

# Phylogenetic relationships within Turneraceae based on morphological characters with emphasis on seed micromorphology

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**Abstract** Genera of Turneraceae differ notably in connation/adnation of calyx, corolla, and androecium. Floral and seed morphology were analyzed in all genera. Phylogenetic analyses were made using a matrix of 91 characters coded for 102 taxa including all genera of Turneraceae and all series of *Turnera*. Our goals were: assessing the impact of morphology in the cladistic analyses of Turneraceae and comparing our results with those based on molecular datasets. Our analyses suggest that all genera are monophyletic. The inclusion of seed micromorphology in the analyses increased resolution within *Turnera*, the strict consensus tree shows four main clades, each gathering two or more current series. A comparison of morphological and molecular trees is difficult to make due to the great differences in taxon sampling. However, some clades or subclades are consistent in both phylogenetic approaches. Apparently, the formation of a floral tube conferred an evolutionary advantage to the Turneraceae, because it developed in 66 % of the genera. The morphological complexity of the tube increased in several steps: (1) adnation

of petal claws to calyx, developing a perianth tube; (2) partial adnation of stamens to the perianth tube; (3) fusion of sepal and petal veins, shaping a 10-veined perianth tube; (4) development of nectar pockets up to the throat turning the tube into an appendicular hypanthium. The reddish-orange aril, associated with ornitochory, is plesiomorphic in Turneraceae, represented only in *Erblichia*; the other genera have white/whitish aril, associated with mirmecochory, except *Mathurina*, with an aril divided into filaments as an adaptation to anemochory.

**Keywords** Adnation · Africa · America · Connation · Floral morphology · Seed micromorphology

## Introduction

The family Turneraceae holds 226 species and 12 genera. *Adenoa* Arbo (1 sp.), *Erblichia* Seeman (1 sp.) and *Piriqueta* Aubl. (45 spp.) occur in the Americas, while *Afroqueta* Thulin & Razafim. (1 sp.), *Hyalocalyx* Rolfe (1 sp.), *Loewia* Urb. (1 sp.), *Stapfiella* Gilg (6 spp.), *Streptopetalum* Hochst. (6 spp.) and *Tricliceras* Thonn. ex DC. (16 spp.) occur in Africa. *Arboa* Thulin & Razafim. (4 spp.) is endemic to Madagascar, and *Mathurina* Balf. f. (1 sp.) is endemic to the Mascarene Islands. *Turnera* L. has 141 native species in the Americas and two in Africa, which are at present arranged into 11 series, hereafter abbreviated as “ser.” (Table 1). Turneraceae is closely related to Passifloraceae and Malesherbiaceae, and the three of them are treated together as Passifloraceae s.l. in the APG III (2009). Sequences from 25 genera and 42 species of Passifloraceae s.l. were analyzed phylogenetically by Tokuoka (2012). According to his results, the monophyly of Passifloraceae s.l. and that of Turneraceae, Malesherbiaceae and

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**Table 1** Comparison of molecular and morphological analyses

Position of genera and ser. of <i>Turnera</i>	Molecular trees		Morphological trees
	Chafé (2009) 9 genera, 68 spp.	Thulin et al. (2012) 12 genera, 29 spp.	Arbo and Espert (2009) 3 genera, 95 spp.
Genera from continental Africa	Monophyletic: <i>Afroqueta</i> , <i>Stapfiella</i> , <i>Streptopetalum</i> , <i>Tricliceras</i> , <i>Hyalocalyx</i> ; basal position	Monophyletic: <i>Afroqueta</i> , <i>Stapfiella</i> , <i>Streptopetalum</i> , <i>Tricliceras</i> , <i>Hyalocalyx</i> , <i>Loewia</i> ; derived position	Not resolved, <i>Stapfiella</i> , <i>Hyalocalyx</i> and <i>Tricliceras</i> not associated; <i>Loewia</i> , <i>Afroqueta</i> , <i>Streptopetalum</i> in one clade sister to American genera
<i>Arboa</i> (4 spp.)	Not treated	3 spp., associated with <i>Mathurina</i>	2 spp., not associated with other genera
<i>Adenoa</i> (1 sp.)	Associated with <i>Erblichia</i> and <i>Mathurina</i>	Associated with <i>Piriqueta</i> and <i>Turnera</i>	Associated with <i>Piriqueta</i> , sister to <i>Turnera</i>
<i>Erblichia</i> (1 sp.)	Associated with <i>Adenoa</i> and <i>Mathurina</i>	Sister to all African genera	Basal position, sister to all other genera
<i>Mathurina</i> (1 sp.)	Associated with <i>Adenoa</i> and <i>Erblichia</i>	Associated with <i>Arboa</i>	Not associated with other genera
<i>Piriqueta</i> (45 spp.)	24 spp., monophyletic, sister to <i>Turnera</i>	4 spp., monophyletic, sister to <i>Turnera</i>	12 spp., monophyletic, sister to <i>Adenoa</i>
<i>Turnera</i> (143 spp.)	35 species, monophyletic, sister to <i>Piriqueta</i>	6 spp., monophyletic, sister to <i>Piriqueta</i>	73 spp., monophyletic, sister to <i>Adenoa</i> and <i>Piriqueta</i>
Ser. <i>Annulares</i> (4 spp.)	Not treated	Not treated	Monophyletic (4 spp.), nested in clade I with ser. <i>Capitatae</i> , <i>Stenodictyae</i> and <i>Salicifoliae</i>
Ser. <i>Anomala</i> e (14 spp.)	2 spp., associated with <i>T. calyptrocarpa</i> (ser. <i>Microphyllae</i> )	Not treated	7 spp., nested with ser. <i>Turnera</i> ; sister to ser. <i>Conciliatae</i> group in clade IV
Ser. <i>Capitatae</i> (10 spp.)	2 spp., in a clade sister to all other species of <i>Turnera</i>	1 sp. sister to the other 5 spp. of <i>Turnera</i>	6 spp., nested with ser. <i>Annulares</i> , <i>Salicifoliae</i> and <i>Stenodictyae</i> in clade I
Ser. <i>Conciliatae</i> (1 sp.)	Not treated	Not treated	Associated with <i>T. calyptrocarpa</i> + <i>T. hebeptetala</i> (ser. <i>Microphyllae</i> ); sister to ser. <i>Turnera</i> + <i>Anomala</i> e in clade IV
Ser. <i>Letocarpae</i> (56 spp.)	Not resolved, <i>T. sideoides</i> separated; 5 spp. in one clade sister to ser. <i>Turnera</i>	1 sp. sister to ser. <i>Turnera</i>	20 spp., resolved, including <i>T. sideoides</i> , with ser. <i>Sessilifoliae</i> nested in clade IV; sister to clade III
Ser. <i>Microphyllae</i> (5 spp.)	Not resolved	Not treated	Not resolved
	1 sp. associated with 1 sp. of ser. <i>Papilliferae</i>	1 sp. associated with 1 sp. of ser. <i>Papilliferae</i> (2 spp.)	2 spp., associated with ser. <i>Papilliferae</i> (2 spp.) in clade I
	1 sp. associated with 2 spp. of ser. <i>Anomala</i> e	1 sp. associated with 2 spp. of ser. <i>Anomala</i> e	2 spp., sister to ser. <i>Conciliatae</i> and ser. <i>Turnera</i> + <i>Anomala</i> e in clade IV
Ser. <i>Papilliferae</i> (2 spp.)	1 sp. associated with 1 sp. of ser. <i>Microphyllae</i>	Not treated	2 spp., associated with 2 spp. of ser. <i>Microphyllae</i> in clade II
Ser. <i>Salicifoliae</i> (12 spp.)	3 spp., not resolved	Not treated	8 spp., nested with ser. <i>Annulares</i> , <i>Capitatae</i> , <i>Stenodictyae</i> in clade I

Table 1 continued

Position of genera and ser. of <i>Turnera</i>	Molecular trees		Morphological trees	
	Chafe (2009) 9 genera, 68 spp.	Thulin et al. (2012) 12 genera, 29 spp.	Arbo and Espert (2009) 3 genera, 95 spp.	This study, 12 genera, 100 spp.
Ser. <i>Sessilifoliae</i> (2 spp.)	Not treated	Not treated	1 sp. associated with 1 sp. of ser. <i>Leiocarpae</i>	Nested with ser. <i>Leiocarpae</i> in clade III
Ser. <i>Stenodictyae</i> (10 spp.)	Not treated	Not treated	5 spp., monophyletic, associated with ser. <i>Capitatae</i> and <i>Salicifoliae</i>	4 spp., nested with ser. <i>Annulares</i> , <i>Capitatae</i> and <i>Salicifoliae</i> in clade I
Ser. <i>Turnera</i> (27 spp.)	19 spp., monophyletic	4 spp., monophyletic	19 spp., associated with ser. <i>Anomalaе</i>	15 spp., associated with ser. <i>Anomalaе</i> ; sister to ser. <i>Conciliatae</i> group in clade IV

Passifloraceae sensu stricto, are strongly supported. Urban's monography (1883) and the revisions of the neotropical genera (Arbo 1977, 1979, 1995, 1997, 2000, 2005, 2008) highlighted the relevance of seed characters for taxonomy. In these studies, the authors described seed shape, size, color, curvature, type of coat (episperm), as well as the degree of chalaza development and orientation, shape of the exostome, and relative length and width of the aril. Some details were also considered, such as the presence of outstanding knots on the seed coat reticulate and of punctiform cavities in each areole, and the type of cells forming the aril.

The first molecular phylogenetic study of Turneraceae included 5 species of *Piriqueta* and 35 American species belonging to seven series of *Turnera* (Truyens et al. 2005). According to their results, *Turnera* was monophyletic, and series *Turnera* was the only monophyletic series. Afterwards, cladistic analyses of the genus *Turnera* (92 species) based on morphological characters and chromosome numbers were made to test the monophyly of the series, and to assess biogeographic patterns (Arbo and Espert 2009).

Chafe (2009) conducted further analyses including all the species of *Turnera* sequenced in Truyens et al. (2005), *Adenoa*, *Erblichia*, *Mathurina*, some additional species of *Turnera*, many other species of *Piriqueta* and species of four African genera. His results varied depending on the method used to analyze the data (parsimony, maximum likelihood, or Bayesian inference). Nevertheless, some clades as *Piriqueta* and *Turnera* ser. *Turnera* were consistent across methods, as well as the isolated position of *Piriqueta capensis*, which is currently treated as the monotypic genus *Afroqueta* (Thulin et al. 2012). The molecular phylogenetic tree that so far includes the largest number of species of Turneraceae is illustrated in Chafe's thesis (2009, Fig. 11): 68 species of 9 genera, missing only the African genera *Arboa*, *Hyalocalyx* and *Loewia*. The tree comprises 24 species of *Piriqueta* and 35 of *Turnera*, corresponding to seven series. The African genera *Afroqueta*, *Stapfiella*, *Streptopetalum* and *Tricliceras* are found in a clade sister to all other Turneraceae. *Adenoa*, *Mathurina* and *Erblichia* appear closely related, and *Turnera* and *Piriqueta* are sister clades.

The most recent molecular phylogenetic study (Thulin et al. 2012) includes 29 species representing all the genera. Their Bayesian majority rule consensus tree shows two well-supported clades structured geographically: the basal one with the American taxa and the two African *Turnera* species, and the other with the African genera and the American *Erblichia odorata*. Unfortunately, this study, the only one including all the genera of the family, sampled very few species of *Turnera*.

