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# Chromosome variation in *Araceae*: V\*

## ACOREAE TO LASIEAE

C. J. MARCHANT†

*Summary.* Somatic chromosome numbers are reported for six tribes of the *Araceae* and for miscellaneous genera which were not included earlier in the series. While much chromosome diversity is evident in the family, the chromosome data do not support completely the major *Araceae* classifications of either Hutchinson (1959) or Engler (1920). Evolutionary relationships in the family are considered from the viewpoint of known basic chromosome numbers and suggestions are made concerning the taxonomic positions of some critical genera especially in the proposed *Arophyteae* of Bogner (1972).

### INTRODUCTION

This is the last of a series of five papers describing karyotypes of miscellaneous genera in the *Araceae*. The purposes of this chromosome survey of an entire family were set out in the introduction to the first paper (Marchant, 1970) and the very diverse karyotypes and basic numbers, with occasional intra-specific diversity, have become apparent (Marchant, 1970, 1971a & b, & 1972). The cytological examination of genera is far from complete, there being some 54 genera not available at Kew, but it is hoped that these data for a little over 50 per cent. of the total genera will provide a stimulus for further work at the same comparative level.

The present paper deals with the first six tribes in the classification of Hutchinson (1959), most of which are considered by him to be the more primitive in the family. There are also a number of miscellaneous additions from tribes already treated in this series. These plants have had their chromosomes counted since the results of the survey first began to be published.

### MATERIALS AND METHODS

Materials and methods were both as previously described (Marchant, 1970). Voucher specimens are deposited in the Kew Herbarium.

### RESULTS

#### *Acoreae*

*Acorus calamus* L. has a curiously mixed record of chromosome numbers accorded to it by numerous authors (listed in Table 1). These range from  $2n = 18$  ( $x = 9$ ) to  $2n = 24$  ( $x = 12$ ) and  $2n = 36, 44$  and  $48$  ( $x = 11$  or  $12$ ). The majority, with  $2n = 36$ , correspond with my own count of  $2n = c. 36$  very small chromosomes (Fig. 1/A, p. 202). It is important that chromosomes of other species in the genus should be counted to help verify the correct basic number. The most likely explanation seems to be that widespread European  $2n = 36$  plants, which are sterile, are triploids based on  $x = 12$  (Palmgren, 1943) in the same chromosome series as  $2n = 24$  and  $48$ . The

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† Now at the University of British Columbia, Vancouver 8, British Columbia, Canada.

