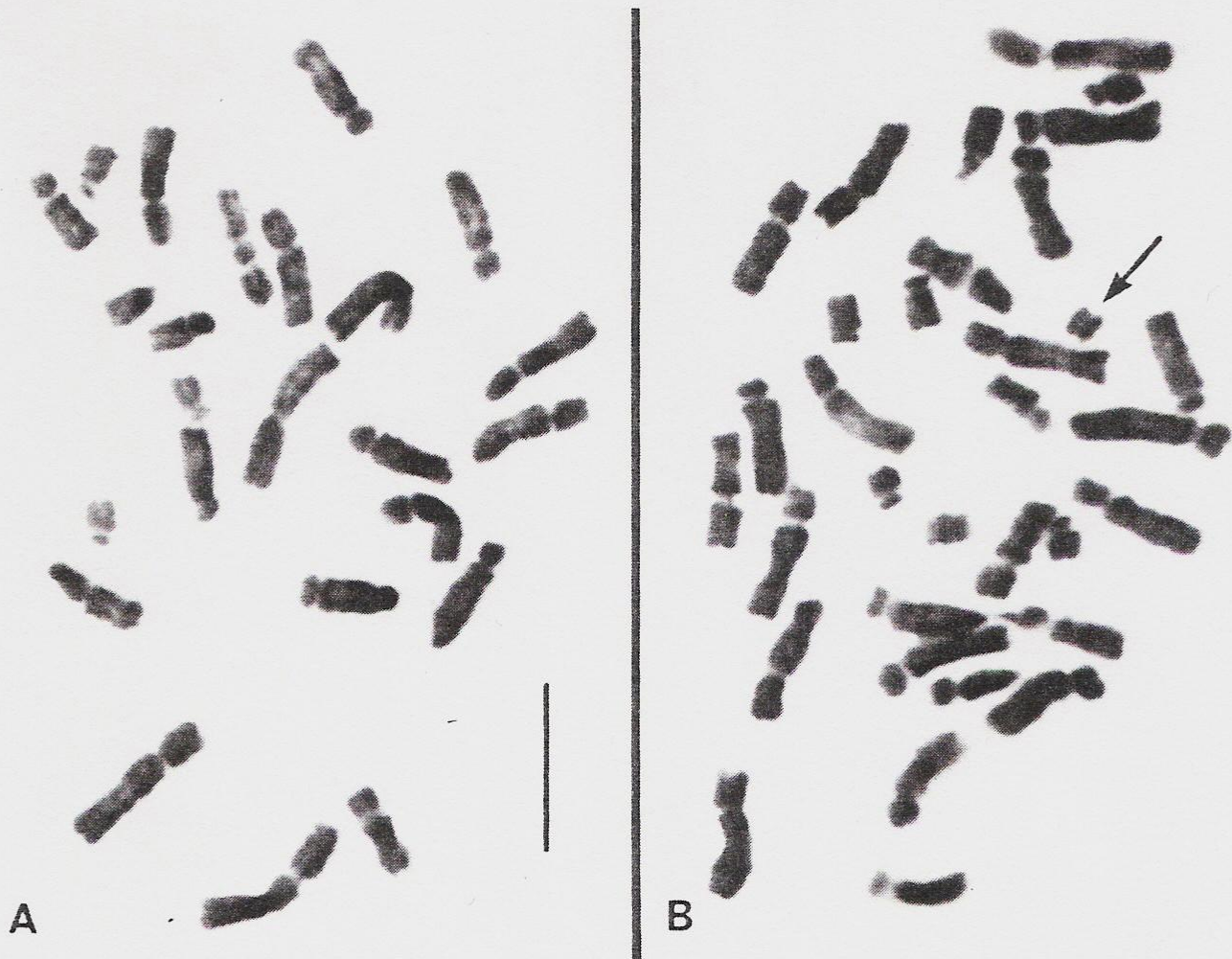


FIG. 9. Somatic chromosomes of *Narcissus* cultivars in division 8, the *N. tazetta* hybrids. **A** 'Golden Dawn', $2n = 24$; **B** 'Matador', $2n = 32 + B$ (arrow). Scale = $10\mu\text{m}$.

