

The first Loranthaceae fossils from Africa

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Abstract

An ongoing re-investigation of the early Miocene Saldanha Bay (South Africa) palynoflora, using combined light and scanning electron microscopy (single grain method), is revealing several pollen types new to the African fossil record. One of the elements identified is Loranthaceae pollen. These grains represent the first and only fossil record of Loranthaceae in Africa. The fossil pollen grains resemble those produced by the core Loranthae and are comparable to recent Asian as well as some African taxa/lineages. Molecular and fossil signals indicate that Loranthaceae dispersed into Africa via Asia sometime during the Eocene. The present host range of African Loranthaceae and the composition of the palynoflora suggest that the fossil had a range of potential host taxa to parasitise during the early Miocene in the Saldanha Bay region.

Keywords: Santalales, diagnostic pollen, host plants, Miocene, palaeoecology, palaeophytogeography, parasitic plants, pollen morphology

The Loranthaceae is a large family with c. 76 genera and at least 1000 species divided into five tribes (Nickrent 1997-onwards; Nickrent et al. 2010). The family is widely distributed and occurs in tropical to temperate regions of Australasia, Asia, the Middle East, Africa, Europe, and Central and South America (e.g. Barlow 1983; Polhill & Wiens 1998), showing a clear geographic split between a New World group (Psittacanthinae Engl.) and Old World-Australasian lineages (Elythrantheae Engl. and Lorantheae Rchb.; e.g. Nickrent et al. 2010; Grímsson et al. 2017, 2018). The c. 238 species and 21 genera occurring in Africa (Table I) are considered to be the most derived in the family. Most of the African genera/species are endemic, with only Helixanthera and Taxillus extending into Asia. Helixanthera, occurring from Africa to Indonesia, is regarded as the most primitive Lorantheae genus thriving in continental Africa (Polhill & Wiens 1998). Even though Lorantha-

ceae are currently found all over Africa (except the Sahara desert), it has been suggested that they dispersed to the continent during the Cretaceous (Gondwanan derivation) or Eocene times (Asian derivation) (Barlow 1983, 1990; Polhill & Wiens 1998), however, no fossil Loranthaceae have ever been reported from this part of the world. The fossil record of Loranthaceae, recently summarised by Grímsson et al. (2017, figures 10, 11, and file S4), shows that the family already had a global distribution during the Eocene, occurring on all continents except Africa and Antarctica. The Loranthaceae have a fragmentary fossil record composed solely of fossil pollen (Grímsson et al. 2017), most likely due to the ecology and life cycle of Loranthaceae (small woody plants, with relatively few leaves, their fruits are ingested by birds, and seeds germinate immediately after regurgitation; see Polhill & Wiens 1998). Therefore, the only way to trace the origin and evolution of this family, in