Evolution is intimately linked to changes in chromosome number and karyotype. The ancestral base number in

Turneraceae might be  $x = 7$ , found in *Piriqueta*, *Turnera* and *Adenoa* (Fernández 1987; Gonzalez et al. 2012). However, chromosome numbers are unknown for the African genera. The genus *Turnera* including 66 % of the species of the family has three base chromosome numbers:  $x = 7$ ,  $x = 5$  and  $x = 13$  (Fernández 1987), and polyploidy has played an important role in speciation, given that several species, including *T. ulmifolia* L. (type of the genus) are allopolyploids (Shore et al. 2006). At present, *Turnera* is divided into 11 series, but critical information to change taxonomic circumscriptions is still missing, since the base number of ser. *Annulares*, ser. *Capitatae*, ser. *Conciliatae* and ser. *Sessilifoliae* is unknown. Finally, none of the species of four series: ser. *Annulares*, ser. *Conciliatae*, ser. *Sessilifoliae* and ser. *Stenodictyae* has yet been included in a molecular phylogenetic analysis.

The present research focused on the micromorphology of seeds in all genera of Turneraceae, as well as the comparison of floral and fruit traits important in the delimitation of genera. The goals of our study were: (1) to assess the impact of morphological attributes and seed micromorphology in the cladistic analyses of the family; (2) to compare our results with those based on molecular datasets.

## Materials and methods

This study includes 101 taxa representing all twelve genera in the family Turneraceae, and all the series of *Turnera*. The species and specimens analyzed with collector, number, source, and herbaria are listed in Online Resource 1. All known allopolyploid species were excluded, as in Arbo and Espert (2009), because they are the products of reticulate evolution. A species of *Passiflora* was used as outgroup. Seeds were gold/palladium coated for the scanning electron microscope (SEM) analysis. We used the Jeol 5800 LV SEM of the electron microscopy service of the ‘Universidad Nacional del Nordeste’, Corrientes. Some photographs were taken at Washington University, St. Louis, USA and York University, Toronto, Canada. Seeds of *Erblichia* were microtome sectioned to verify the nature of the endotesta, following techniques described in Gonzalez et al. (2012) and Gonzalez and Arbo (2013).

The list of characters and character states is provided in Online Resource 2. Each time a character is mentioned in the text, the corresponding number is shown between square brackets. We considered all morphological characters used by Arbo and Espert (2009) in *Turnera* and *Piriqueta* and base chromosome numbers. To cover generic variability, it was necessary to add states in some characters. Variable characters at generic level [19, 27, 29, 33, 35,

36, 39, 41, 44, 45] and 34 features resulting from the micromorphological analysis of seed were included. In the case of continuous characters, three ranges (minimum–medium–maximum) were settled and the average was considered for each species.

Areoles were measured at the region of maximum seed diameter, but given the variable areole size in individual seeds, only the areoles of largest size, corresponding to the antiraphe, were measured and the average was calculated. Whenever possible, SEM photographs were used for character coding; they were supplemented with observations at the maximum magnification (40×) of a stereoscopic microscope Leica Wild M3C equipped with double incident light (lateral and vertical).

The morphological matrix was analyzed using the maximum parsimony criterion implemented in the software TNT ver. 1.1 (Goloboff et al. 2008a). The states of the characters [18, 19, 25, 26, 90] related with the evolution of the floral tube, present in the majority of the genera, were ordered according to its development. The missing characters were coded with a question mark, and the inapplicable ones with a hyphen. The data matrix is available upon request to the corresponding author.

Analyses considering all characters with equal weights detected high levels of homoplasy. Consequently, analyses with different implied weighting schemes were carried out. Different characters do not provide equally strong evidence (some display high levels of homoplasy while others are perfectly hierarchical). Therefore, trees obtained from properly weighted characters are desirable (Goloboff 1993). Moreover, down-weighting morphological characters according to their homoplasy produce more strongly supported groups and more stable results (Goloboff et al. 2008b). The implied weighting method consists of estimating character weights according to their homoplasy. The “character fit”, as defined by Goloboff (1993), is a concave function of the character’s homoplasy (i.e., number of extra steps), and trees with maximum total fit are searched for. The “fittest” tree implies that the characters are maximally reliable and, given character conflicts, they are solved in favor of those characters that have less homoplasy. The degree of the concave function can be modified in TNT through the constant  $k$ , where lower  $k$  values penalize homoplastic characters harder than higher values of  $k$ . When using implied weights, it is not clear how much a homoplastic character should be down-weighted; therefore, we explored different values of  $k$  and chose 2–4, because with these values, homoplasy was hardly penalized and very similar trees were retrieved. The searching procedure involved a driven search finding the minimum length 10 times with default settings for sectorial searches and tree fusing (Goloboff 1999). The resulting trees were

additionally TBR swapped. Bremer support (Bs; Bremer 1994) and Jackknife (JK; Farris et al. 1996) were used to estimate branch support. Bs was estimated on the basis of 10,000 suboptimal trees of 0.01–0.5 less fit and for each value of  $k$ . For the JK analyses, the matrix was resampled 1000 times, with a probability of character removal  $p = 0.36$ . Values were expressed as absolute frequencies.

Morphological characters, including homoplastic ones, were optimized on one of the most parsimonious trees randomly selected from the analysis. The homoplastic characters were optimized because they define groups within the family or represent diagnostic features for *Turnera*.

## Results

The numbers shown between square brackets indicate the corresponding number of each character listed in Online Resource 2.

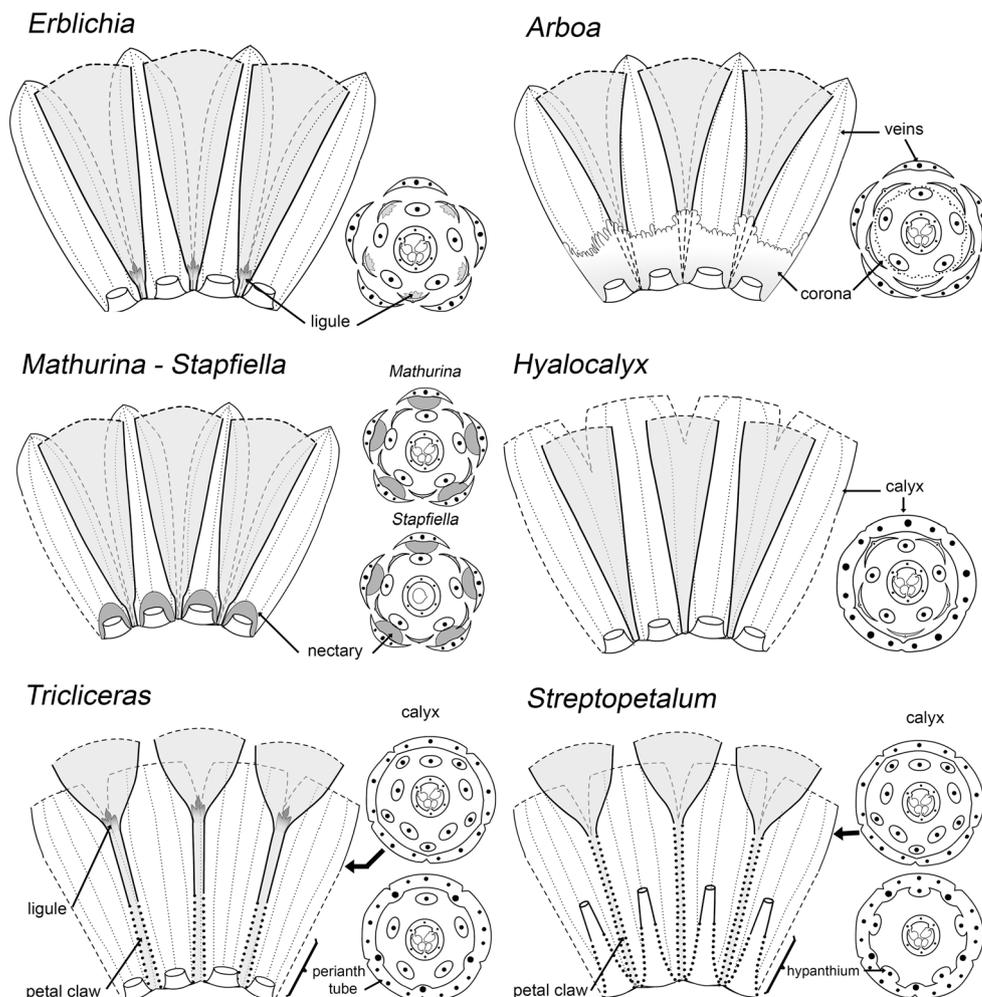
## Adnation and connation of floral whorls

Within Turneraceae, sepals, petals and staminal filaments show noteworthy variation in connation (merging of pieces of the same verticil) and adnation (adherence, fusion of pieces from different whorls).

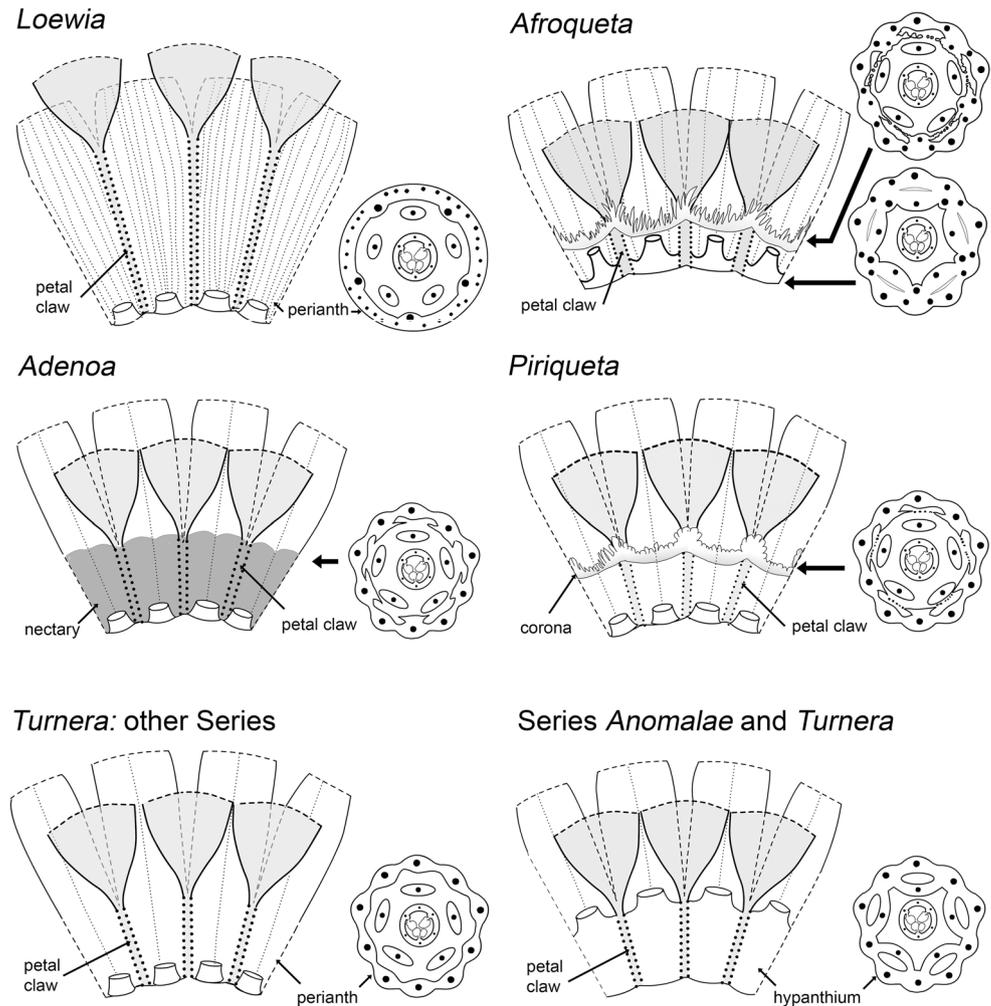
*Erblichia*, *Arboa*, *Mathurina* and *Stapfiella* (Fig. 1) have free or almost free sepals [18]. All the other genera possess a tube of variable nature, length of connation [18] and number of major veins [19]. *Hyalocalyx* (Fig. 1) is the only genus in which the tube is formed only by the calyx, because the petals are adnate [90] only at the base to the calyx tube; in the remaining genera, the petal claws are adnate to the calyx, developing a perianth tube.