'Golden Dawn' ($2n = 24$) and 'Matador' ($2n = 32 + B$). Both were introduced by J. de Graaff of Oregon Bulb Farms in 1958 as hybrids between 'Admiration' and ?'Op' (A.D.S. 1981), so these two will be considered together. Their claimed common origin cannot be reconciled with the chromosome data now available. 'Op' is probably a diploid, as can be implied from the ancestry of 'Tête-à-Tête' (see above), but 'Admiration' is chromosomally unknown. It is clearly a *N. tazetta* form or hybrid (A.D.S. 1981), so could have $2n = 17, 20, 22$ or 24 chromosomes. A significant fact relevant to this problem is that 'Matador' was said by Jefferson-Brown (1969) to set seeds occasionally, which is borne out by our finding quite a high pollen fertility (66%) in this cultivar (Table 2), which is unlike any other hybrid between subgenus *Hermione* and subgenus *Narcissus*. This would suggest that 'Matador' is chromosomally similar to 'Scilly White' (also $2n = 32$), being an allotriploid between two closely related subgenus *Hermione* species or hybrids having $2n = 20$ and 22 chromosomes with a B chromosome additionally present in one of them, the $2n = 20$ plant being 'Admiration'. Thus 'Matador' is $2n = 10$ (from 'Admiration', 20) + $22 + B$ (from $22 + B$ with non-reduction) = $32 + B$ (Fig. 9B).

Because of the close affinity of the $x = 10$ and $x = 11$ genomes of subgenus *Hermione*, 'Matador' (which has $2n = 10 + 11 + 11 + B$) would behave as a cytological autotriploid and would be slightly fertile, as are other autotriploids (Brandham 1982). Clearly no subgenus *Narcissus* plant with $x = 7$ (e.g. 'Op') could have taken part in its parentage.

In the light of this suggested origin for 'Matador' the origin of the other plant produced by de Graaff, 'Golden Dawn' can be explained easily on the basis of the claimed cross, 'Admiration' (now probably $2n = 20$) \times 'Op' ($2n = 14$), with non-reduction in 'Op'. Chromosomally, 'Golden Dawn' is therefore $2n = 10$ (from 'Admiration', 20) + 14 (from 'Op', 14) = 24 (Fig. 9A).



FIG. 10. Somatic chromosomes of *Narcissus* cultivars with *N. tazetta* in their ancestry. **A** *N. x compressa*, $2n = 31$; **B** 'Silver Chimes', $2n = 28 + B$ (arrow); **C** 'Killara', $2n = 45 + B$ (arrow). Scale = $10\mu\text{m}$.

'Silver Chimes' ($2n = 28 + B$) and 'Killara' ($2n = 45 + B$). These two cultivars have one parent in common, the *N. tazetta* hybrid 'Grand Monarch' (origin and chromosome number unknown), said to be crossed with diploid *N. triandrus* to produce 'Silver Chimes' and with triploid 'Empress' to produce 'Killara' (A.D.S. 1981), but as will be shown below at least one of these crosses cannot be as claimed. The affinity of these two hybrids is nevertheless demonstrated by the fact that both carry what seems to be the same B chromosome, presumably inherited from 'Grand Monarch'. This B is very small, submetacentric, and is unusual in the genus *Narcissus* in being heterochromatic, staining much less intensely than the A chromosomes (Figs. 10B, 10C).