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Table I. African	n Loranthaceae	genera	and	their h	osts.
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Genus	Number of species	African species	Occurrence in Africa	Recorded host families in Africa	See Table
Helixanthera	c. 45	12	Tropical, scattered around edge of continent	Anacardiaceae, Bignoniaceae, Boraginaceae, Burseraceae, Combretaceae, Euphorbiaceae, Fabaceae, Lauraceae*, Loranthaceae, Malvaceae*, Moraceae, Ochnaceae, Phyllanthaceae, Rhamnaceae, Bubiceae* Euroceae Sanctaceae	S1
Plicosepalus	12	12	Eastern side of Africa to Angola and South Africa	Anacardiaceae, Apocynaceae, Burseraceae, Combretaceae, Fabaceae, Rubiaceae	S2
Emelianthe	1	1	Drier parts of E. and NE. Africa	Anacardiaceae, Burseraceae, Euphorbiaceae, Malvaceae	S 3
Pedistylis	1	1	Southern Africa	Anacardiaceae, Combretaceae, Ebenaceae, Fabaceae, Meliaceae, Moraceae	S4
Actinanthella	2	2	SE. and S. Africa	Capparaceae, Erythroxylaceae, Oleaceae	S5
Oncocalyx	13	13	Drier forests and bushland of eastern and southern Africa	 Anacardiaceae, Apocynaceae, Boraginaceae, Burseraceae, Cannabaceae, Capparaceae, Celastraceae, Combretaceae, Ebenaceae, Euphorbiaceae, Fabaceae, Malvaceae, Pittosporaceae, Rhamnaceae, Salicaceae, Salvadoraceae, Tamariaceae, Zygophyllaceae 	S 6
Spragueanella	2	2	E. and SC. Africa along coast and extending into mountains in dryer forest	Podocarpaceae, Putranjivaceae	S 7
Oliverella	3	3	Eastern and south-central Africa in coastal and deciduous bushland and mixed woodland	Combretaceae, Euphorbiaceae, Fabaceae, Malvaceae	S8
Berhautia	1	1	Senegal and Gambia	Combretaceae	S 9
Englerina	25	25	Tropical Africa	Achariaceae, Asteraceae, Bignoniaceae, Boraginaceae, Buddlejaceae, Burseraceae, Clusiaceae, Combretaceae, Ebenaceae, Fabaceae, Loganiaceae, Malvaceae, Oleaceae, Primulaceae, Proteaceae, Rhamnaceae, Rubiaceae, Rutaceae*, Sapindaceae	S10
Agelanthus	59	59	Africa south of the Sahara	 Anacardiaceae, Apocynaceae, Asteraceae, Boraginaceae, Burseraceae, Cannabaceae, Cappaeaceae, Celastraceae, Chrysobalanaceae, Combretaceae, Convolvulaceae, Erythroxylaceae, Euphorbiaceae, Fabaceae, Iteaceae, Juglandaceae*, Lamiaceae, Loranthaceae, Lythraceae*, Malvaceae, Meliaceae, Moraceae, Olacaceae, Meliaceae, Moraceae, Olacaceae, Oleaceae, Phyllanthaceae, Plumbaginaceae, Rhamnaceaee, Rosaceae*, Rutaceae*, Proteaceae, Salicaceae, Solanaceae*, Limaceae, Urticaceae, Vitaceae 	511
Tapinanthus	30	30	Tropical and southern Africa	Anacardiaceae, Orncaceae, Vitaceae Anacardiaceae, Apocynaceae, Asphodelaceae, Asteraceae, Burseraceae, Celastraceae, Combretaceae, Crassulaceae, Ebenaceae, Euphorbiaceae, Fabaceae, Juglandaceae*, Kirkiaceae, Lamiaceae, Loranthaceae, Malvaceae, Meliaceae, Melianthaceae, Moraceae, Myrtaceae, Ochnaceae, Phyllanthaceae, Proteaceae, Rosaceae, Rhamnaceae, Rutaceae, Salicaceae, Salvadoraceae, Tamaricaceae	S12

Table I. (Continued).

Genus	Number of species	African species	Occurrence in Africa	Recorded host families in Africa	See Table
Moquiniella	1	1	Southern Namibia and the Cape Province of South Africa	Anacardiaceae, Apocynaceae, Ebenaceae, Fabaceae, Hypericaceae, Malvaceae, Moraceae, Rosaceae*, Salicaceae	S13
Globimetula	13	13	Tropical Africa	 Anacardiaceae*, Burseraceae, Chrysobalanaceae, Combretaceae, Fabaceae, Malvaceae*, Meliaceae, Moraceae, Myrtaceae*, Phyllanthaceae, Proteaceae, Rutaceae* 	S14
Taxillus	35	1	Coast of Kenya	Fabaceae	S15
Vanwykia	2	2	Eastern and south-eastern Africa	Fabaceae, Moraceae	S16
Septulina	2	2	Western Cape Province of South Africa and southern Namibia	Aizoaceae, Anacardiaceae , Fabaceae , Solanaceae, Tamaricaceae	S17
Oedina	4	4	Montane forests from Tanzania to northern Malawi		S18
Oncella	4	4	Montane and coastal areas of eastern Africa	Phyllanthaceae, Malvaceae, Meliaceae	S19
Erianthemum	16	16	Eastern and southern Africa	 Anacardiaceae, Archariaceae, Asteraceae, Bignoniaceae*, Burseraceae, Celastraceae, Combretaceae, Ebenaceae, Euphorbiaceae, Fabaceae, Lamiaceae, Loganiaceae, Malvaceae, Meliaceae, Myrtaceae*, Phyllanthaceae, Proteaceaea, Rhamnaceae, Rosaceae*, Rutaceae*, Sapotaceae 	S20
Phragmanthera	34	34	Tropical forests of Africa, few extend into dry habitats in south-central and southern Africa	 Anacardiaceae*, Annonaceae, Boraginaceae, Burseraceae, Casuarinaceae*, Combretaceae, Euphorbiaceae, Fabaceae, Irvingiaceae, Lauraceae*, Malvaceae*, Melianthaceae, Moraceae, Myrtaceae*, Rhamnaceae, Rubiaceae, Rutaceae*, Tamaricaceae, Phyllanthaceae, Proteaceae, Sapotaceae 	S21