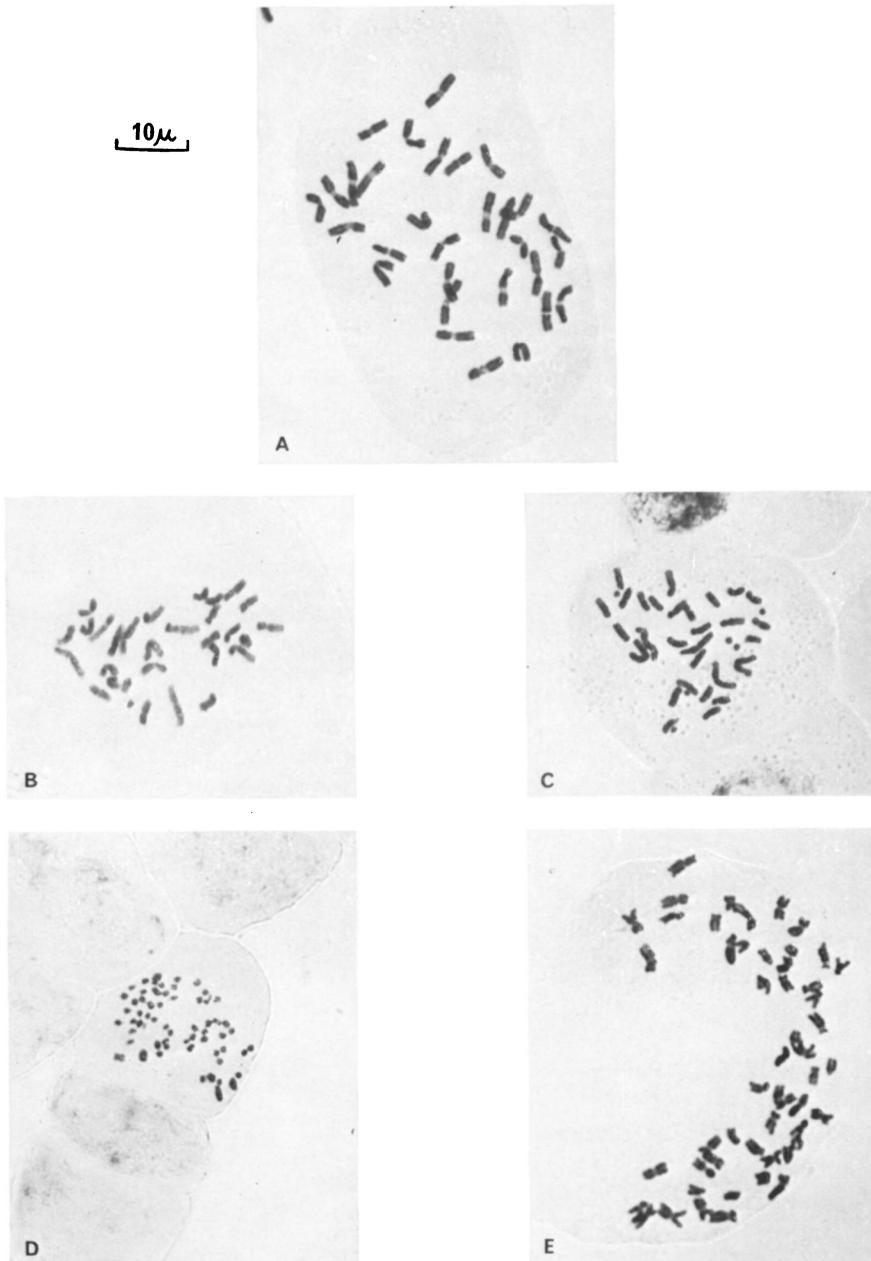






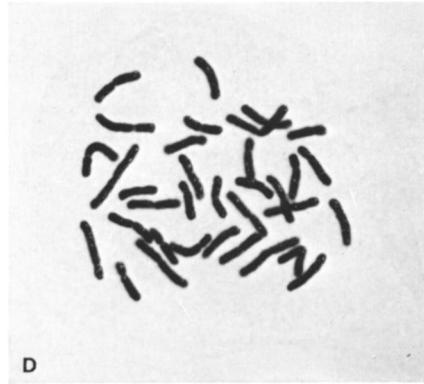
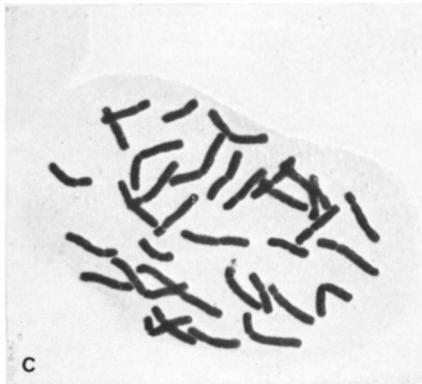
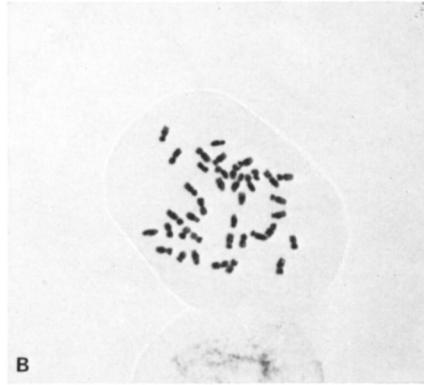
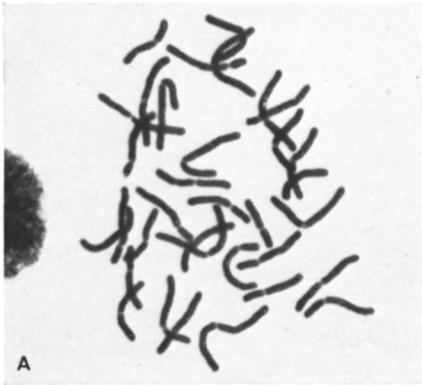
FIG. 1. Mitotic chromosome complements in some *Araceae*. **A**, *Acorus calamus* ( $2n = \text{ca. } 36$ ); **B**, *Lysichiton americanum* ( $2n = 28$ ); **C**, *Orontium aquaticum* ( $2n = 26$ ); **D**, *Spathiphyllum wallisii* ( $2n = 30$ ); **E**, *Anthurium microphylla* ( $2n = 30 + 1f$ ); **F**, *Urospatha* sp. ( $2n = 52$ ).