Taking firstly 'Silver Chimes', its $2n = 28 + B$ chromosomes could have been inherited as $n = 7$ from *N. triandrus*, which is known to be only diploid (Fernandes 1967), leaving $n = 21 + B$ as the chromosome constitution of the gamete from 'Grand Monarch'. This could also be the somatic chromosome number of this cultivar (assuming that non-reduction had taken place), since the $2n = 20$ and 22 forms of *N. tazetta* are known to hybridise to produce $2n = 21$ plants (Fernandes 1966), this one containing a B chromosome also. Thus 'Silver Chimes' is $2n = 21 + B$ (from *taz.* $20 + B \times taz.$ $22 =$ 'Grand Monarch', $21 + B$) + 7 (from *N. triandrus*, 14) = $28 + B$ (Fig. 10B). Chromo-

somally it is a very unusual plant, being a tribasic allotriploid ($2n = 3x = 11 + 10 + 7$ in addition to the B).

If 'Grand Monarch' is indeed taken to be $2n = 21 + B$ two alternatives for the production of the $2n = 45 + B$ chromosomes of 'Killara' (Fig. 10C) can be suggested as follows, neither of which can possibly involve the known $2n = 21 + B$ chromosomes of 'Empress', either non-reduced or otherwise: either a, 'Killara', $2n = 10 + B$ (from 'Grand Monarch', $21 + B$) + 35 (from an unknown subgenus *Narcissus* pentaploid with non-reduction) = $45 + B$, or b, 'Killara', $2n = 21 + B$ (from 'Grand Monarch' $21 + B$, with non-reduction) + 24 (a viable non-reduced gamete from an unknown allotriploid *N. tazetta* hybrid with $2n = 24$, similar to 'Aspasia', 'Kingcraft' etc., as detailed above) = $45 + B$.

Clearly a close investigation of 'Silver Chimes', 'Killara' and all of their putative parents is required before a definite answer to this particular problem can be reached.

Division 10.

N. × biflorus Curt. and *N. × medioluteus* Mill. (both $2n = 17$). Natural hybrids between diploid *N. tazetta* ($2n = 20$) and *N. poeticus* ($2n = 14$).

N. × intermedius Lois. ($2n = 17$). Natural hybrid between diploid *N. tazetta* ($2n = 20$) and *N. jonquilla* ($2n = 14$).

N. canaliculatus Guss. ($2n = 30$). A synonym of *N. tazetta* subsp. *lacticolor* (Steud.) Baker, and presumably autotriploid ($x = 10$).

N. × compressa Haw. ($2n = 31$). Reported by Jefferson-Brown (1969) to be a hybrid between *N. tazetta* and *N. jonquilla*. Chromosomally $2n = 10$ (from *taz.*, 20) + 21 (from triploid *N. jonquilla*, 21) = 31 (Fig. 10A).

N. × dubius Gouan ($2n = 50$). A well-known allohexaploid derived from *N. papyraceus* ($2n = 22$) and *N. juncifolius* ($2n = 14$). It arose from the fusion of a non-reduced *juncifolius* gamete ($n = 14$) with a normal *papyraceus* gamete ($n = 11$) to give a hybrid with $2n = 25$ which doubled its chromosome number to produce a plant with $2n = 50$. The genomic constitution of this fertile allohexaploid is Ju-Ju-Ju-Ju-Pa-Pa. It is widespread in north-east Spain and southern France (Fernandes 1967). Chromosomally it is $2n = [11$ (from 22) + 14 (from 14) = 25] $\times 2 = 50$.

Division 11.

'Modesta' ($2n = 30$). Affinities and origin not known. Either an autotriploid form of *N. tazetta* ($x = 10$) or an aneu-tetraploid ($2n = 4x + 2$) of a subgenus *Narcissus* plant ($x = 7$).

Pollen fertility

Fertility levels in the cultivars range between zero and well over 90% (Table 2). Plants falling into the zero category are of two types. The first is hybrids between subgenus *Hermione* and subgenus *Narcissus* (i.e. most of division 8 and a few others). These hybrids, with $2n = 17, 18, 24, 31, 45$ etc. (see above) display very little affinity between their component genomes and their resulting aberrant meiosis (Fernandes 1950) makes them almost completely

sterile. Only a few non-reduced or doubled gametes (e.g. $n = 17$ or 34 from a $2n = 17$ plant) are viable in these plants (Wylie 1952).