Notes: Families with introduced host taxa are marked with asterisk*. Host families known from the fossil palyno-assemblage appear in **bold**. Loranthaceae systematics and distribution summarised from Polhill and Wiens (1998), data on host taxa compiled from Wiens and Tölken (1979), Visser (1981), Dean et al. (1994), Polhill and Wiens (1998, 1999), Dzerefos et al. (2003), Roxburgh and Nicolson (2005), Veste (2007), Didier et al. (2008), Ogunmefun et al. (2013), Dlama et al. (2016) and Okubamichael et al. (2013, 2016). See also Tables S1–S21 in Supplemental data.

time and space, is to study fossil Loranthaceae pollen in relation to phylogeny. Grímsson et al. (2018) evaluated the correlation of pollen morphology and molecular phylogenetic relationships within Loranthaceae and discovered that most pollen types in this family are linked to a single genus or discrete evolutionary lineages. Since pollen types produced by most extant members of the Loranthaceae are distinct (Feuer & Kuijt 1978, 1979, 1980, 1985; Kuijt 1988; Liu & Qiu 1993; Han et al. 2004; Roldán & Kuijt 2005; Caires 2012; Caires et al. 2012, 2014, 2017; Grímsson et al. 2017, 2018) and cannot be confused with pollen from other related families, fossil Loranthaceae pollen give the potential to trace modern lineages back in time.

Here we describe a new fossil Loranthaceae pollen type from the earliest Miocene of Saldanha Bay, South Africa. These fossils are the first representatives of this family in the fossil record of Africa. The diagnostic light microscopy (LM)- and scanning electron microscopy (SEM)-based features of the pollen provide sufficient support to assign the fossils to a distinct lineage within the Loranthaceae. Based on the taxonomic affiliation to extant taxa the palaeophytogeographic signals and palaeoecological aspects of these fossil grains are discussed and potential host taxa are suggested from the currently known palaeo-palynoflora.

Material and methods

The sedimentary rock containing the fossil Loranthaceae pollen is from core sample #114755 collected at Saldanha Bay, South Africa. The sediments are believed to be of earliest Miocene age. A Chattian to early Miocene age for the Saldanha Bay deposits is suggested on the base of the dinoflagellate indicator taxa Distatodinium craterum Eaton, Chiropteridium lobospinosum Gocht, Homotryblium plectilum Drugg et Loeblich Jr. as well as Impagidinium paradoxum (Wall 1967) Stover et Evitt 1978 (see details in Roberts et al. [2017] including supplements). For a full geological, stratigraphic, palaeontological and palaeoenvironmental background of this locality/core see Roberts et al. (2017). The sedimentary rock sample was processed and fossil pollen grains extracted according to the protocol outlined in Grímsson et al. (2008). The fossil Loranthaceae pollen grains were investigated both by LM and SEM using the single grain method as described in Zetter (1989). The description of fossil Loranthaceae pollen includes diagnostic features observed both in LM and SEM. Pollen terminology follows Punt et al. (2007; LM) and Hesse et al. (2009; SEM). Loranthaceae fossil material (SEM stubs) from Saldanha Bay, South Africa, are stored in the collection of the Department of Palaeontology, University of Vienna, Austria under the accession numbers IPUW 7513/ 211 and IPUW 7513/216.

Systematic palaeontology

The fossil pollen described here falls within the variation of Pollen Type B defined by Grímsson et al. (2018). Pollen of this type is oblate (to various degrees), triangular to trilobate in polar view and shows a \pm psilate sculpturing in LM. Usually, further sculpture details are not observed in LM, but some pollen grains show a clear exine thickening or thinning at the pole and along the colpi or in the mesocolpium. The pollen is syn(3) colpate, see figure 1 in Grímsson et al. (2018) for a schematic drawing. Since the fossil pollen grains described here show combining features known from four extant Loranthaceae genera (see later), we classify this fossil taxon as a morphotype (MT) named after the locality where the pollen occurs.