## PLATE 14



Somatic chromosome complements from root tips in the tribes *Spathiphyllae*, *Anthurieae*, *Dracontieae* and *Lasieae*. **A**, *Spathiphyllum* × *hybridum*, 69.791 ( $2n = 30$ ); **B**, *Anthurium signatum*, 68.1642 ( $2n = 30 + 1f$ ); **C**, *A. harrisii*, 69.103 ( $2n = 30 + 5f$ ); **D**, *Symplocarpus foetidus*, 69.456 ( $2n = 60$ ); **E**, *Urospatha sagittifolia*, 68.1411 ( $2n = 52$ ).

## PLATE 15



10  $\mu$

Somatic chromosome complements from root tips in the tribes *Stylochitoneae*, *Pothoeae* and *Colocasieae*. **A**, *Microculcas marattioides*, 69.785 ( $2n = 34$ ); **B**, *Arophyton humbertii*, 70.64 ( $2n = 38$ ); **C**, *Gonatopus boivinii*, 70.27 ( $2n = 34$ ); **D**, *Zamioculcas zamifolia*, 70.30 ( $2n = 34$ ).

counts of  $2n = 44$  by Larsen (1969) for *Acorus calamus* in Thailand and by Ito (1942) for *A. asiaticus* Nakai apparently have a distinct basic number of  $x = 11$ .

#### *Oronteeae*

A specimen of *Lysichiton americanum* Hulten & St. John had  $2n = 28$  small chromosomes (Fig. 1/B) in agreement with the  $2n = 28$  report of Löve & Kawano (1961). In contrast in the same tribe, *Orontium aquaticum* L. (Fig. 1/3) has  $2n = 26$  and large chromosomes agreeing with Cave (1967). However, Delay (1951) reports  $2n = 24$ , and Huttleston (in Darlington & Wyllie, 1955)  $2n = 28$ .

#### *Spathiphyllaeae*

Three species of *Spathiphyllum* Schott and a hybrid have chromosome numbers based on  $x = 15$ , all with  $2n = 30$  medium-sized chromosomes (Fig. 1/D, p. 202) except the tetraploid *S. floribundum* (Linden & André) N. E. Br. with  $2n = 60$ . The hybrid, *S. × hybridum* N. E. Br., also has  $2n = 30$  (Plate 14/A) but its parental species were not available for examination and comparison. An interesting and distinctive cytological feature in *Spathiphyllum* is the pericentric region of heterochromatin in several chromosomes of the complement (Plate 14/A). Such pericentric heterochromatin has been reported in the insect, *Drosophila* (Hannah, 1951) but is not commonly so clearly visible and this is the first example observed in the *Araceae*.

#### *Anthurieae*

The genus *Anthurium* Schott clearly has a base number of  $x = 15$ , usually with  $2n = 30$  or  $2n = 60$  small chromosomes. However, there are two species (*A. gracile* (Rudge) Lindl. and *A. scolopendrinum* (W. Ham.) Kunth) with  $2n = 40$ . There are also two polyploids with  $2n = c. 124$  ( $x = 15$ ), *A. lucidum* Kunth and *Anthurium* sp., not previously recorded. The most extensive list of previous chromosome counts is that of Gaiser (1927) who cites approximate counts for about 37 species, all of them based on  $x = 15$ . In addition to their counts of  $2n = 30$  and 60, Sharma & Bhattacharya (1966) cite two species with  $2n = 28$ , but their poor karyotype illustrations cast doubt on the accuracy of these numbers. Other authors cite  $2n = 34$  (Mookerjea, 1955) for *A. crystallinum* Linden & André and *A. signatum* C. Koch & Mathieu and  $2n = 24$  (Pfitzer, 1957) for *A. scandens* (Aubl.) Engl. Such aneuploid variation is almost certainly due in this case to the presence of B-chromosomes, ranging from 1 to 5 in number, which I have recorded in five species with  $2n = 30$  chromosomes (Plate 14/B & C and Fig. 1/E, p. 202) and in one  $2n = 60$  species. However, in the two species recorded as  $2n = 40$  there is no evidence for the presence of B-chromosomes and the basic number difference seems to be a shift in the A-chromosome complement, possibly an upward trend to  $x = 20$ .