The second type with zero pollen fertility is division 4, the double-flowered cultivars. Most of these plants have no anthers and are obviously totally pollen-sterile. It is nevertheless possible that they can possess a fully functional and fertile female reproductive system. Those plants of division 4 which do have some anthers in the flower (e.g. 'Riotous' and 'White Lion'), even if they are distorted or partially converted into corolla tissue, achieve a reasonable degree of pollen fertility. The total amount of pollen per flower may, however, be too small for the flower to be used easily by breeders as a pollen source.

Other cultivars with very low pollen fertilities are the triploids with $2n = 21$ chromosomes. It has been established (Brandham 1982) that autotriploids can have a fairly efficient meiosis and can produce a reasonable number of viable haploid, near haploid, near diploid and diploid gametes (i.e. with $n = 7, 8, 13, 14$, less commonly 9 or 12 from a triploid with $2n = 3x = 21$). Triploid cultivars of this type are 'Bartley' (34% pollen-fertile), 'Cornet' (15%) and 'Jana' (39%). These are evidently autotriploids derived from fusion of a normal gamete with a non-reduced one either within *N. cyclamineus* or between *N. cyclamineus* and a plant very closely related to it chromosomally.

Allotriploids have two similar genomes and a third which is genetically quite different, and usually arise from a tetraploid/diploid hybridisation involving distantly related entities. During meiosis many chromosomes do not pair in allotriploids and the resulting disruption of the division causes a high level of sterility. Examples of allotriploid *Narcissus* are 'St. Keyne' (one *N. tazetta* genome and two from subgenus *Narcissus*) which is only 3% pollen-fertile, also the cross between 'Mitylene' (division 2 tetraploid) and *N. cyclamineus* which has given rise to 'Charity May' (0% pollen-fertile), 'Dove Wings' (3%) and 'Jenny' (10%) and the cross between 'Gulliver' (division 3 tetraploid) and *N. jonquilla* which has produced 'Golden Incense' (0% pollen-fertile) and 'Tittle Tattle' (8%). These data are from Table 2 and A.D.S. (1981).

Other triploids of which the pollen fertility is known (Table 2) are of obscure origin, but in the light of the above examples it may be taken that those with well over 10% pollen-fertility are fundamentally autotriploid, having a fertility similar to the known autotriploids *Aloe jucunda* (15%, Brandham 1982) and *Tradescantia acaulis* (22%, Martinez, pers. comm.), while those with 10% pollen fertility or less are more probably allotriploids with a wider hybrid origin.

The remainder of the cultivars, diploids ($2n = 14$), tetraploids ($2n = 28$) and aneuploids ($2n = 26, 27, 29$ etc.) are remarkably fertile, with pollen fertilities well over 50% in most cases and very few that are below 30% (Table 2). Indeed the only members of this particular chromosome group that we have found to have pollen fertilities which are too low to be of any practical value to the breeder are 'Fermoy' (6%), 'Golden Lacquer' (2%), 'Kilrea' (3%) and 'Mr Jinks' (2%). All of these are tetraploid, with a B chromosome in 'Kilrea' and an extra A chromosome in 'Mr Jinks'. The precise cause of the sterility of these plants cannot be established at present in the absence of information on their meiotic behaviour.

In the majority of the tetraploids which have increasing levels of fertility, moderate fertility (i.e. $\pm 50\%$) would be indicative of autotetraploidy (i.e. they have four very similar sets of chromosomes that can be expressed as AAAA). These plants form large numbers of quadrivalents during meiosis and consequently the later stages of meiosis display a high frequency of numerical

aberrations (i.e. gametes with a range of numbers which could be as wide as from $n = 10$ to $n = 18$). Abortion will result in those gametes that are excessively unbalanced chromosomally (i.e. with more than two more or fewer than the expected $n = 14$ chromosomes), with the result that overall pollen fertility is only moderate.