> Family Loranthaceae Juss. Tribe Lorantheae Rchb. Saldanha MT, aff. Lorantheae Figures 1–3

Description. — Pollen, oblate, concave-triangular to trilobate in polar view, elliptic in equatorial view, equatorial apices obcordate to T-shaped; size small, polar axis 8.8–12.5 μ m long in LM, equatorial diameter 20–25 μ m in LM, 15–22 μ m in SEM; syn(3) colpate; exine 0.8–1.0 μ m thick, nexine thinner than sexine (LM), triangular intercolpial nexine thickenings in polar area (LM); tectate; sculpture psilate in LM, nanoverrucate to granulate in area of mesocol-

pium in SEM, nanoverrucae $0.2-0.5 \mu m$ in diameter, verrucae composed of conglomerate granula; margo well developed, margo psilate or partly granulate, margo with triangular protrusions in polar area (SEM); colpus membrane nanoverrucate and granulate (SEM).

Remarks. — Compared to extant pollen this fossil MT shows a suite of features found only within the tribe Lorantheae. This combination of outline, size, colpi arrangement, thickening of nexine (LM), and sculpture observed under SEM is typical for taxa placed in the Subtribes Dendropthinae Nickrent & Vidal-Russell (e.g. Tolypanthus, Dendrophthoe), Scurullinae Nickrent & Vidal-Russell (e.g. Taxillus), and partly Emelianthinae Nickrent & Vidal-Russell (Phragmanthera). Both Tolypanthus maclurei (Merr.) Danser and Dendrophthoe pentandra (L.) Mig. pollen is very similar to the Saldanha MT (see Table II). The Tolypanthus maclurei pollen (see figure 38 in Grímsson et al. 2018) is usually larger than the fossils, and the D. pentandra pollen (see figure 36 in Grímsson et al. 2018) tends to have a slightly thicker nexine, and wall peculiarities in the polar area are hard to distinguish (LM). Otherwise the pollen of these two taxa is almost identical to the Saldanha MT. Pollen of Taxillus caloreas (Diels) Danser (see figure 49 in Grímsson et al. 2018) has a more striking and larger, hexagonal in outline, thickening of nexine in the polar area that differs from that observed in the Saldanha MT. Most other features are comparable to those observed in the fossils. Pollen of Phragmanthera rufescens (DC.) Balle (see figure 43 in Grímsson et al. 2018) is more or less identical to the Saldanha MT, except the P. rufescens pollen is slightly larger and the exine is thicker (LM).

Despite the number of fossil Loranthaceae pollen reported so far only few grains/types have been studied using SEM (see Grímsson et al. 2017). Of those studied using SEM only two MTs, the Changchang MT form the middle Eocene of China, and the Altmittweida MT from the late Oligocene-early Miocene of Germany (Table II; see also Grímsson et al. 2017), indicate a possible lineage relation (Lorantheae) with the African fossils. The broadly rounded apices, the rhombic structures covering equatorial apertures, and the merely granulate sculpture clearly distinguishes the Chinese Changchang MT from the African Saldanha MT (see Table II). Still, the minute sculpture and the basic form of the Changchang MT also link it to the Lorantheae, especially to the Scurrulinae (Taxillus, Scurulla) and Amyeminae (Amyema), and Grímsson et al. (2017, p. 21) described this pollen MT as 'a Scurrulinae pollen with an Amvema-like margo'. Therefore, the Changchang MT most likely belongs to an extinct or ancestral Lorantheae lineage related to



Figure 1. LM (**A**) and SEM (**B**) micrographs of fossil Loranthaceae pollen from the early Miocene of Africa. **A.** Saldanha morphotype (MT) pollen grains in equatorial and polar view. Note triangular intercolpial nexine thickenings in polar area. **B.** Saldanha MT pollen grains in polar view. Equatorial apices are obcordate to T-shaped and the margo is psilate or partly granulate and with triangular protrusions in polar area. Scale bars $-10 \mu m$ (A, B).

the core Lorantheae. The emarginate outline in equatorial view, the reduced sexine in the polar area, the microverrucate sculpture in SEM, and the pollen size clearly distinguishes the German Altmittweida MT from the African Saldanha MT. Extant pollen very similar to the Altmittweida MT can be found in two



Figure 2. SEM micrographs of fossil Loranthaceae pollen from the early Miocene of Africa. **A–D.** Close-ups of central polar area showing margo with triangular protrusions in polar area. **E–H.** Close-ups of apex showing obcordate to T-shaped apices, and psilate or partly granulate margo. Scale bars – $1 \mu m$ (A–H).