#### *Dracontieae*

Two of the four tribal genera had their chromosomes counted. *Dracontium foecundum* Hook. f. had  $2n = 26$  medium sized chromosomes, no previous counts being available for comparison. *Symplocarpus foetidus* (L.) Salisb. has two previously recorded ploidy levels with numbers of  $2n = 30$  (Ito, 1942)

and  $2n = 60$  (Mulligan, 1965). My own result of  $2n = 60$  small chromosomes (Plate 14/D) verifies the higher level of polyploidy already recorded in the northern collection of Mulligan (*l.c.*).

#### *Lasieae*

Only two of the seven genera in this largely tropical tribe were studied. Two accessions of *Cyrtosperma johnstonii* (Bull) N. E. Br. had  $2n = 26$  small chromosomes in accord with a previous count of  $2n = 26$  for *C. senegalense* (Schott) Engl. by Mangenot & Mangenot (1962). These indicate a basic number of  $x = 13$ , a base which appears again in *Urospatha* Schott in two species, *U. sagittifolia* (Rodsch.) Schott and *Urospatha sp.* both with  $2n = 52$  medium sized chromosomes (Plate 14/E and Fig. 1/F). No previous counts are available.

#### *Tribe not specified by Hutchinson (1959)*

A relatively recently described genus, *Microculcas* Peter had a chromosome count showing  $2n = 34$  large chromosomes in *Microculcas marattioides* Peter (Plate 15/A). This is an interesting link with the  $2n = 34$  large chromosomes in *Zamioculcas* Schott (Plate 15/D), with which there is considerable morphological affinity.

### ADDITIONAL RESULTS THROUGHOUT THE FAMILY

During the course of this long-term survey a number of counts have accumulated for additional species not already included in earlier papers of the series. They are presented in Table 2 and discussed below.

#### *Pothoeae*

*Pothos chapelieri* Schott from Madagascar has  $2n = 24$  small chromosomes with a base number of  $x = 12$ . This is in accord with  $2n = 24$  in *P. aff. scandenti* L. reported previously (Marchant, 1970).

#### *Calleae*

*Pycnospatha soerenseii* S. Y. Hu has  $2n = 26$  ( $x = 13$ ). This is a very different basic number from the only other member of this tribe, *Calleae*, which, in *Calla palustris* with  $2n = 72$  (Marchant, 1970), has  $x = 9$  (or possibly 6).

#### *Stylochitoneae*

Some members of this group of genera from tropical Africa and Madagascar, were discussed in Paper 1. Further counts for *Arophyton tripartitum* Jumelle and *A. humberii* Bogner, are  $2n = 38$  with small chromosomes (Plate 15/B) indicating a secondary basic number of  $x = 19$ . This basic number was also recorded in *A. tripartitum* previously with  $2n = ca. 76$  (Marchant, 1970), but it is not in agreement with  $2n = 40$  ( $x = 10$  or  $20$ ) in *A. buchettii* Bogner (Marchant, *l.c.*). There may well be simple duplication (tetrasomy), or two B-chromosomes in the latter species, but this has not been established. It is of interest that two other genera of the tribe, *Carlephyton madagascariense* Jumelle with  $2n = 108$  (Marchant, 1970), *C. glaucophyllum* Bogner with  $2n = 54$  (Table 2) and *Colletogyne perrieri* Buchet have a base of

TABLE 2. Additional counts from tribes covered in earlier parts (Marchant, 1970, 1971a, 1971b and 1972).