In *Hyalocalyx*, *Tricliceras*, *Streptopetalum* (Fig. 1) and *Afroqueta* (Fig. 2), the tube has 15 veins [19], i.e., sepals connation does not involve the fusion of the sepals lateral veins. In *Tricliceras* (Fig. 1), the petal claws are adnate to the calyx tube along 1/3–1/2 of its length, while in *Streptopetalum* (Fig. 1) they are adherent to the calyx tube up to

**Fig. 1** Floral morphology in Turneraceae: *Erblichia*, *Arboa*, *Mathurina* and *Stapfiella* with free sepals, *Hyalocalyx* with calyx tube, *Tricliceras* and *Streptopetalum* with perianth tube. On the left, part of the calyx inner face with basal portions of petals and staminal filaments; the adnate portions are indicated with thick dotted lines. On the right, cross section of flower showing the relationships between whorls; the principal veins are represented with dots



**Fig. 2** Floral morphology in Turneraceae: *Loewia*, *Afroqueta*, *Adenoa*, *Piriqueta* and *Turnera* with perianth tube, *Turnera* ser. *Anomalae* and ser. *Turnera* with hypanthium. On the left, part of the calyx inner face with basal portions of petals and staminal filaments; the adnate portions are indicated with thick dotted lines. On the right, cross section of flower showing the relationships between whorls; the principal veins are represented with dots



the throat [90]; this perianth tube is 6–18 mm long. The same happens in *Afroqueta* (Fig. 2), but the perianth tube is only 0.5–3 mm long [18].

In *Loewia* (Fig. 2), the calyx tube does not show major veins, according to Urban (1896) there are 35–40 delicate veins [19], and the petals' claws are adnate all along the calyx tube [90].

Connation and adnation are deeper in *Adenoa*, *Piriqueta* and *Turnera* (Fig. 2), because the perianth tube has only 10 veins [19], 5 of which represent the fusion of the lateral veins of the sepals with the vein along the claw of each petal (Gonzalez 2001; Gonzalez et al. 2012).

Staminal filaments of *Erblichia* (Fig. 1) are adnate at the base to a sepal and are free from each other [25, 26, 27]. In *Mathurina* and *Stapfiella*, there are conspicuous nectaries at the insertion of the staminal filaments on the sepals. In *Arboa*, the staminal filaments are not adnate to the sepals, because there is an annular free corona between perianth and androecium [23].

In *Hyalocalyx* (Fig. 1), the staminal filaments are adnate to the calyx tube only at the base, on the external face [26, 27] and are free from one another [25]. In *Triliceras* (Fig. 1), *Loewia*, *Adenoa*, *Piriqueta*, and many species of *Turnera* (Fig. 2), they are adnate to the perianth tube in the same way. The stamens of *Streptopetalum* (Fig. 1) are attached to the perianth tube 20–33 % of its length [26]; therefore, the basal portion of the tube is in fact an appendicular hypanthium.

The staminal filaments of *Adenoa* (Fig. 1) and a few species of *Turnera* are almost free, barely adnate to the perianth tube at the base, but in *T. calyptrocarpa* and *T. hebetata* (ser. *Microphyllae*) they are also briefly connate [26], developing a slender annular structure 0.25–0.40 mm long (Arbo 2000; Gonzalez 2001).

*Turnera rubrobracteata* (ser. *Conciliatae*) and *T. reginae* (ser. *Anomalae*) display staminal filaments connate at different heights above adnation to calyx or perianth [26] (Arbo 2005, 2008).

In *Afroqueta* (Fig. 2), the margins of the staminal filaments are shortly adhering (0.4 mm) to the petal claws and connate at the base [25, 26, 27]. In the species of *Turnera* ser. *Anomalae* and ser. *Turnera* (Fig. 2), the staminal filaments show marginal adnation to petal claws and conation usually up to the throat of the flower, developing nectar pockets [28] and defining an appendicular hypanthium (Arbo 1986, 2005; Gonzalez 2001).

Perianth appendices (coronas, ligules, emergences) [23]

The only genus of Turneraceae with a free corona, inserted between perianth and staminal filaments, is *Arboa* (Fig. 1). *Afroqueta* and *Piriqueta* (Fig. 2) have an annular corona, laciniate, fixed at the perianth throat, on the base of the petal blades and on the sepals. In one brevistylous flower of *Turnera reginae*, a streak of filiform appendices was found at the same location (Arbo 2005). The petals of *Erblichia* and *Tricliceras* (Fig. 1) have a small ligule, which can also be found in a few species of *Turnera* ser. *Capitatae* and ser. *Salicifoliae* (Arbo 1997, 2000; Gonzalez 2001).

Placentation

The Turneraceae have a tricarpeolar unilocular ovary, usually with a few to many ovules arranged in parietal placentation. The only exception is the genus *Stapfiella* (Figs. 1, 3c), in which each ovary has only one ovule and basal placentation [35].

In most genera, each placenta is linear. In *Turnera* and *Piriqueta*, the placental bundles often merge into one bundle that runs parallel to a sutural bundle originated by the fusion of the marginal bundles of carpels (Gonzalez 2001). The length of the placenta is variable and in relation to the number of ovules developed. In pauciovulate ovaries, the placenta is short, less than half of the ovary length.

The ovules are usually inserted along a slender stripe on the placental bundle. In mature fruits, the placental bundle appears as a prominent line along the middle vein on the inner face of each valve (Fig. 3d, e). In a few genera: *Erblichia*, *Mathurina*, *Arboa* and *Adenaea*, the placenta is wide, the ovules are inserted along many rows [36]. The funicles or the scars of their insertion can be observed on the inner face of the dehiscent fruit valves (Fig. 3b).

Size, shape, and parts of the seed

Turneraceae seeds are small, only a few species have seeds that are more than 4 mm long [52]. The largest seeds are found in *Erblichia* (4–5 mm) and *Stapfiella* (3–4.6 mm), while the smallest are found in *Hyalocalyx* (1.5–1.6 mm). In *Turnera*, seeds vary across the whole size range: large seeds occur in *T. glaziovii* (4.9 mm), and small ones in *T.*

*argentea*, *T. diamantina* Arbo and *T. curassavica* Urb. (1.2–1.6 mm). Ripe seeds are usually very dark, almost black (Fig. 3c).

The seeds of Turneraceae develop from anatropous ovules, with the hilum and the micropyle located at the base, the chalaza at the apex, and the linear raphe along one side (Gonzalez and Arbo 2013). In some cases, a cellular proliferation [50] can be observed at the chalazal end of the raphe. The proliferation is notable for its dark color in immature seeds with light brown surface. This contrast disappears in ripe seeds, which become completely dark. The proliferation is found in seeds of *Mathurina* (Fig. 4d) and various species of *Turnera* and *Piriqueta* (Gonzalez and Arbo 2013).

Seeds are straight [46, 48] in *Mathurina* (Fig. 4d), *Stapfiella* (Fig. 4e, f) and *Tricliceras* (Fig. 5b). They are curved in other genera, such as *Adenaea* (Fig. 4a), *Hyalocalyx* (Fig. 5a), *Afroqueta* (Fig. 5d), *Streptopetalum* (Fig. 5e), and many species of *Piriqueta* and *Turnera*; when the seed is curved, the raphe is located at the concavity (Fig. 5a, d–f). In rapheal view [47, 49], they are usually obovoid and slendering towards the exostome (Fig. 5c). When the seed is markedly curved, as in *Hyalocalyx*, the rapheal view allows a frontal sight of the exostome (Fig. 6a).

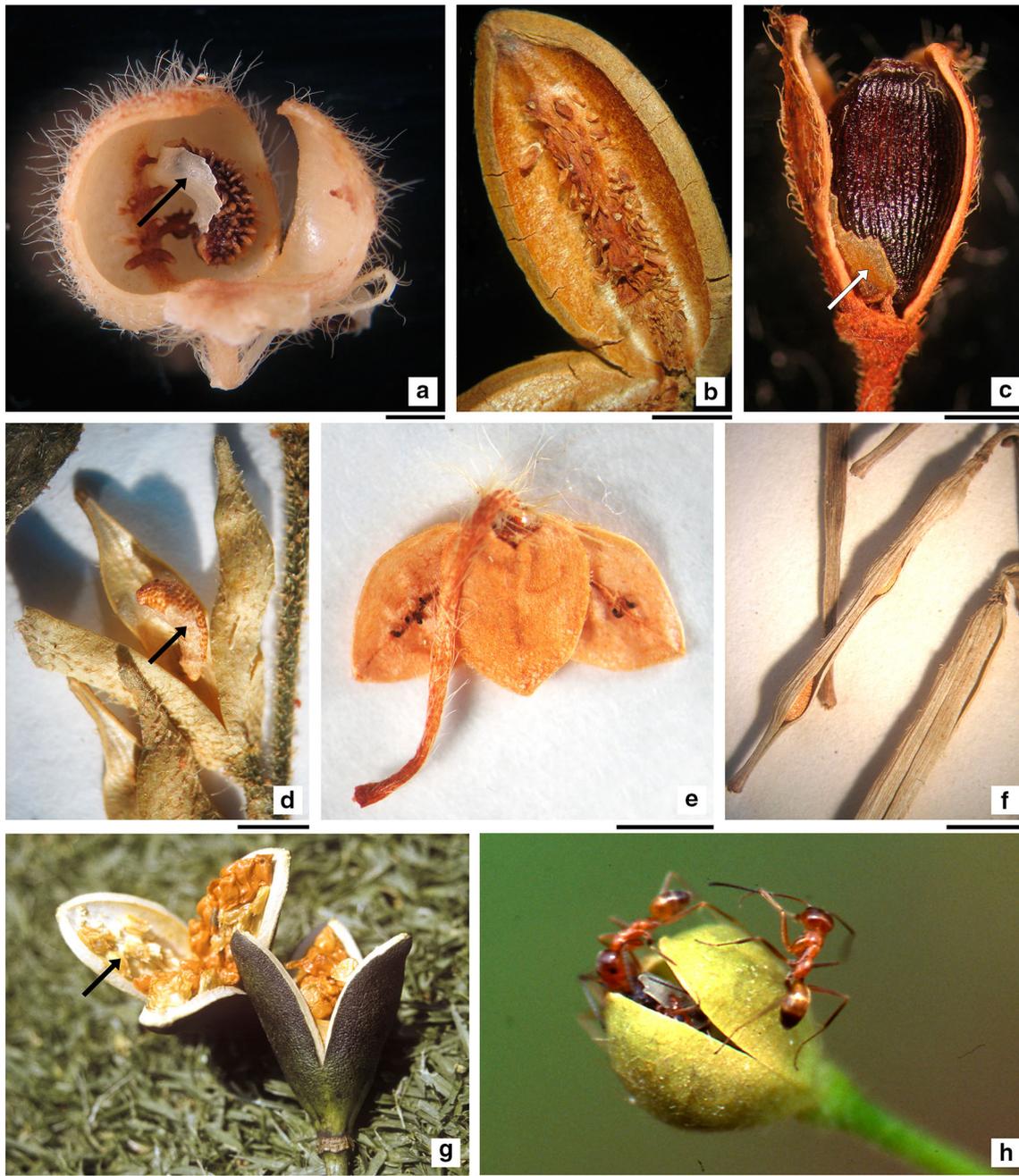
The exostome [72–75] is conical in *Arboa*, *Erblichia* (Fig. 4c), *Stapfiella lucida* (Fig. 4e) and *Mathurina* (Fig. 8f), hemisphere shaped in *Adenaea*, *Hyalocalyx*, *Tricliceras*, *Piriqueta* and species of *Turnera* (Fig. 6b, d), and it is parrot-beak shaped in *Streptopetalum serratum* (Fig. 6e) and *Afroqueta*. The exostome is usually shorter in length than in diameter, but it is longer in *Stapfiella lucida* (Fig. 6c) and *Streptopetalum serratum* (Fig. 5e).