Figure 3. SEM micrographs of fossil Loranthaceae pollen from the early Miocene of Africa. A–D. Close-ups of mesocolpium showing nanoverrucate to granulate sculpture (SEM). Scale bars – $1 \mu m$ (A–D).

extant species of Lorantheae, *Amyema gubberula* Danser and *Helixanthera kirkii* (Oliv.) Danser. It is therefore also likely that the Altmittweida MT belongs to a lineage related to the core Lorantheae.

Discussion

The African Lorantheae fossils in a global (time and space) context

The fossil pollen record of Loranthaceae (e.g. *Gotha-nipollis*) recently summarised by Grímsson et al. (2017) shows that the family had a worldwide distribution already during the Eocene, with representatives found in South America, North America, Europe, East Asia, and Australasia. Based on this palaeo-phytogeographic pattern it is most likely that Loranthaceae were also present in Africa during that time. The lack of fossil Loranthaceae pollen in the African record should be considered an artefact caused primarily by preparation techniques and study methods, or palynologists working on African material not knowing this typical *Gothanipollis* type. Accepting this, the dispersal of Loranthaceae into

Africa might have occurred in the Southern Hemisphere before the final phases of the Gondwana breakup (Late Cretaceous) or in the Northern Hemisphere via Asia (early Eocene). Unfortunately, the majority of Eocene fossil Loranthaceae pollen found in the Southern Hemisphere (South America, Australasia) has mostly been studied using LM only (e.g. Romero & Castro 1986; Raine et al. 2011) and is therefore of very limited use for interfamilial segregation. In a molecular phylogenetic context (see figure 2 in Grímsson et al. 2018) the present African Loranthaceae show a closer relation to South, Southeast and East Asian lineages than any other, and are clearly most distantly related to American Loranthaceae. It is interesting, based on pollen morphology, that the earliest Miocene fossils from Saldanha Bay suggest the same close relation to Asian taxa (Table II) and 'no' relation to any of the American lineages. Grímsson et al. (2017) established that several major lineages of Loranthaceae were present during Eocene in the Northern Hemisphere, with records including representatives of extinct or ancestral lineages with affinities to both root-parasitic genera (Nuytsia/Nuytsieae) and epiphytic lineages