|   | 677.67       | 70.958  | Madagascar, Bogner s.n.   | 24 | 12 | S |        | Larsen          | Unpub. |
|---|--------------|---------|---|----|----|---|--------|-----------------|--------|
| Tribe <b>Pothoeae</b><br><i>Peltos chapetieri</i> Schott                                      |              |         |   |    |    |   |        |                 |        |
| Tribe <b>Callieae</b><br><i>Pycnospatha soerenseni</i> S. Y. Hu                               | 094-71-01039 | 71.1584 | Thailand, Makam-Dist. Village, Makam-Dist. Province Chantaburi, Bogner 395      | 26 | 13 | ? | 26     | Larsen          |        |
| Tribe <b>Stylochitonaceae</b><br><i>Arophyton tripartitum</i> Jumelle var. <i>tripartitum</i> | 129.69-00982 | 69.714  | Madagascar, Presqu'île de Masoala, Mont Ambato, Bogner 272                      | 38 | 19 | S | 38     | Kress in Bogner | 1972   |
| <i>A. humbertii</i> Bogner  | 639.69-5385  | 70.64   | Madagascar, 38 Reserve Naturelle No. 12 (Massif du Marojezy), Bogner 163        | 38 |    |   |        |                 |        |
| <i>A. crassifolium</i> (Buchet) Bogner ( <i>Humbertina crassifolia</i> Buchet)                | 639.69-5386  | 70.63   | Madagascar, Massif de l'Ankarana, 20-30 m. (type locality), Bogner 276          | 54 | 9  |   |        |                 |        |
| <i>Colletogyne perrieri</i> Buchet  | 711.67       | 69.96   | Madagascar, Montagne des Français, Bogner 165                                   | 54 | 9  |   |        |                 |        |
| <i>Carléphyton glaucophyllum</i> Bogner   | 719.67       | 69.658  | Madagascar, Reserve Naturelle No. 12, Massif de l'Ankarana, Bogner 167          | 54 | 9  | S | c. 108 | Kress in Bogner | 1972   |
| <i>Zamiocalcas zamitifolia</i> (Lodd.) Engl.  | 630.67       | 70.30   | Tanzania, limestone formation between Kange & Amboni (Tanga Prov.) Bogner, s.n. | 34 | 17 | L |        |                 |        |
| <i>Zamiocalcas zamitifolia</i> (Lodd.) Engl.  | 494.67       | 70.28   | Usambara mountains, Tanzania, Bogner, s.n.                                      | 34 |    |   |        |                 |        |
| Tribe <b>Diefenbachieae</b><br><i>Gonatopus boivinii</i> (Decne.) Hook. f.                    | 301.68       | 70.27   | Tanzania, near Kwamtili, East Usambaras, Bogner 219                             | 34 | 17 | L |        |                 |        |
| Tribe <b>Colocastieae</b><br><i>Xanthosoma brasiliense</i> (Desf.) Engl.                      | 797.58       | 69.97   | Dominica, Campbell  | 26 | 13 | M |        |                 |        |
| Tribe <b>Areaceae</b><br><i>Arum cyrenaicum</i> Hruby   |              |         | Cyrenaica, Gebel Akhdar, L. Boudos  | 56 | 7  | S |        |                 |        |

$x = 9$  but it would be premature to suggest, without further research that the  $x = 19$  of most *Arophyton* species is in some way derived from a combination of  $x = 10$  as in *A. buchettii* and  $x = 9$  in related *Carlephyton*. Further study is needed in this group.

Two further counts of  $2n = 34$  in *Zamioculcas zamiifolia* (Lodd.) Engl. confirm my earlier citation for this genus (Marchant, 1970). *Microculcas marattioides* described above with a closely similar karyotype and the same chromosome number, appears to be closely related on this basis and should be included in the same tribe. In the same way *Gonatopus boivinii* (Decne.) Hook. f. with  $2n = 34$  large chromosomes (Plate 15/C) and *Heterolobium petiolulatum* Peter cited previously (Marchant, 1971a) seem to belong in the *Stylochitoneae* and not the *Dieffenbachieae*.

A count of  $2n = 54$  was reported previously (Marchant, 1972) for *Humbertina crassifolia* Buchet. Bogner (1972) has decided to move this species to the genus *Arophyton* as *A. crassifolium* (Buchet) Bogner. This involves a change of tribe, for the genus *Humbertina* was placed in the *Areae* by Hutchinson (1959). This change is in better agreement with the basic number of  $x = 9$  which, though rare in the *Areae* (only *Typhonium* and a few *Cryptocoryne* spp.), is common in the *Stylochitoneae*. However, at the generic level the switch to *Arophyton* does introduce yet another basic number ( $x = 9$ ) to an already confused mixture of  $x = 19$  and  $x = 10$  or  $20$  (Marchant, 1970). It seems to me that  $2n = c. 76$  in *A. tripartitum* Jumelle is  $x = 19$ , in accord with *A. hundertii* and *A. rhizomatosum*. This is probably a secondary basic number derived at some point in evolution from combining  $x = 9$  with  $x = 10$  in other species.