High fertility in tetraploids (80%+) is more indicative of allotetraploidy, in which the four genomes may be termed AABB. These tend to form bivalents at meiosis, with resultant more regular chromosome segregation and consequent high frequency of viable gametes with $n = 14$ chromosomes.

Not many diploid cultivars ($2n = 14$) derived solely from the subgenus *Narcissus* are listed in Table 2, but those whose pollen fertilities were assessed are all at least 50% fertile, with most much higher, e.g. 'Dallas' (79%), 'Glory of Lisse' (90%), 'Milan' (96%), 'Pepper' (53%), 'Sonata' (87%), the only low fertilities occurring in double-flowered forms having no anthers, e.g. 'Doublet', 'Irene Copeland' and 'White Sail'. The diploid cultivars ($2n = 17$) involving subgenus *Hermione* crossed with subgenus *Narcissus* are quite sterile apart from some non-reduced gametes that are viable.

Polyploid frequency in Narcissus cultivars compared with that in other ornamentals

As can be seen from the information given in this article the optimum chromosome number for cultivated hybrid forms of *Narcissus* is clearly at or near the tetraploid level, a trend noted as early as 1923 by de Mol who was working on a relatively small amount of chromosomal information. This tendency towards the selection of polyploid forms of ornamental plants from diploid or lower polyploid wild ancestors is quite widespread, with the following being typical examples: *Dahlia*, cultivars octoploid ($2n = 64$), from tetraploid wild ancestors; *Chrysanthemum*, all high polyploids, with large-flowered cultivars having more chromosomes, approximately $5x-7x$ with much aneuploidy; Roses, largely polyploid; bearded *Iris* (*Pogoniris* group), originally diploid but now almost entirely tetraploid (these examples from Darlington 1963); *Kalanchoe*, wild species diploid and tetraploid ($2n = 34, 68$) in similar numbers but the great majority of cultivars circa-tetraploid or higher polyploids ($2n = 66, 67, 68, 69, 72, 75, 84, 85, 96$) (van Voorst & Arends 1982).

The optimal level of polyploidy of cultivars of all of the above plants is tetraploid, as in *Narcissus*, or higher, but some cultivated ornamentals do not follow this pattern, with chromosome numbers below the tetraploid level being more common. Cultivars of *Canna* and *Bougainvillea*, for instance, are mostly diploid (Khoshoo & Zadoo 1969, Khoshoo & Mukherjee 1970), and also the sweet pea *Lathyrus odoratus*, in which tetraploid forms occur occasionally but are almost totally sterile and quite useless for breeding purposes (Darlington 1963). *Tulipa*, which might be regarded as a close equivalent to *Narcissus* with respect to its uses in cultivation and its many hundreds of named cultivars, is in fact very different from *Narcissus* as far as its optimum level of polyploidy in cultivation is concerned. In a major study of this genus Plavcova (1969) counted the chromosomes of 560 cultivars, finding 507 diploids ($2n = 24$), 51 triploids ($2n = 36$) and only two tetraploids ($2n = 48$), the last being the cultivars 'Sunburst' and 'Mrs. John T. Scheepers'. Clearly tetraploids are not a great success in *Tulipa* and the triploid cultivars have arisen through non-reduced gametes from diploids as in many instances in *Narcissus* rather than through diploid-tetraploid hybridisation.

Crocus cultivars appear largely to be diploid selections from diploid parental species or hybrids, with polyploids being known but uncommon (Brighton *et al.* 1980). *Primula sinensis* is known as a series of diploid and tetraploid cultivars (Darlington 1963), as is *Pelargonium*, but in the latter the preferential selection of tetraploids seems to have occurred only in part of the genus. Daker (1969) showed that 91% of the *Pelargonium* cultivars based on $x = 9$ were diploids, a frequency identical to the percentage of diploids among his sample of *Pelargonium* species with $x = 9$, but in $x = 11$ forms of the same genus he found 62% diploids among the species, but only 22% diploids among the cultivars, the majority of the cultivars being tetraploid. In *Nerine*, wild species are diploid only ($2n = 22$), and most cultivars are also diploid, although a small number of tetraploids ($2n = 44$) and triploids or near-triploids have been selected (Roberts 1984).