	Amon (III) of frontions a	men commun ommun bonnen					
	Tolypanthus maclurei	Dendrophthoe pentandra	Taxillus caloreas	Phragmanthera rufescens (s.l.) ^a	Saldanha MT (this study)	Changchang MT	Altmittweida MT
Age/epoch	Recent	Recent	Recent	Recent	Early Miocene	Middle Eocene (Lutetian- Bartonian)	Late Oligocene- early Miocene (Chattian- Aouiranian)
Distribution/ locality	East Asia	South, East and Southeast Asia	East Asia	Tropical Africa	Saldanha Bay, South Afruca, core sample #114 755	Changchang Basin, close to Jiazi Town, Qiongshan County, Hainan, China	Altmittweida, Saxony, Germany
P/E ratio Outline p.v.	oblate trilobate to straight-	oblate concave-triangular to	oblate concave-triangular	oblate concave-	oblate concave-triangular to	oblate concave-triangular	oblate convex-triangular
	triangular	trilobate		triangular	trilobate	to broadly trilobate	
Outline eq. v.	elliptic	elliptic	elliptic	elliptic	elliptic		emarginate
Equatorial apices P in LM (1111)	obcordate 8.3–15.8	obcordate 13_3-15	obcordate	T-shaped 15–18.3	obcordate to T-shaped 8 8–12.5	broadly rounded	broadly obcordate 4 4–5 5
E in LM (µm)	25-30	21.7–25.8	23.3–30	26.7–31.7	20-25	21.1 - 24.4	14.4–17.8
Aperture Exine thickness in LM (µm)	syn(3)colpate 0.8–1.3	syn(3)colpate 1.1–1.3	syn(3)colpate 1.0–1.3	syn(3)colpate 1.1–1.4	syn(3)colpate 0.8–1.0	syn(3)colpate 0.9–1.1	syn(3)colpate 0.9–1.1
Wall peculiarities	triangular intercolpial	sexine partly reduced in	hexagonal nexine	triangular	triangular intercolpial	rhombic structures	intercolpial nexine
	thickening of nexine	polar area, colpi	thickening in	intercolpial	thickening of nexine	(opercula)	thickening at
	in polar area	widening to a small neid	polar arca	unickening of nexine in polar area	in polar area	covering equatorial apices	pole, sexme partly reduced in polar area
Sculpture (SEM)	nanoverrucate to	nanoverrucate to granulate	nanoverrucate to	nanoverrucate	nanoverrucate to	granulate	nano- to
	granulate		granulate	to granulate	granulate		microverrucate to granulate
Type and size of sculpture elements (µm)	verrucae 0.2–0.5 (–0.8)	verrucae 0.2–0.6	verrucae 0.1–0.5	verrucae 0.1– 0.6	verrucae 0.2–0.5		verrucae 0.2–1.3
Margo (SEM)	well developed, psilate	well developed, psilate or	well developed,	well · · ·	well developed, psilate	well developed,	psilate to
	or paruy granuate, with triangular	paruy granulate to nanoverrucate, with	psuate with lew nanoverrucae or	aevelopea, psilate or	or paruy granuate, with triangular	psuate	microverrucate, granulate
	protrusions in polar	triangular protrusions in	granula in polar	partly	protrusions in polar		
	area	polar area	area	granulate	area		
Colpus membrane (SEM)	nanoverrucate and granulate	nanoverrucate and granulate	nanoverrucate and granulate	nanoverrucate and	nanoverrucate and granulate	granulate	nanoverrucate and granulate
				granulate			
					-		•

Note: Distribution of extant taxa from Qui and Gilbert (2003) and Polhill and Wiens (1998). Pollen morphology of extant taxa summarised from Grimsson et al. (2018). Pollen morphology of fossil mophotypes summarised from Grimsson et al. (2017). ^a*Phragmanthera rufescens* has been widely applied as an aggregate for tropical African *Phragmanthera*. According to Polhill and Wiens (1998) *P. rufescens* is only known from Guinée and the Casamance region of southern Senegal, but the sample figured in Grimsson et al. (2018) is from Gameroon and might therefore represent *P.* kamerunensis or another Phragmanthera species. (Lorantheae, Psittacantheae, *Notanthera*, Elytrantheae). In this scenario, it seems more likely that the ancestor(s) of the fossils described here and the current African lineages dispersed into Africa from Asia (northern route) during the Eocene. For now, dispersal into Africa in the Southern Hemisphere during the final phases of the Gondwana breakup cannot be ruled out. If some Loranthaceae were dispersed via a southern route then those lineages became extinct in Africa during the Cainozoic.

Time of origin and divergence of African Loranthaceae lineages

Fossil constrained molecular dating by Grímsson et al. (2017) suggests that *Tubeia* (A-type pollen) and Loranthaceae with B-type pollen diverged in the early Eocene (~50 Ma). A primary radiation is believed to have followed shortly thereafter involving the formation of both 'New World' (root parasites, Elytrantheae, Psittacantheae) and 'Old World' (Lorantheae) clades. Crown group radiation in the Lorantheae is then believed to have started at the latest in the late Eocene (\geq 38 Ma), with a second major radiation taking place ~10 million years later (latest in the Oligocene) involving among others the 'Old World' core Lorantheae (Grímsson et al. 2017). Unfortunately, there are no current African records from the Eocene or Oligocene so far, but one could expect to find Loranthaceae pollen showing morphology characteristic for the core crown Lorantheae (e.g. Amyeminae, Dendropthinae) in such samples. Still, the Saldanha MT suggests that until the earliest Miocene pollen producing Loranthaceae in Africa (at least southern Africa) still had the basal Lorantheae pollen form. More diverged lineages/genera must therefore have evolved no earlier than during the latest part of the early Miocene. The age and pollen morphology of the Saldanha MT fit perfectly with the suggested core crown group radiation of Lorantheae and the alleged formation of extant lineages/modern genera lasting until the middle Miocene (\geq 9 Ma; see figure 9 in Grímsson et al. 2017). Fossil pollen showing derived features within the Lorantheae, e.g. Emelianthinae and Tapinanthinae, are most likely to be found in sediments younger than earliest Miocene.