The upper edge of the exostome is marked by a rim (Figs. 5b, e; 6a, b) in all genera except for *Arboa*, *Erblichia* (Fig. 4c), *Mathurina* (Fig. 8f) and *Stapfiella* (Fig. 6c). In *Turnera sidoides*, the rim develops into an annular crest (Gonzalez and Arbo 2013). The ratio rim/exostome diameter is variable [75], sometimes the exostome is bulky (Fig. 6d), but when it is small, its diameter may correspond to just 1/3 of the rim (Fig. 6a).

The hilum [51] is usually located on the rapheal side next to the exostome rim (Fig. 6b), but in *Adenaea*, *Arboa*, *Erblichia* (Fig. 4a–c) and *Mathurina* (Fig. 8f), it is properly on the exostome, close to the base.

The chalaza [76] is obtuse in *Adenaea*, *Arboa*, *Erblichia*, *Mathurina* (Fig. 4a–d), *Hyalocalyx* (Fig. 5a), and *Piriqueta*. It is prominent in *Tricliceras* (Fig. 5b) with the surface markedly concave in *Loewia*, *Afroqueta* (Fig. 5c–d), *Stapfiella* (Fig. 4e) and *Streptopetalum*. *Turnera* shows all the range of variation (Gonzalez and Arbo 2013).

Sometimes, when the chalaza is prominent [77–80], an outcropping develops with the same sculpturing of the seed



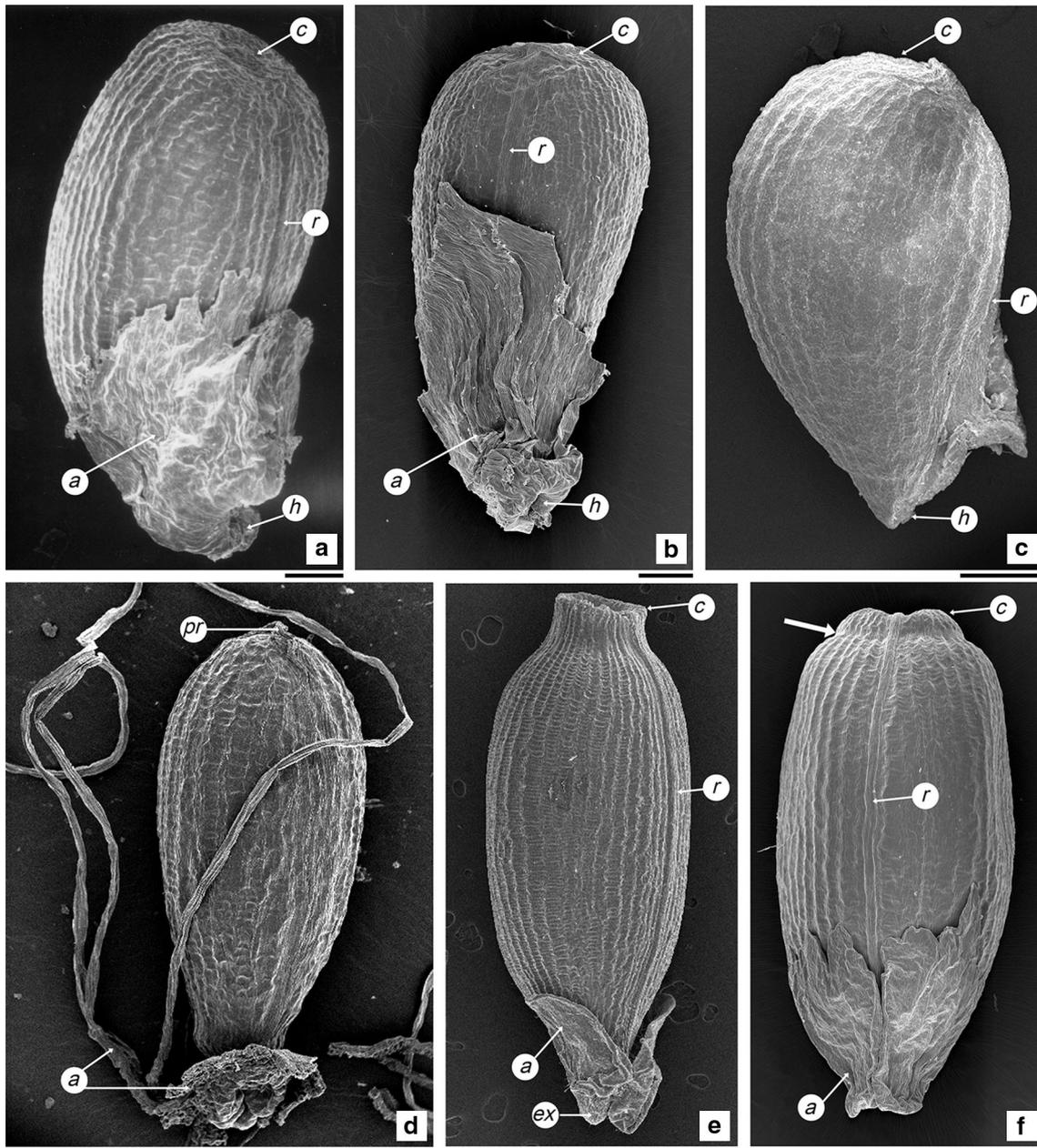
**Fig. 3** Dehiscent fruits of Turneraceae: **a** *Turnera sidoides* subsp. *carnea*, almost ripe fruit, valve inner face showing insertion of a crested seed (Solís Neffa et al. 271); **b** *Erblichia odorata*, valve inner face, funicles and scars arranged in several rows (Jiménez et al. 1531); **c** *Stapfiella ulugurica*, ripe fruit without a valve, the only seed inserted at the base (Mlangwa et al. 1548); **d** *Streptopetalum serratum*

(Dinter 7530); **e** *Hyalocalyx setiferus*, inverted fruit (Schlieben 6338); **f** *Tricliceras longipedunculatum*, siliqua-shaped capsule (Schlieben 9283); **g** *Erblichia odorata*, seeds covered with orange-red aril (photo A. García Mendoza); **h** *Piriqueta cistoides* subsp. *caroliniana*, ants collecting seeds (photo T. Feldman). Scale bars **a** = 2.5 mm, **b**, **f**, **g** = 6 mm, **c** = 1 mm, **d**, **e**, **h** = 2 mm; arrow indicates aril

(Fig. 4e). The base of the chalaza is occasionally marked by a slight constriction (Fig. 4f, arrow). The orientation of the chalaza surface is variable [80], towards the apical pole (Fig. 5b), intermediate (Fig. 5c–d), or towards the raphe (Fig. 5f).

#### Micromorphology of seed coat (episperm)

Considering structure and sculpturing, there are two basic types of seed coat: reticulate and crested [55–56]. The reticulate seed coat and its variants, striate–reticulate and



**Fig. 4** Seeds of Turneraceae with striate seed coat: **a**, **c–e** lateral view, **b**, **f** rapheal view; **a** *Adenoa cubensis* (Acuña 12577); **b** *Arboa antsingyae* (Leandri 2173); **c** *Erblichia odorata*, aril insertion along the raphe (Mc Pherson 9999); **d** *Mathurina penduliflora* (Friedman 2444); **e** *Stapfiella lucida* with an outcropping chalaza (Lewalle

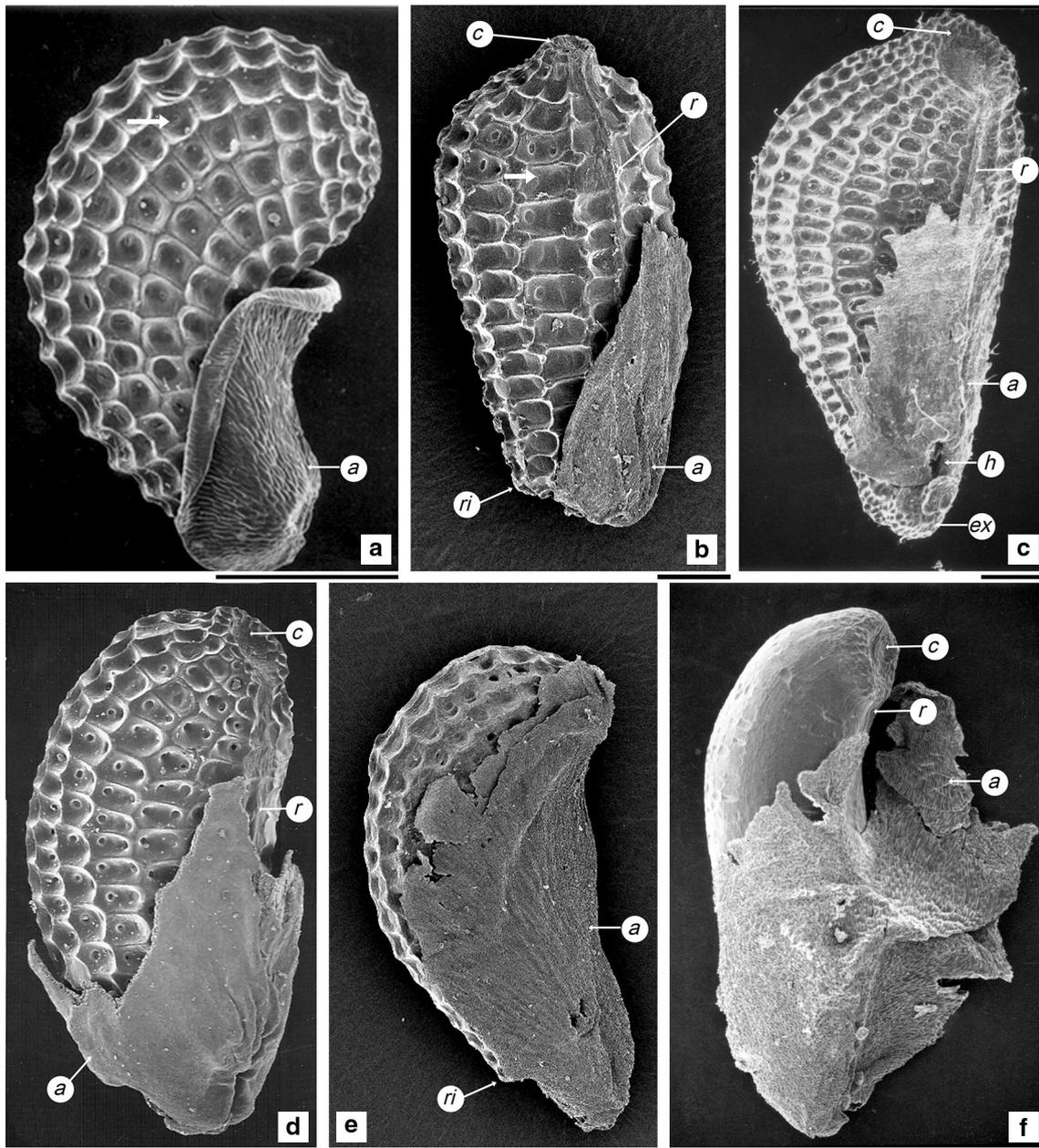
3494); **f** *Stapfiella ulugurica* with a slight constriction at the base of the chalaza (arrow) (Mlangwa et al. 1548). Scale bars **a**, **b**, **d–f** = 0.5 mm, **c** = 1 mm. *a* aril, *c* chalaza, *ex* exostome, *h* hilum, *pr* proliferation, *r* raphe

striate, are due to the interaction of endotesta and exotegmen; the crested seed coat is exclusive to *Turnera sidoides* (ser. *Leiocarpae*, Fig. 3a), and it is mostly originated by the exotesta (Gonzalez and Arbo 2013).