#### *Colocasieae*

The count of  $2n = 26$  for *Xanthosoma brasiliense* (Desf.) Engl. accords with my earlier report of  $2n = 26$  for three other species of the genus (Marchant, 1970).

#### *Areae*

A specimen of *Arum cyrenaicum* Hruby from Cyrenaica has  $2n = 28$  small chromosomes in accord with the basic number ( $x = 7$ ) of my previous counts for the genus (Marchant, 1972) and with those of other authors.

### DISCUSSION

In this final chapter of this series tabulating mitotic karyotypes and basic numbers in the *Araceae* I intend to make some attempt to relate the cytological data to existing taxonomic classifications. It is impossible without a specialist knowledge to suggest profound changes in arrangement, though these may well be necessary in this poorly understood family. Instead I will indicate instances where widely separated genera have similar karyotypes or where cytological data support or contradict changes suggested by taxonomists.

I would first like to correct a misconception in Paper 1 (Marchant, 1970) where I erroneously concluded that *Scindapsus pictus* Hassk. ( $2n = ca. 110$ ) had a base number of  $x = 10$ . This is a poor numerical interpretation of the facts. In accord with chromosome data from other sources cited in that

paper it now seems much more likely that the base number is  $x = 7$  and the specimen of *S. pictus* with  $2n = ca. 110$  would then be 16-ploid.

In the *Richardeae*, the genera *Nephtytis* Schott and *Anchomanes* Schott, both with  $2n = 40$ , have extremely similar chromosome size and karyotype agreeing with Hutchinson's classification. However, it seems much more logical to separate *Homalomena*, again with  $x = 10$  but with small chromosomes, as Engler (1920) has done.

The genus *Protarum* Engl. in the *Areae*, with  $2n = 28$  seems to relate chromosomally to *Alocasia* Neck. ( $2n = 28$ ) in the *Colocasieae*, as Bogner (unpublished) suggests on morphological grounds.

As has already been mentioned above, the genera *Zamioculcas* and *Gonatopus* are separated by Hutchinson in the *Stylochitoneae* and *Dieffenbachieae* respectively, while *Microculcas* was not placed by him in a tribe; yet each has  $x = 17$  large chromosomes and closely similar karyotypes. So far, in Hutchinson's *Dieffenbachieae*, *Dieffenbachia* Schott is the only other genus recorded with  $x = 17$  and is clearly not closely related morphologically to *Gonatopus*. It therefore seems logical to remove *Gonatopus* from the *Dieffenbachieae* and put it with *Microculcas* and *Zamioculcas*. These three would fit best in Engler's subfamily *Pothoideae*, tribe *Zamioculcaseae*.

From the chromosome data cited in this paper, and from the conclusions of Bogner (personal comm.) it appears that *Zamioculcas*, *Gonatopus* and *Microculcas* should be removed as a group from the *Stylochitoneae* as defined by Hutchinson (1959), while *Humbertina* must be included in the genus *Arophyton* and in the tribe *Arophyteae*; the latter genus seems misplaced in the *Stylochitoneae* by Hutchinson. Several of the genera now included by him in the *Stylochitoneae* were not known at the time of Engler's publication (1920) and have since been assigned to the tribe *Arophyteae* Lemée (formerly *Synandrodieae* Buchet). The *Arophyteae* would fit neatly next to the *Stylochitoneae* of Engler's subfamily *Aroideae*.

It is clear that neither of the major classifications of the *Araceae* are adequate to circumscribe the family. Whereas Hutchinson's treatment of the family is too simple, with too many genera lumped into too few groups at the tribal level, Engler (1920) goes to the opposite extreme with a breakdown into many small and diversified units. However, Engler's system, using floral characters, vegetative morphology and anatomy as criteria, was much more broadly based than that of Hutchinson who relies solely on floral structure. This is borne out also by the comprehensive pollen morphological study of Thanikaimoni (1969) which also supports Engler's system.