Ecology and potential hosts of the Saldanha MT

Loranthaceae are currently found in all parts of Africa except the Sahara desert where there is little vegetation. They occur in various habitats, ranging from sea-level to mountain tops, in grasslands as well as rainforests and semi-deserts. Their only requirements seem to be the presence of suitable host plants and dispersal mechanisms (e.g. birds) to carry them between hosts (e.g. Visser 1981; Polhill & Wiens 1998). It is hard to pinpoint the preferred host of a fossil taxon and if it had a narrow (specialist) or wide (generalist) host range. Based on the available host ranges of recent African Loranthaceae (Table II; Tables S1-S21 in Supplemental data) it seems that most of the genera are generalists and parasitising many species/genera/families. The fossil palynoassemblage containing the Saldanha MT is extremely taxon rich (Roberts et al. 2017) and composing pollen from at least 150 different angiosperms verified using SEM (Grímsson et al. unpublished data). Many of the fossil pollen types belong in families that are known to be parasitised by recent African Loranthaceae. These include Anacardiaceae, Asteraceae, Casuarinaceae, Euphorbiaceae, Fabaceae, Myrtaceae, Oleaceae, Proteaceae, Santalaceae, Sapindaceae, Sapotaceae, and the gymnosperm family Podocarpaceae. Based on the recorded host families listed in Table II, it is likely that every recent Loranthaceae genus would find a suitable host plant in the palaeo-vegetation at Saldanha Bay during the earliest Miocene. The palaeo-vegetation units in the Saldanha Bay region are believed to have been very diverse (Roberts et al. 2017), composed, e.g. of various lowland wetland and marshland forests (mangrove, riparian/swamp) and different mixed evergreen broad-leaved and/or coniferous forests stretching into the surrounding highlands (see figure 13 in Roberts et al. 2017). All potential vegetation units (habitats) recorded by Roberts et al. (2017) would be suitable for Loranthaceae based on the current vast habitat range of the family in Africa (e.g. Polhill & Wiens 1998).

Conclusion and outlook

Despite the numerous palaeopalynological investigations on African Cretaceous to Miocene microflora there are no comprehensive high resolution SEMbased studies so far. Even though the potential for studying pollen using combined LM and SEM from African sediments was already established by Coetzee in the 1980s (Coetzee 1981, 1983; Coetzee & Muller 1984; Coetzee & Praglowski 1984, 1988), she only presented a handful of fossil taxa using SEM and no other African palynologist has used this combined method since. Based on the current Loranthaceae fossil record it is very unlikely that the family should be absent from Eocene and Oligocene sediments in Africa. It is more likely that the preparation methods (including sieving) and study techniques (counting up to 300 grains in LM) biased the outcome, or/and when present, the palynologist did not know this distinct pollen type and disregarded it or simply misidentified it. Our combined LM and SEM based study shows that the early Miocene South African Loranthaceae fossils resemble the core Lorantheae (Dendrophae, Scurullae) and the more derived lineages (Tapinanthae, Emelianthae) were not present in this area at the time of accumulation. Molecular and fossil signals suggest that Loranthaceae dispersed into Africa via Asia (northern route) during the Eocene. Also, the recent host range of African Loranthaceae and the palaeo-palynological spectrum suggest that the fossil would have no problems finding a host plant during the early Miocene in the Saldanha Bay region. To fully enlighten the African palaeophytogeographic history of this family, including a more precise 'time of origin' in both time and space for the derived lineages, Eocene to Pliocene sediments in other parts of Africa as well as middle Miocene to Pleistocene sediments in South Africa must be screened for Loranthaceae pollen and studied using SEM. Upper Cretaceous to Paleocene sediments (e.g. McLachlan & Pieterse 1978; Partridge 1978; Scholtz 1985; Sandersen et al. 2011) should also be screened for Loranthaceae-type pollen to establish if South American/Australasian basal lineages dispersed into Africa via a southern route, but went extinct during the early Cainozoic.

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Supplemental data

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