The reticulate seed coat (Fig. 5) is present in *Afroqueta*, *Hyalocalyx*, *Loewia*, *Piriqueta*, *Streptopetalum*, *Tricliceras* and most species of *Turnera* (ser. *Annulares*, ser. *Capitatae*, ser. *Conciliatae*, ser. *Leiocarpae*, ser. *Microphyllae*,

ser. *Papilliferae*, ser. *Sessilifoliae*, and ser. *Turnera*). The reticulate knots [59] are sometimes prominent; they are acute in *Piriqueta racemosa*, and rounded in *Tricliceras* (Figs. 5b, 6f) and some species of *Turnera* ser. *Leiocarpae* (Gonzalez and Arbo 2013).

The areoles [60] may be outstanding (Fig. 5d) or hardly perceptible (Fig. 5f). They are usually transverse-rectangular (Fig. 5b, arrow) or square, sometimes penta-

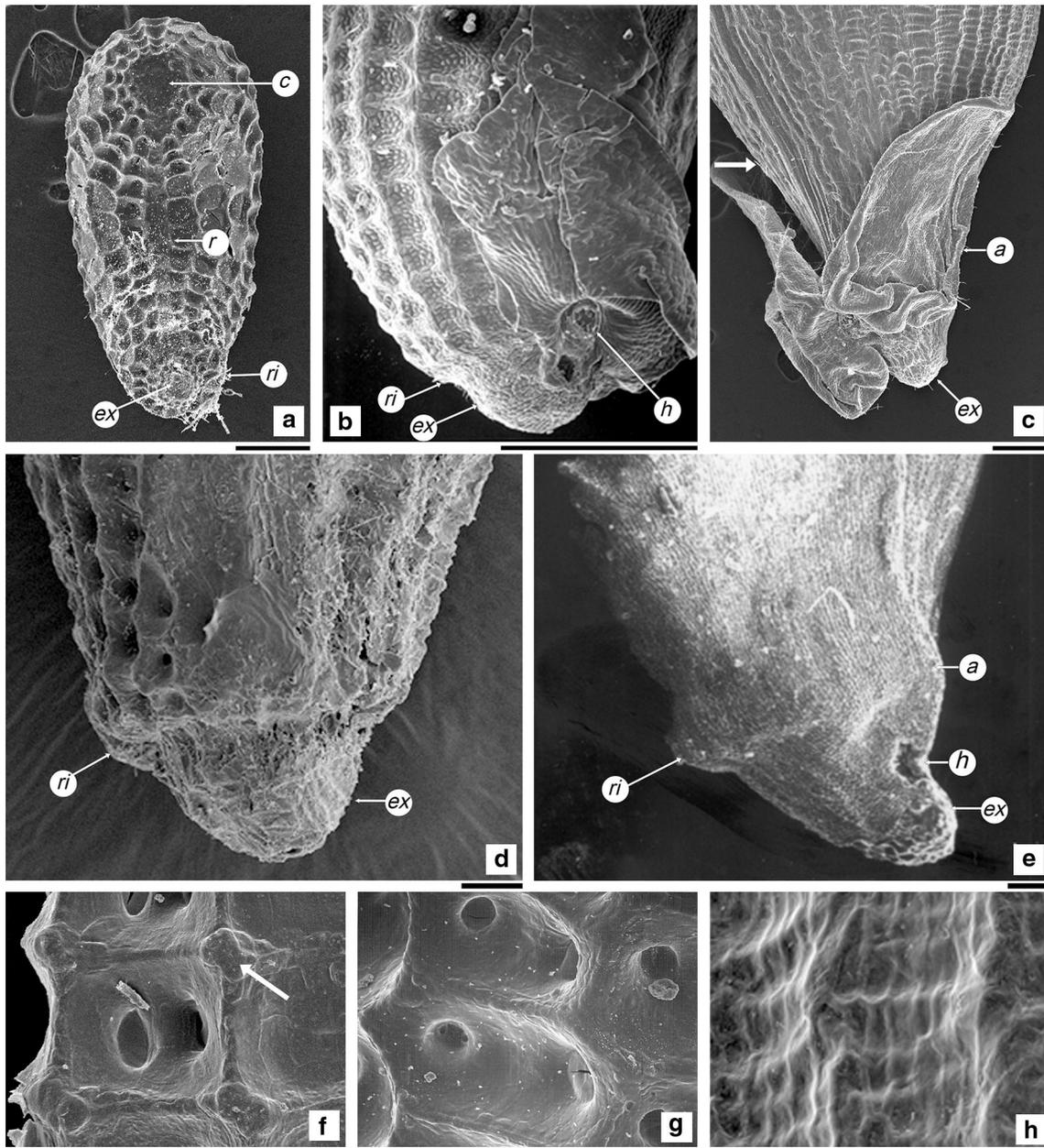


**Fig. 5** Seeds of Turneraceae with reticulate seed coat: **a, d–f** lateral view, **b, c** rapheal view. **a** *Hyalocalyx setiferus* with some pentagonal areoles (arrow) (Decary 8041); **b** *Tricliceras pilosum*, rectangular areoles (arrow) (Dinklage 8); **c** *Loewia glutinosa* with concave chalaza (Gillett and Newbould 19170); **d** *Afroqueta capensis* (Wall

sn); **e** *Streptopetalum serratum* exostome upper edge with a rim (Stannard and Gilbert 1107); **f** *Turnera hermannioides* (Harley et al. 15618). Scale bars **a–f** = 0.5 mm. *a* aril, *c* chalaza, *ex* exostome, *h* hilum, *r* raphe, *ri* rim

hexagonal (Fig. 5a, arrow). Areoles shape and depth are set by the endotesta cells (Gonzalez and Arbo 2013). This stratum of the seed coat [57] consists of only one layer of cells in *Adenoa* (Gonzalez et al. 2012), *Arboa* (Fig. 7a), *Hyalocalyx* (Fig. 7b), *Loewia*, *Stapfiella*, *Streptopetalum* (Fig. 7c), *Tricliceras* (Fig. 7d), *Piriqueta* and *Turnera* (Gonzalez and Arbo 2013); while in *Erblichia* and *Mathurina* (Fig. 7e, f), it has several layers.

The reticulate areoles [61] lacking punctiform cavities (just concaves) occur in *Hyalocalyx* (Fig. 7b) and species of *Piriqueta* and *Turnera*; areoles with one punctiform depression are exclusive to species of the American genera *Piriqueta* and *Turnera* (Gonzalez and Arbo 2013). The areoles with two punctiform cavities characterize the African genera *Loewia* (Fig. 5c), *Afroqueta* (Fig. 6g), *Streptopetalum* (Fig. 7c), and *Tricliceras* (Figs. 5b, 7d),



**Fig. 6** Seeds of Turneraceae: **a** *Hyalocalyx setiferus*, rapheal view, small exostome (Schlieben 6338); **b** *Turnera argentea*, aril insertion (Huber and Tillet 2809); **c** *Stapfiella lucida*, exostome upper edge without a rim (arrow) (Lewalle 3494); **d** *Turnera rubrobacteata*, bulky exostome (Kuhlmann 6656); **e** *Streptopetalum serratum*, exostome parrot-beak shaped (Dinter 7530); **f** *Tricliceras*

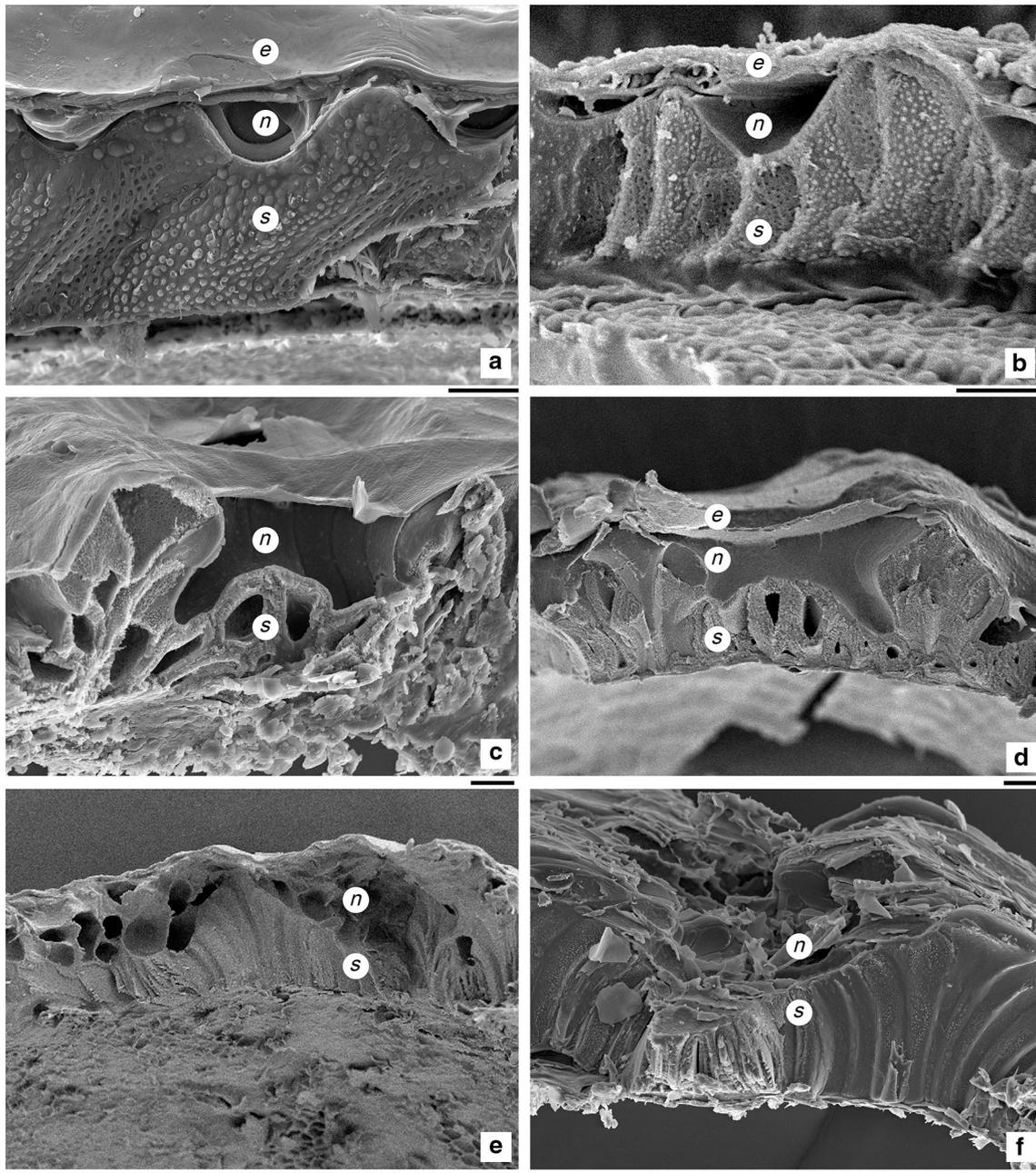
*longipedunculatum*, prominent reticulate knots (arrow) (Stuhlmann 957); **g** *Afroqueta capensis*, areoles with two punctiform cavities (Wall sn); **h** *Stapfiella ulugurica*, detail of striate seed coat (Mlangwa et al. 1548). Scale bars **a–c** = 0.25 mm, **d, e** = 0.1 mm, **f, g** = 50  $\mu$ m, **h** = 20  $\mu$ m. *a* aril, *c* chalaza, *ex* exostome, *h* hilum, *r* raphe, *ri* rim

even though they are also found in some shrubby species of *Piriqueta* (Arbo 1995; Gonzalez and Arbo 2013).