Hyacinthus is unique as far as chromosome number distribution among its cultivars is concerned, and Darlington *et al.* (1951), in a survey of 106 cultivars, demonstrated a complete range, all chromosome numbers occurring between $2n = 2x = 16$ and $2n = 4x = 32$ with major peaks of frequency at the diploid and triploid levels ($2n =$ circa 16 and circa 24 respectively). Early cultivars (dating from 1550 to 1850) were diploid only, but triploids arose from these through non-reduction (1700–1900). The autotriploids ($2n = 24$) are unusually fertile; indeed there is some unpublished suspicion based on their chromosome morphology that they are actually hexaploids, which would explain their high fertility. They could be crossed successfully with diploids to produce a range of hypotriploids (1800–1920) with $2n = 17 - 23$, a result similar to that observed in triploid \times diploid crosses in the *Liliaceae/Aloineae* (Brandham 1982). They were also crossed with each other to produce the lower hypertriploids (1850–1920), with perhaps $2n = 25 - 28$, and the upper end of the $2n = 25 - 28$ range could be intercrossed to give chromosome numbers approaching the tetraploid level of $2n = 32$ (1900–1950) but so far only one plant has actually reached it, the cultivar 'Blue Giant' (Darlington 1963).

The reason why the selection of cultivars of *Narcissus* has evidently stabilised largely at the tetraploid level with very few higher polyploids is not completely clear. Presumably the larger size and greater vigour characteristic of most tetraploids applies in this genus also, resulting in their being selected preferentially over diploids, but it is quite evident that most of the higher polyploids that doubtless arise from the tetraploids show a reduction in size and/or vigour (Wylie 1952) and are not usually chosen for vegetative propagation by breeders when they appear in the progeny of their plants. In view of the comment by Darlington (1963: 107) that the only species of *Narcissus* to have autopolyploid races rising frequently above the tetraploid level (i.e. excluding the allohexaploid *N. dubius* and hexaploids occurring very rarely in other species) is *N. bulbocodium*, which has among the smallest chromosomes in the entire genus, it is possible that the generally large chromosomes in the rest of the genus constitute a nucleus that is too large for the efficiency and normal speed of DNA replication and/or mitosis to be maintained fully at levels above the tetraploid. The result of the loss of efficiency caused by exceeding this tetraploid nuclear volume is a slowing down of the mitotic system and growth rate and hence a smaller, weaker plant of little value to the breeder.

The chromosomes of *Narcissus* and *Tulipa* are of the same order of size, but the basic numbers of the two genera differ widely ($x = 7$ in most *Narcissus* species and $x = 12$ in *Tulipa*). It is therefore not surprising that the most

efficient nuclear size, found at the tetraploid level ($2n = 4x = 28$) in *Narcissus*, should be almost matched by a similarly critical size at the diploid level in *Tulipa* ($2n = 2x = 24$), since the chromosome numbers and nuclear volumes of tetraploid *Narcissus* and diploid *Tulipa* are comparable. It is therefore a tenable hypothesis that in these genera and in related monocotyledons there is a critical DNA level that cannot be exceeded without a severe impairment of mitotic efficiency, this level coming into operation at different ploidy levels depending on absolute chromosome size and basic number. Critical nuclear size could perhaps explain why no wild plants of *Narcissus* subgenus *Hermione* have been found at the tetraploid level ($2n = 40$ or 44). These would greatly exceed the optimal DNA amount and would thus be too slow-growing to compete with diploids when they arise.

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