# CHROMOSOME NUMBERS IN TASMANIAN GOODENIACEAE AND BRUNONIACEAE

Ву

W. D. JACKSON

Department of Botany. University of Tasmania

(WITH 1 TEXT FIGURE)

#### ABSTRACT

Species of Goodenia, Velleia, Selliera and Scaevola have a basic chromosome number of 8, whilst Dampiera and Brunonia show a basic number of 9. With the exception of Selliera radicans which has diploid and hexaploid races, all others are diploid.

#### INTRODUCTION

Thirteen members of the Goodeniaceae and the monotypic *Brunonia australis* have been recorded from Tasmania. None are endemic. Four species are rare or are doubtful records. Four are restricted in distribution, while the other five are common species with wide distributions. With the possible exception of the sub-herbaceous Goodenias, the Tasmanian members offer few taxonomic difficulties. The principle interest lies in the relationships between genera and the position of *Brunonia* which has in the past been grouped with the Goodeniaceae.

At present there is no published account of chromosome numbers in this distinctive Australian group. However, P. Martin (University of Adelaide) is at present working on the Australian members.

#### MATERIALS AND METHODS

Only Tasmanian specimens were studied. Where possible material was obtained from different environments and from a wide range in the Tasmanian distribution. Morphological variants common in Goodenia humilis, G. elongata and G. lanata were investigated.

Meiosis in pollen mother cells was studied using aceto-orcein and aceto-lacmoid smear methods (Darlington and La Cour, 1947). Root tips were used for the study of mitosis, both untreated and with treatments of aqueous solutions of paradichlorobenzene (Meyer, 1945) before fixation in acetic-alcohol. Acetic stains were used for most of the investigation. Feulgen staining was tried, but no satisfactory stain intensity was obtained.

#### OBSERVATIONS AND DISCUSSION

A summary of the counts determined and the origin of the material is given in Table I. Drawings of representative plates are provided in the text figure.

In the Goodeniaceae the chromosome complement is surprisingly uniform in both number and morphology. Apart from *Dampiera stricta* with 2n = 18, the Tasmanian members investigated all have

the diploid number 2n=16, except in Selliera radicans where a hexaploid 2n=48 was found in addition to the diploid form. There do not appear to be any significant morphological differences between these two chromosome races, which were found in the same locality. The chromosome number of Brunonia australis is 2n=18. Meiosis was normal in all specimens and pollen fertility, as measured by stainable pollen, was high. There is no evidence of apomixis in the members studied.

The chromosomes are fairly small. In mitotic metaphase the chromosomes range from 2·5-4  $\mu$  in length. Mitotic plates are small and compact. Narrow meiotic spindles are a feature of the Goodeniaceae, with cross sectional areas 1/20th that of the pollen mother cell section. This feature while common to all species of the Goodeniaceae studied is not found in Brunonia australis.

As mentioned above, the chromosome complement in the Goodeniaceae is markedly uniform. Although Dampiera stricta has a haploid complement of nine, studies of chromosome morphology indicate that eight of these are indistinguishable from the genome in other genera. All the chromosomes are metacentric, with four shorter chromosomes with a 5/4 arm ratio. One of the four long members has a 1/1 arm ratio while the ratio in the other three is approximately 3/2. The long arm of one of these has a prominent trabant. In Dampiera stricta the complement appears to be identical apart from the addition of another short chromosome with a 1/1 arm ratio. Martin (unpub.) has found polyploid series based on x = 9in mainland members of Leschenaultia and Dam-Brunonia australis appears to possess piera. eight members similar to that in the eight membered Goodeniaceae. There the additional member to the haploid set is a long chromosome with a 5/4 arm ratio and a heterochromatic segment near the distal end of the short arm.

Practically all chiasmata are terminal or nearly so. In Brunonia the distal localization of chiasmata is most pronounced. Brunonia australis also differs from the members of the Goodeniaceae studied in having a larger cell size and a relatively larger spindle. With these exceptions the group appears to be homogeneous cytologically, based on x=8 and x=9, with some apparent uniformity in chromosome morphology.

I wish to acknowledge the kind assistance of Mr. T. E. Burns in collecting material in Northern Tasmania.

TABLE I

Chromosome numbers of Tasmanian Goodeniaceae and Brunoniaceae.

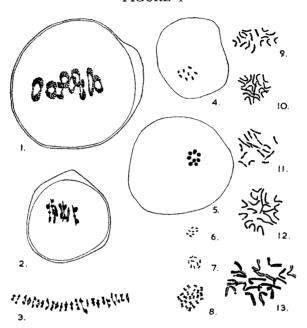
Species	Collecting locality altitude	and	Remarks	Chromosome number	
Goodeniaceae Velleia, Sm.				2n	n
V. montana, Hook. f.	King William Saddle Interlaken	$\frac{2,800}{2,600}$	robust lvs. 10 cm. reduced lvs. 4-5 cm.	16 16	8 8
V. paradoxa, R.Br.	Springfield, Glenorchy Georgetown	200 100	robust lvs. 14 cm. glabrous lvs. 7 cm. hairy entire ±	16 16	8 8 8
Goodenia, Sm.	desigetown	100	175. 1 cm. nany choice	- 10	
G. ovata, Sm.	Tolosa St., Glenorchy	200	typical	16	8
G. amplexans, F.v.M. G. barbata, R.Br. G. geniculata, R.Br.	St. Marys Not collected Not collected Not collected	800	typical	16	h
G. lanata, R.Br.	Mt. Nelson Murdunna	300 50	typical lys. 3 cm. x 2·5 cm.	16 16	8 8
G. elongata, Labill.	Low Head St. Helens	50 50	lvs. 4 cm. x 1.5 cm. hairy reduced	16 16	8
G. humilis, R.Br.	Low Head Low Head	100 20	± glabrous robust typical lvs. 3 cm.	16 16 16	8 8 8
Selliera, Cav.	Low Head	20	robust lvs. 5 cm. x 0.5 cm.	16	<b>8</b>
S. radicans, Cav.	Glenorchy Glenorchy		typical typical	16 48	8 24
Scaevola, Linn. S. hookeri, F.v.M.	Mt. Wellington	4,000	reduced	16	8
Dampiera, R.Br. D. stricta, R.Br.	Collins Cap Bicheno	1,900 50	robust     typical	16 18	9
BRUNONIACEAE Brunonia, Sm. B. australis, Sm.	Queechy Hill, Laun.	50	typical	18	9

#### REFERENCES

Darlington, C. D. and La Cour, L. F., 1947.—The Handling of Chromosomes. London, Allen and Unwin.

MEYER, J. R., 1945.—Prefixing with Paradichlorobenzene to Facilitate Chromosome Study. Stain Technology, 20, 121-125.

#### FIGURE 1



401

#### Side View

#### Metaphase I

- 1. Brunonia australis n=9 2. Goodenia ovata n=8 3. Selliera radicans n=24

### Polar View

## Metaphase II

- 4. Goodenia kumilis n = 8
  5. Velleia montana n = 8
  6. Goodenia ovata n = 8
  7. Dampiera stricta n = 9
  8. Selliera radicans n = 24.

#### Polar View

#### Mitotic Metaphase

- 9. Velleia montana 2n = 16 10. Goodenia lanata 2n = 16
- 11. Goodenia elongata 2n = 16
  12. Dampiera stricta 2n = 18
  13. Scaevola hookeri 2n = 16