The size of the areoles [62–64] is variable in each seed, the maximum is found on the antiraphe side, in the region of largest seed diameter. The average maximum surface of the areoles ranges from 0.006 mm<sup>2</sup> in *Turnera stenophylla* to 0.1 mm<sup>2</sup> in *Tricliceras pilosum* (Willd.) R. Fern.

The striate–reticulate seed coat is present in most of the species of *Turnera* ser. *Anomalae*, ser. *Salicifoliae* and ser. *Stenodictyae* (Gonzalez and Arbo 2013).

The striate seed coat (Fig. 4), in which areoles can not be recognized clearly or measured on the seed surface, is present in *Adenoa*, *Arboa*, *Erblichia*, *Mathurina* and *Stapfiella*. The seed coat of *Adenoa* was described as



**Fig. 7** Seed coat sections: **a, b** concave endotesta cells; **c, d** endotesta cells with two punctiform cavities; **e, f** endotesta of several cell layers. **a** *Arboa antsingyae* (Leandri 2173); **b** *Hyalocalyx setiferus* (Schlieben 6338); **c** *Streptopetalum serratum* (Stannard and Gilbert 1107);

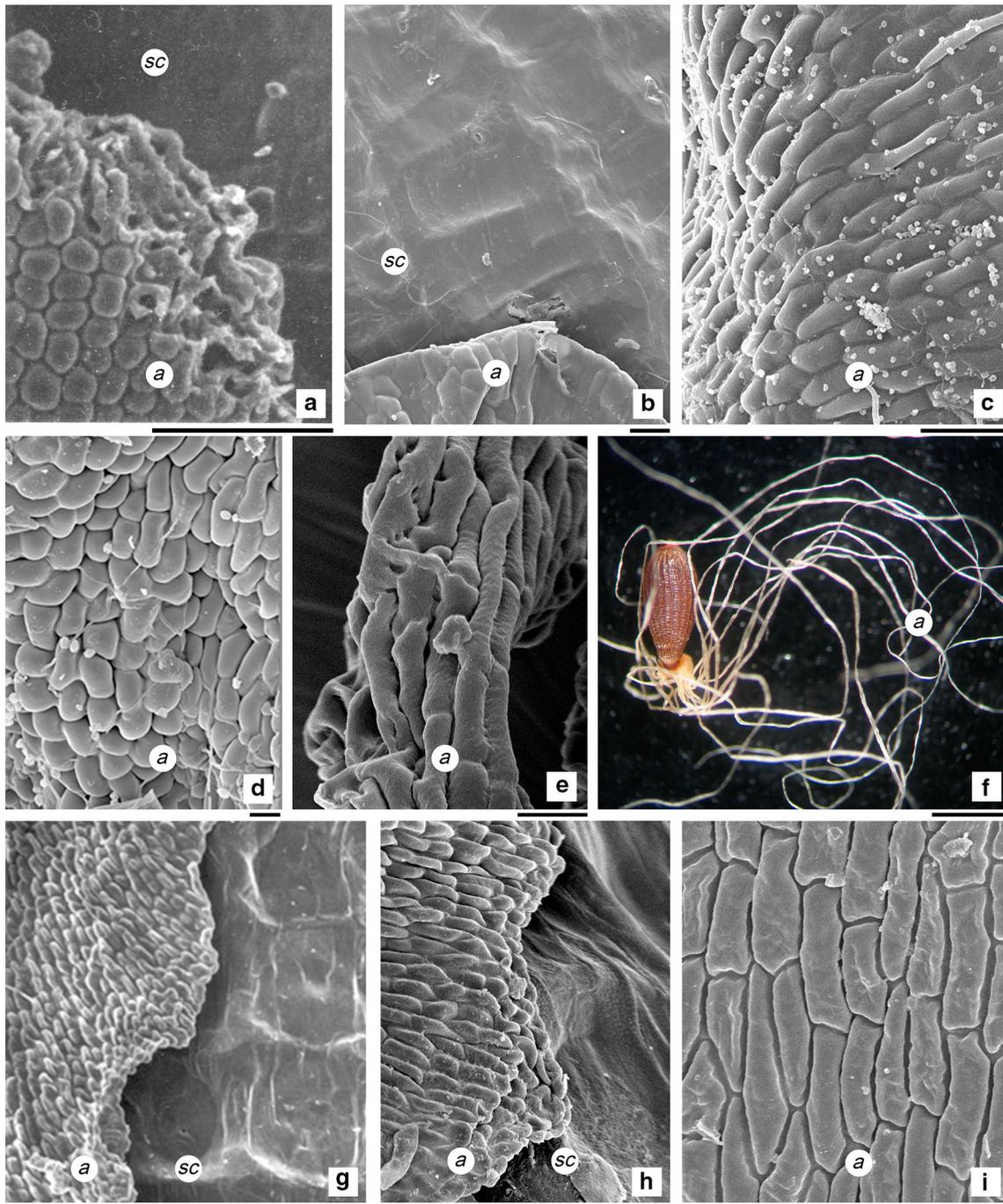
**d** *Triliceras longipedunculatum* (Stuhlmann 957); **e** *Erblichia odorata* (McPherson 9999); **f** *Mathurina penduliflora* (Friedman 2444). Scale bars **a–d, f** = 20  $\mu\text{m}$ , **e** = 0.2 mm. *e* epidermis (exotesta), *n* endotesta, *s* exotegmen sclereids

striate–reticulate (Gonzalez et al. 2012), but not even with SEM was it possible to measure the surface of the areoles (Fig. 8b).

The seed surface appearance is determined by the structure and variability of the epidermal cells [65–66]. The shape of epidermal cells is the same all over the seed sculpture except in some species of *Turnera* ser. *Leiocarpae*, where they have different size and shape: cells

located on the reticule ridges are large with smooth or convex outer wall, while cells located on the areoles are smaller and papillose (Fig. 6b). This group includes *Turnera sidoides*, which has a crested seed coat; epidermal cells are different in the crests and depressions (Gonzalez and Arbo 2013).

Commonly, the outer walls of the epidermal cells are flat (Fig. 6g) or convex, with smooth or striated cuticle.



**Fig. 8** Aril: **a, i** flat cells; **c, d** dome shaped cells; **g, h** papillose cells. **a** *Afroqueta capensis* (Wall sn); **b** *Adenoa cubensis* (Acuña 12577); **c** *Hyalocalyx setiferus* (Schlieben 6338); **d** *Loewia glutinosa* (Gillett and Newbould 19170); **e** *Mathurina penduliflora*, aril thread; **f** seed

and part of aril (Friedman 2444); **g, h** *Streptopetalum serratum* (Dinter 7530-S, Stannard and Gilbert 1107); **i** *Tricliceras longipedunculatum* (Stuhlmann 957). Scale bars **a–c, g** = 50  $\mu\text{m}$ , **d** = 10  $\mu\text{m}$ , **e, h, i** = 20  $\mu\text{m}$ , **f** = 1 mm. *a* aril, *sc* seed coat

However, in many species of *Turnera* and *Piriqueta*, the epidermal cells have papillae of different shapes [67–68]. In most species of *Piriqueta* there are finger-like papillae, while filiform papillae are found in species of *Turnera*, ser. *Salicifoliae*, ser. *Stenodictyae* and ser. *Anomala*e. Hemispheric

and mammiform papillae occur in species of *Turnera* ser. *Anomala*e, ser. *Annulares*, ser. *Capitatae*, ser. *Leiocarpae* (Fig. 6b) and ser. *Salicifoliae* (Gonzalez and Arbo 2013).

The presence of epicuticular wax [71] shaped as thin sticks (Gonzalez and Arbo 2013) is exclusive to several

species of *Turnera* ser. *Leiocarpae* including *Turnera si-doides*. It is mainly observed in immature seeds.

## Aril

All the seeds of Turneraceae possess a live fleshy aril (Fig. 3a), which is membranous when dry (Fig. 3c, d). Live aril is white (Fig. 3a) or whitish except in *Erblichia odorata*, where it is reddish-orange (Fig. 3g).

The aril is inserted around the hilum (Fig. 6b), except for *Erblichia* (Fig. 4c) and some species of *Turnera* (*T. blanchetiana*, *T. hermannioides*, *T. joelii*) where it is also inserted on the basal portion of the raphe. It is glabrous except in a few species of *Turnera* (Gonzalez and Arbo 2013).

The aril may be shorter or many times longer than the seed [81–88]. It is very short in some species (Fig. 4 e), sometimes it is unilateral, limited to the rapheal area (Fig. 5 b), or may cover also the sides of the seed (Fig. 5e). In *Erblichia*, the aril almost doubles the seed length and it is very broad, its edges overlap on the opposite side of the raphe, so that the seed is completely enveloped (Fig. 3g); its cells have a high content of fatty substances.

The aril may be entire (Fig. 5a), lobed (Fig. 5d) or lacinate (Gonzalez and Arbo 2013); in *Mathurina* (Fig. 8f), with pendulous dehiscent fruits, the aril has some particular features and it is many times longer than the seed and also unusually wide. However, it does not cover the seed at all [87] because it is divided into thread-like segments almost to the base [84].

The external cells of the aril [81] may be flat (Fig. 8a, i), dome shaped (Fig. 8c, d) or papillose. In *Piriqueta* and *Turnera*, each cell may have one or several papillae (Gonzalez and Arbo 2013). Aril cells of *Mathurina* are smooth (Fig. 8e). *Streptopetalum* is the only African genus which shows some papillose cells (Fig. 8g, h).

## Phylogenetic analyses

The morphological dataset consisted of 91 characters examined in 102 taxa. The matrix contained 1.63 % missing data and all characters were parsimony informative. Nine trees were obtained using  $k = 4$ ,  $k = 3$ , and  $k = 2$ . A strict consensus of the 27 most parsimonious trees is shown in Fig. 9. All genera were inferred as monophyletic.

Most of the nodes at the base of the tree are low to moderately supported (JK values from 52 to 78; Bs values from 0.09 to 0.71). *Piriqueta* and *Turnera* lack JK support, and exhibit a Bs value of 0.02.