Amongst the chromosomal information accumulated in this investigation there is much diversity; yet extreme asymmetry (telocentrics) is absent and strict metacentrics are rare. On the basis of the kind of karyotypes encountered in those genera contrasted by taxonomists as primitive or derived it is hard to formulate any obvious evolutionary trends. This inadequacy of chromosome information serves only to emphasize our present-day lack of knowledge of chromosomes as evolutionary indicators over a broad spectrum of genera, as for example in a family.

Some insight can be gained from the consecutive organization of the basic numbers of the various genera, as indeed has already been done by Larsen (1969) using the data available at that time. Now that just over 50 per cent. of the genera of the *Araceae* have been studied cytologically I agree with Larsen (*l.c.*) that  $x = 7$  is the most common basic number. This is followed

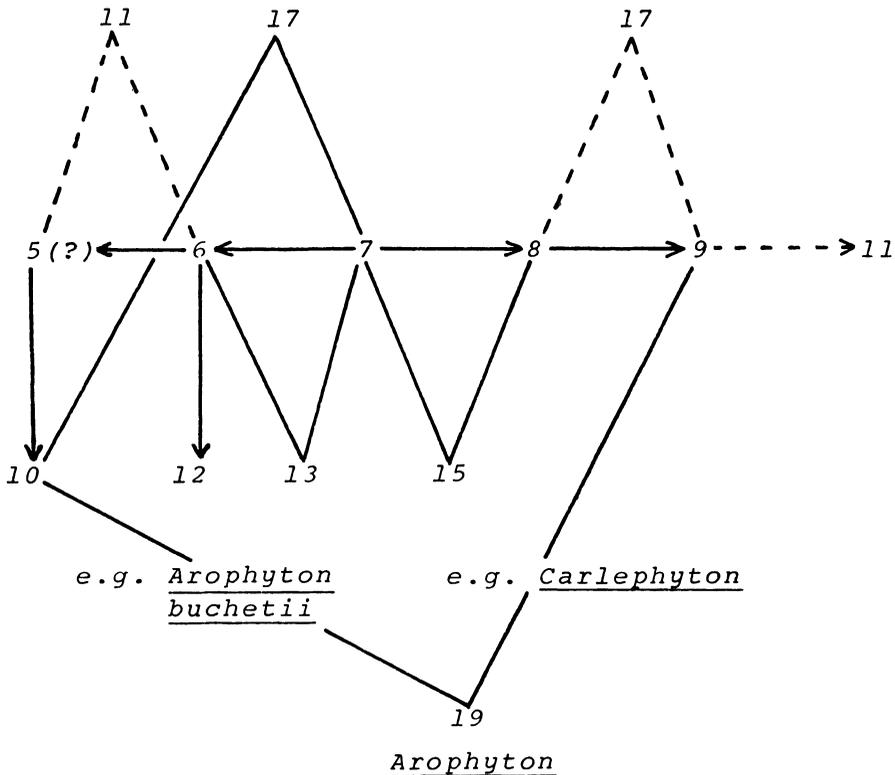


FIG. 2. Possible basic number relationships and direction of their evolution in the *Araceae*. Dotted lines indicate very uncertain derivations.

in frequency by  $x = 13$ . My diagram in Figure 2 (above) reflects the considerations of Larsen for basic number evolution in *Araceae*, while at the same time I have incorporated several higher, secondary, basic numbers. The relationships suggested are only tentative and some alternatives are indicated by broken lines.

Obviously my own survey of Araceous chromosomes, despite its range, is incomplete in terms of coverage of the whole family and in terms of representation within some individual groups. Nevertheless, it has helped to resolve differences in chromosome numbers published by previous authors and gives no support to the claims of widespread intraspecific and intraplant aneusomaty and aneuploidy by A. K. Sharma and his associates (Sharma & Das, 1954; Mookerjea, 1955; Sharma & Mukhopadhyay, 1965; Sharma & Bhattacharya, 1966). I believe this contribution of chromosome numbers to be important in fostering a deeper knowledge of the family through a co-operative multidisciplinary approach. It is intended to encourage a wider interest in, and investigation and interpretation of the phylogeny and evolution of the *Araceae*.

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