PITYOSTROBUS ANDRAEI (PINACEAE) FROM THE BARREMIAN (LOWER CRETACEOUS) OF BELGIUM: A MORPHOMETRIC REVISION

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Important floristic changes took place during the Early Cretaceous (145.0–100.5 Ma). They are notably marked by a peak in conifer diversity, especially within Pinaceae. This diversification is attested to by the numerous ovulate cone taxa found in the western European and North American fossil records. Exceptional deposits in the Wealden facies (Barremian-Albian, 125.0–100.5 Ma) of western Belgium (La Louvière, Houdeng-Aimeries) have yielded hundreds of exceptionally well-preserved pinaceous ovulate cones, yet most of these remain unstudied. These are presently revised. We investigate the morphological variation of the most abundant species, *Pityostrobus andraei* (Coemans) Seward 1919, with a morphometric approach. This work applies statistical tests to a large sample (n = 132 fossil cones) for the first time, using 10 measurements to characterize the morphology of the cones. Our analyses recover eight distinct morphotypes included in the original specific concept of *P. andraei*, and most of the specimens belong to two main morphotypes that represent 50% and 25% of the total specimes analyzed. Our results suggest that the specific diversity of early Pinaceae is underestimated, as it is based on older species concepts. Modern quantitative approaches have the potential to better characterize the breadth and tempo of pinaceous diversification.

Keywords: Wealden, conifers, disparity, cone, morphometry, diversity.

Online enhancements: appendixes.

Introduction

During most of the Mesozoic era (252.6–66.0 Ma), continental ecosystems were dominated by Cycadales, Bennettitales, and ferns. The Lower Cretaceous (145.0–100.5 Ma) was, however, marked by important floristic changes (Watkins and Cardelús 2012). Conifers that appeared in the Pennsylvanian (323.2– 289.9 Ma) were expanding rapidly, increasingly colonizing the Northern Hemisphere continents, especially western Europe and North America (Bonnot et al. 1963; Miller 1977; Taylor et al. 2009; Smith et al. 2017). Some modern families, such as the Cupressaceae and Araucariaceae, were already present at the end of the Jurassic (Taylor et al. 2009; Herrera et al. 2017).

Among living conifers, Pinaceae have a wide ecological range, thriving in continental, mountainous, or subarctic environments (Bonnot et al. 1963; Critchfield and Little 1966; Smith et al. 2017). With 11 genera and about 230 species of trees and shrubs, it is the most diverse conifer family at the species level (Falcon-Lang et al. 2016; Gernandt et al. 2016). In the fossil record, Pinaceae are first found at the end of the Jurassic (*Eathiestrobus mackenziei*; Rothwell et al. 2012) and at the beginning of the Cretaceous (Valanginian, 140–133 Ma; Klymiuk and Stockey

2012; Falcon-Lang et al. 2016). Cones that are morphologically close to current pinaceous cones classified in modern genera such as *Pinus* (found in Belgium and England) appeared as soon as the Early Cretaceous (Alvin 1953, 1960; Ryberg et al. 2012; Falcon-Lang et al. 2016; Smith et al. 2017). The Pinaceae reached their maximal morphological disparity at about the same time (Oyston et al. 2016).

Our understanding of the early evolution and diversification patterns of Pinaceae is mainly based on isolated ovulate cones found in the Cretaceous and the Paleogene of Eurasia and North America (Gernandt et al. 2016). Consequently, as of now there are no studies that have quantitatively examined the inter- and intraspecific variability of these fossil cones. Taxonomic resolution for the form genus Pitvostrobus, erected to encompass isolated pinaceous ovulate cones, is thus likely to be improved by further analyses intended to survey morphological data (Miller 1976; Ryberg et al. 2012). Weakly phylogenetically supported and with no clear diagnosis, Pityostrobus certainly does not form a clade (Smith et al. 2017). It nevertheless represents the most abundant Pinaceae form genus in the Lower Cretaceous, with 28 species of 42 pinaceous cones known for that interval (i.e., two-thirds of the species). Because of the large number of species included in it, this genus is responsible for major biases in phylogenies (Gernandt et al. 2016; Smith et al. 2017).