In the strict consensus tree (Fig. 9), the monotypic American genus *Erblichia* is sister to all the other Turneraceae. The next diverging lineages consist of *Mathurina*, *Arboa* (a clade of two species), *Stapfiella* (a clade of two

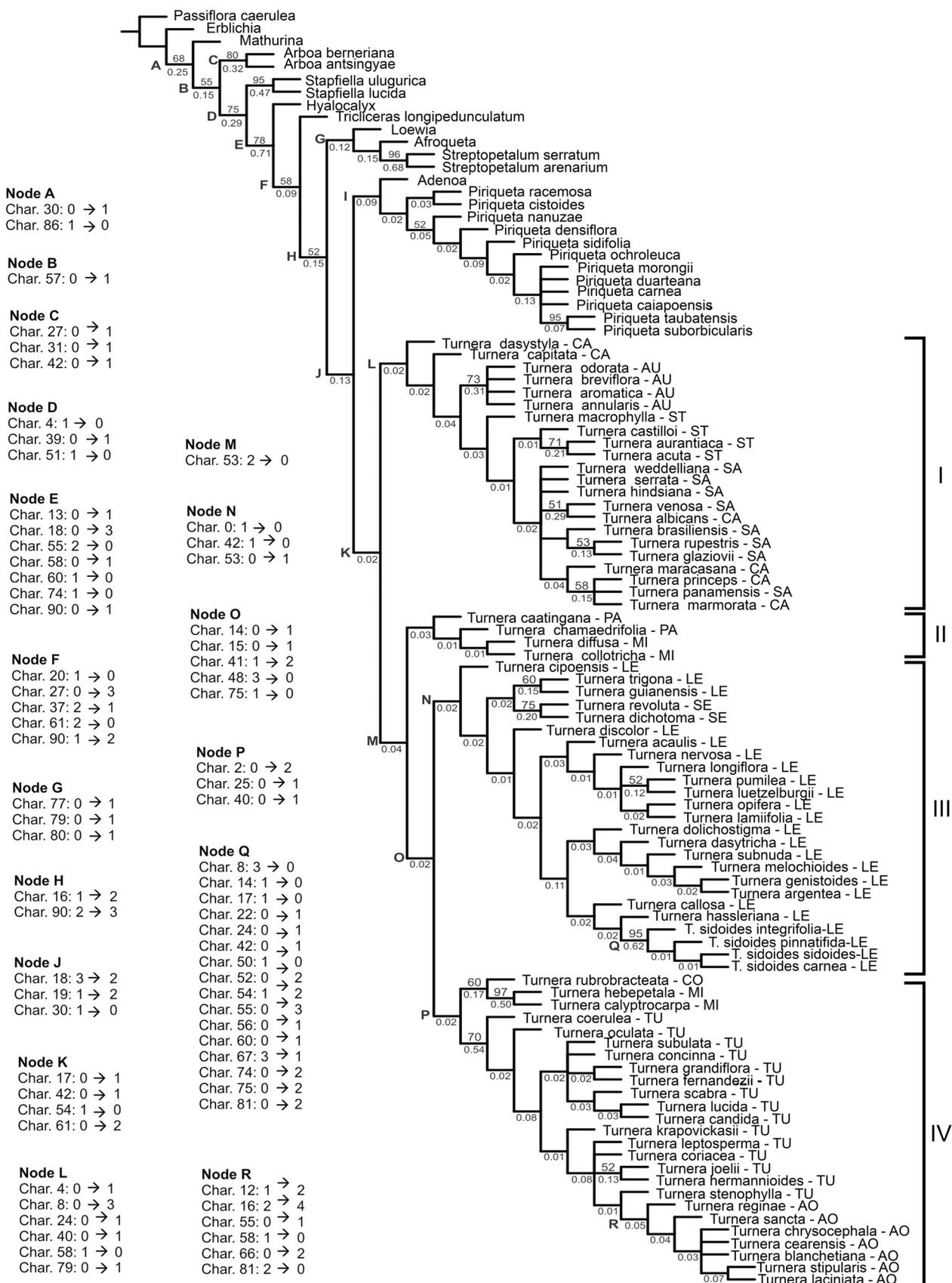
**Fig. 9** Strict consensus of the 27 trees derived from the cladistic analyses using implied weights with  $k = 2, 3$ , and 4. Numbers above and below branches refer to jackknife frequencies and Bremer support values. Letters indicate the nodes for which character substitutions are listed. Within *Turnera* clade, letters indicate AO ser. *Anomalae*, AU ser. *Annulares*, CA ser. *Capitatae*, CO ser. *Conciliatae*, LE ser. *Leiocarpae*, MI ser. *Microphyllae*, PA ser. *Papilliferae*, SA ser. *Salicifoliae*, SE ser. *Sessilifoliae*, ST ser. *Stenodictyae*, TU ser. *Turnera*. Roman numbers refer to clades according to the text

species) and *Hyalocalyx*. The genus *Tricliceras* is sister to the remaining genera; the African genera *Loewia*, *Afroqueta* and *Streptopetalum* (node G) are sister to the clade (node J) that gathers *Adenaea*, *Piriqueta* and *Turnera* with sepals connate up to half of their length [18] and a 10-veined perianth tube [19]. *Adenaea* is sister to *Piriqueta* (node I) and *Turnera* is monophyletic (node K). Within *Turnera*, the position of some species is variable according to the  $k$  value.

*Erblichia* is discriminated by the color of the aril [86] and *Mathurina* by the pendulous fruit [45] and aril divided into filaments almost to the base [84]. *Arboa* is singled out by the free corona [23], stamens insertion [27] and granulate fruit [42]. *Stapfiella* is distinguished by the basal placentation [35], while *Hyalocalyx* has inverted fruit [45] and low seed average diameter [53]. *Tricliceras* bears setiform glandular hairs and emergencies [2, 3], cymose inflorescence [12], ligulate petals [23] and siliqua-like fruit [44]. *Loewia* has a perianth tube with more than 20 delicate veins [19] and a low ratio rim/exostome diameter [75]. *Afroqueta* has nectaries on the abaxial side of leaf blade [8] and a corona fixed on the perianth [23], while *Streptopetalum* has setiform glandular hairs [2] and staminal filaments adnate to the perianth tube up to the throat [26]. *Adenaea* is characterized by a number of apomorphies, such as staminal filaments almost free [26], fruit valves with several rows of seeds [36], gynoecium longer than corolla [38], and high seed average diameter [53]. *Piriqueta* is circumscribed by the porrect stellate hairs [3] and the corona fixed on the perianth [23]. Synapomorphies of *Turnera* include the lack of a floral pedicel [17], warted, granulate or tuberculate fruit [42], lower seed average length/width ratio [54] and the seed areoles without punctiform cavity [61].

Within the monophyletic genus *Turnera*, there are four main groups. Clade I (node L) includes the species of ser. *Capitatae*, ser. *Annulares*, ser. *Stenodictyae* and ser. *Salicifoliae* with pilose staminal filaments [24] and styles [40], longitudinal seed coat ridges prevalence [58] and concave chalaza [79]; it is sister to the remaining three clades (II, III, and IV) gathered together in a large clade (node M).

Clade II is small, assembling both species of ser. *Papilliferae* (*T. caatingana* and *T. chamaedrifolia*) with *T. diffusa* and *T. colotrucha* (ser. *Microphyllae*).



Clade III (node N) includes all the species of ser. *Leiocarpae* and both species of ser. *Sessilifoliae*, with smooth fruits [42]. Clade IV (node P) is integrated by two groups, one including the only species of ser. *Conciliatae*: *T. rubrobracteata* and two species of ser. *Microphyllae* (*T. hebeptala*, *T. calyptrocarpa*). The other group gathers all the species of ser. *Turnera* and ser. *Anomaliae*, with staminal filaments adnate up to the throat [26] and high average seed length/width ratio [54]. Both groups within clade IV are moderately supported (60 and 70 JK, and 0.17 and 0.54 Bs, respectively), as well as some small subclades within *Turnera*.

#### Character optimization

Only ten morphological characters are not homoplastic, eight binary characters and two multistate: three floral characters [28, 35, 90], five seed characters [56, 57, 65, 68, 74], and two aril characters [83, 86]. Base chromosome number [89] is a multistate character not homoplastic either. Optimizations of nine of them are illustrated in Fig. 10.

The basal state of nectar pockets [28] is their absence; they appear only once in the clade gathering the species of *Turnera* ser. *Turnera* and ser. *Anomaliae*.

Placentation [35]: parietal placentation is the ancestral state, which evolves to one ovule of basal placentation only in *Stapfiella*. This is a diagnostic feature for the genus.

Seed coat design due to endotesta and exotegmen or exotesta [56]: the evolution occurred in *Turnera sidoides* (ser. *Leiocarpae*), where the crested seed coat is produced mainly by the exotesta.

The endotesta of seeds [57] is made of several layers in *Erblichia* and *Mathurina*. It evolves into a one-layered stratum in the next node.

The epidermal cells of the seed coat are generally equal all over the surface [65]. The evolution into different cells on ridges and areoles occurred in a group of species of ser. *Leiocarpae*.

The upper edge of the exostome [74] does not have a rim in seeds of *Erblichia*, *Mathurina*, *Arboa* and *Stapfiella*; in the next node it evolves developing a rim. Finally, in *T. sidoides* (node Q), the rim rises into an annular crest.

Aril pubescence [83]. The plesiomorphic state is glabrous; it evolves into pilose only in a subclade integrated by *T. panamensis* (ser. *Salicifoliae*), *T. princeps* and *T. marmorata* (ser. *Capitatae*).

Aril color [86]. It evolves only once from orange–red in the outgroup and *Erblichia*, to white or whitish in the other genera. This is a diagnostic character for *Erblichia*.

Petals degree of adnation [90]. Petals are free (state 0) in the basal genera, and they are adnate at the base to the

calyx tube (state 1) in *Hyalocalyx*; they are adnate along 1/3–1/2 of the calyx tube (state 2) in *Tricliceras*, and fused up to the throat (state 3) in the other genera.

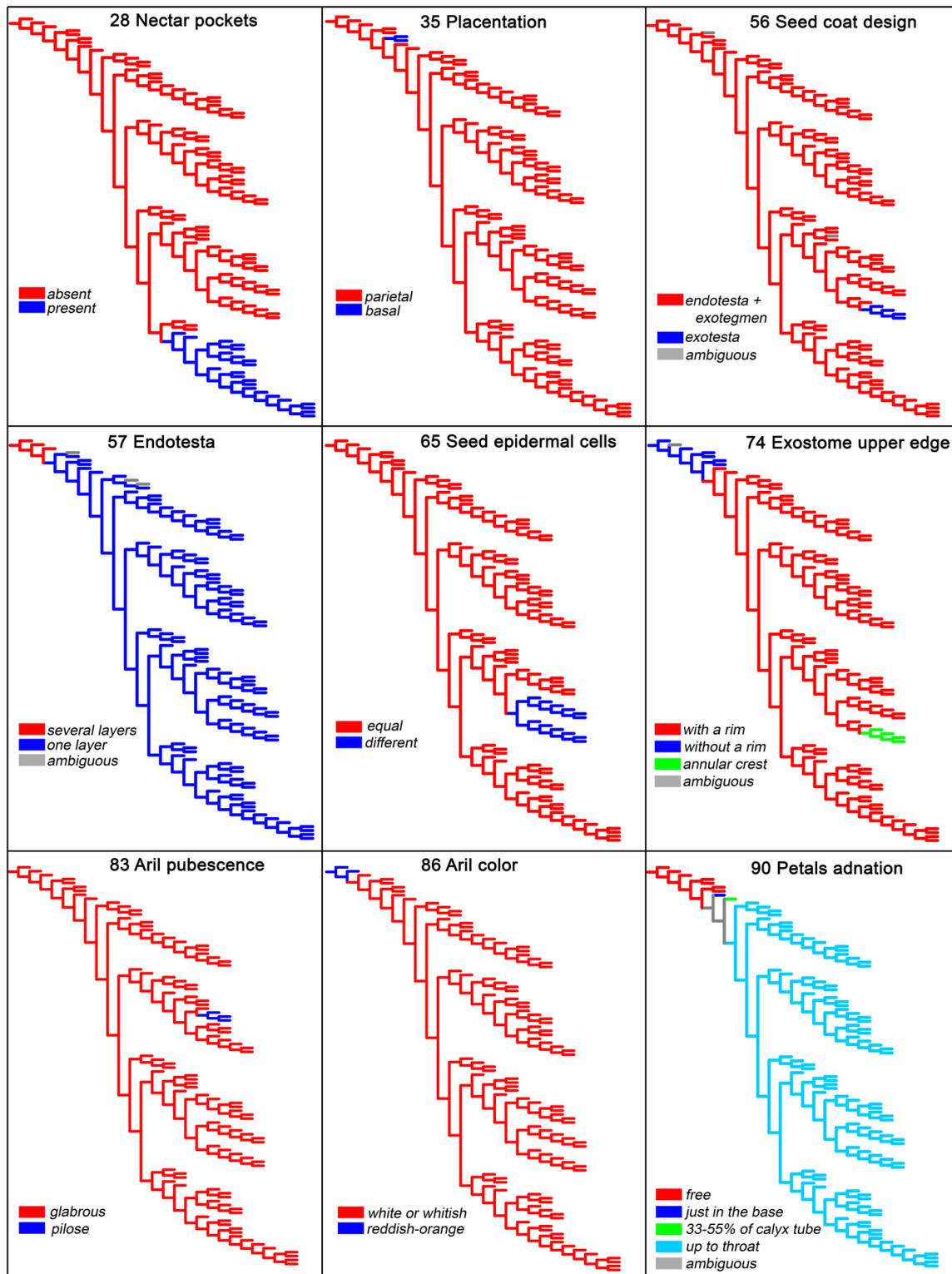
The other 80 characters show homoplasy, optimizations of nine of them are shown in Fig. 11. Some of the characters related with adnation and connation of floral whorls are important to circumscribe genera, i.e., calyx connation [18], the number of veins in the calyx/perianth tube [19], perianth appendices [23] and staminal filaments adnation [26]. Epiphyllous flowers [14], connation of staminal filaments [25], and type of seed coat [55] are valuable characters to infer groups within *Turnera*. Pedicel presence/absence [17] and fruit exocarp [42] are useful characters at both the generic and the infrageneric level.

#### Discussion and conclusions

A comparison of the topologies derived from morphological trees and molecular analyses is difficult to make because of the great differences in taxon sampling between the two type of analyses. However, the main similarities and differences among them are summarized in Table 1; data from Truyens et al. (2005) were not considered because all their results were incorporated in Chafe (2009). The position of the African genera in our analyses is similar to the results of Chafe (2009), they are in a basal position; while in Thulin et al. (2012), they are in a derived position. *Adenaea* is allied with *Erblichia* and *Mathurina* in Chafe (2009). These genera share with *Arboa* some structural features as the styles divergent at the base [39] and the seeds arranged in several rows along each fruit valve [36]. In our trees, *Adenaea* is associated with *Piriqueta* in a clade, which is sister to *Turnera*, like in Thulin et al. (2012). This association also has morphological support because these genera have a 10-veined perianth tube.

It seems like the acquisition of a floral tube is linked with some sort of evolutionary advantage to the Turneraceae, since it occurs in 66 % of the genera: *Erblichia*, *Mathurina*, *Arboa* and *Stapfiella*. The only ones without a calyx tube occupy a basal position in our topology. *Hyalocalyx*, which is sister to the remaining genera, has a 15-veined calyx tube.

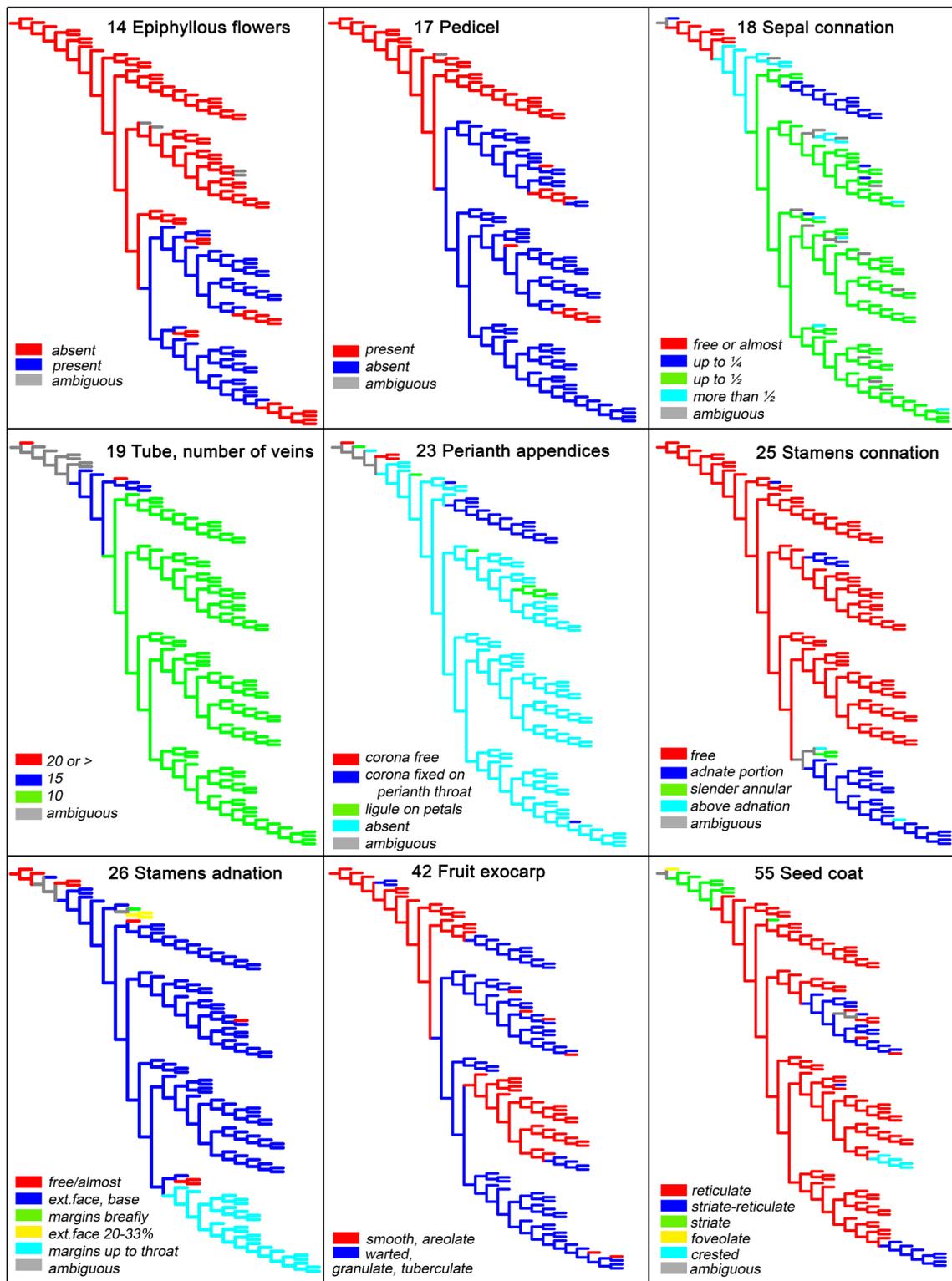
The morphological complexity of the floral tube increased in several steps: (1) the adnation of petal claws to calyx [90] found in *Tricliceras* and *Loewia*, developing a perianth tube. (2) The adnation of staminal filaments [26] to the perianth tube: in *Afroqueta* there is a brief marginal adnation while in *Streptopetalum*, the adnation takes place along 1/3–1/2 of the tube, turning that region into an appendicular hypanthium. (3) The fusion of lateral sepal veins with the main petal vein, shaping a 10-veined



**Fig. 10** Optimization of non-homoplastic characters. The number of each character, its shortened name, and states are indicated to the left of the tree skeletons

perianth tube in *Adenoa*, *Piriqueta* and *Turnera* [19]. (4) The development of nectariferous pockets [25–28] by means of the staminal filaments marginal adnation and

connation up to the throat found in ser. *Turnera* and ser. *Anomalae*; in this case, the whole tube is an appendicular hypanthium.



**Fig. 11** Optimization of homoplastic characters. The number of each character, its shortened name, and states are indicated to the left of the tree skeletons

The brief adnation found in *Afroqueta* seems like a morphological step prior to the one found in *Turnera* ser. *Turnera* and ser. *Anomalae*, producing long nectariferous

pockets. Adnation of staminal filaments [26] to perianth tube is a homoplastic character; marginal adnation appeared in *Adenoa*, where it is very brief, and then in

*Turnera* series *Anomalae* and *Turnera*, where it is well developed.

The color of arils seems to be related to seeds dispersion. The reddish-orange aril of *Erblichia* is associated with ornithochory (Thulin et al. 2012). *Passiflora caerulea* L. has seeds with enveloping red aril (Deginani 2001) similar to the one of *Erblichia*, but the fruit is an indehiscent berry. Probably in both cases, the seeds are dispersed along with birds' feces.

Myrmecochory (Fig. 3h) has been cited for several species of *Turnera* (Barrett 1978), *Piriqueta* (Arbo 1995) and the Malagasy genus *Arboa* (Thulin et al. 2012). It has also been recorded in *Turnera oculata* Story (Jaarsveld 2012), one of the two African species of the genus. In the other genera, the dispersion is probably carried out also by ants, given the characteristics of the aril, which becomes an elaiosome, much appreciated by these insects. Pijl (1982) observed that ants react very quickly to the presence of elaiosomes, probably due to the existence of volatile compounds associated to the lipids. This author also considers that the presence of extrafloral nectaries in *Turnera* would attract the ants to diaspores. Cuautle et al. (2005) analyzed the effects of ant behavior and presence of extrafloral nectaries on seed dispersal of *Turnera ulmifolia* (= *T. velutina* C.Presl).

The red–orange aril of *Passiflora caerulea* looks like a plesiomorphic feature in Turneraceae, since it exists only in *Erblichia*; while all the other genera have a white or whitish aril, typical of mirmecochory. The only exception is *Mathurina*, where the aril, divided into long threads, is an adaptation to anemochory.

The addition of seed characters to the cladistic analysis led to several changes in the morphological trees with respect to *Turnera*. There is a better resolution of the series, which are distributed in four main clades. In molecular phylogenetic trees (Truyens et al. 2005; Chafe 2009), the only monophyletic series was ser. *Turnera*, characterized by a base chromosome number  $x = 5$ .

Series *Capitatae*, ser. *Stenodictyae* and ser. *Salicifoliae* form clade I including also ser. *Annulares*, which was associated with ser. *Turnera* and ser. *Anomalae* in the previous morphological analysis (Arbo and Espert 2009).

*Turnera chamaedrifolia*, with  $x = 13$  (ser. *Papilliferae*), is associated with *T. diffusa* with  $x = 7$  (ser. *Microphyllae*) in molecular analyses (Truyens et al. 2005; Chafe 2009). In our clade II, both species of ser. *Papilliferae* (*T. caatingana* and *T. chamaedrifolia*) are related to *T. diffusa* and *T. colotracha*, two species of ser. *Microphyllae*. The base number  $x = 13$  found in *T. chamaedrifolia* probably originated by aneuploidy from a species with  $2n = 4x = 28$  chromosomes (Fernández 1987).

Series *Leiocarpae* was paraphyletic in both the previous morphological (Arbo and Espert 2009) and molecular

phylogenetic trees (Truyens et al. 2005; Chafe 2009); here, all the species are grouped in clade III where both species of ser. *Sessilifoliae* are nested. *Turnera sidoides* has several apomorphies and, in particular, its seeds are matchless in the family. In the molecular phylogenetic analyses (Truyens et al. 2005; Chafe 2009), the subspecies of *T. sidoides* formed a separate clade, unresolved among the first diverging lineages of the tree, like in the previous cladistic analyses (Arbo and Espert 2009). In the present study, this species fits in clade III ser. *Leiocarpae*-ser. *Sessilifoliae*, and interestingly, it is in a rather derived position.

In a previous morphological analysis (Arbo and Espert 2009), all species of ser. *Microphyllae* were gathered in one clade, whereas in this study *T. calyptrocarpa* and *T. hebetata*, with a slender basal annular connection of staminal filaments, are gathered with *T. rubrobracteata*, the only species of ser. *Conciliatae*. They are integrated in clade IV with ser. *Turnera* + ser. *Anomalae*, which present nectar pockets. This is consistent with molecular phylogenetic analyses (Truyens et al. 2005; Chafe 2009), in which *T. calyptrocarpa* is related to *T. cearensis* and *T. bahiensis*, the only species of ser. *Anomalae* analyzed.

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