Descriptions of seed cone morphotaxa are subject to several inherent identification biases: quality of preservation, number of specimens available, and subjectivity of identification, depending

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on the author. From a taxonomic point of view, *Pityostrobus* is a "dustbin" genus that encompasses multiple different species concepts. A standardization of the genus is needed to clarify the identifications of Pinaceae morphospecies and to develop reproducible methods of analysis for fossil ovulate cones. To this end, we tested the combination of descriptions and quantitative analyses presented here.

The current research was triggered by the reinvestigation of the Wealden collections housed at the Royal Belgian Institute of Natural Sciences (RBINS; Brussels, Belgium). At the end of the nineteenth century, several excavations in the Belgian Wealden (Lower Cretaceous) continental facies yielded hundreds of exceptionally well-preserved ovulate cones. Nine species have been described based on this material, and most of them have been included in comparative and phylogenetic studies of fossil and extant Pinaceae on multiple occasions (Smith and Stockey 2002; Gernandt et al. 2008, 2018; Smith et al. 2017). This unique collection is remarkable for the number of specimens of each species that were collected, which provides an opportunity to explore quantitatively the variability of individual characters and discuss the validity of fossil species.

This report represents the first part of the revision of this entire collection from the Belgian Lower Cretaceous. Here we focus on one of the most abundant *Pityostrobus* species found in this collection: *Pityostrobus andraei* (Coemans) Seward 1919. Its intraspecific morphological disparity has been reassessed, and it has been compared with other *Pityostrobus* species. As a result, eight different morphotypes are recognized.

Material and Methods

Localities

Studied fossils were collected from two closely located Belgian localities: La Louvière and Houdeng-Aimeries (fig. 1). The first



Fig. 1 Geographical map showing the localities of La Louvière and Houdeng-Aimeries (Roeulx 1 [R1] and Roeulx 5 [R5]), Belgium.

collection was built by Coemans (1867) from a now-abandoned clay quarry in the former hamlet of Baume in La Louvière. The collection was further increased by C. Bommer, who in 1892 collected abundant plant specimens during the construction of floodgate 3 of the "canal du centre" (Bommer 1892; Alvin 1953). These localities are currently not accessible.

Both La Louvière and Houdeng-Aimeries are situated at the eastern limit of the Mons Basin. This basin is part of the Anglo-Paris Basin, representing its northeast rim. The Mons Basin constitutes a 40-km-long and 10-km-wide subsiding zone that accumulated 300 m of Meso-Cenozoic sediments (Yans 2007; Yans et al. 2010). The Wealden facies is exposed as "pockets" at the north of the Mons Basin. They correspond to Cretaceous continental siliciclastic sediments deposited in deltaic complexes. Highly diachronic across the basin, they range in age from middle Barremian (127.3 Ma) in the west to middle-late Albian (100.5 Ma) in the east (Yans et al. 2006; Dejax et al. 2007, 2008; Robaszynski et al. 2007; Gernandt et al. 2008). Although this west-east Barremian-Albian age gradient is supported by some data, several eastern localities of the Wealden facies-Soignies, Houdeng-Aimeries (Roeulx 5), and La Louvière (Roeulx 1)-have never been precisely dated.

Material

All ovulate cones were found isolated in a sandy clay matrix in association with leaves, twigs, and wood fragments, which were not collected (Bommer 1892). They were mechanically extracted and were not chemically prepared. The cones are lignitized (Alvin 1953, 1971, 1974) and preserve most of their original tridimensional shape, being only slightly compressed. Anatomy and tissues are well preserved (Alvin 1953).

In Belgium, the Wealden facies occurs in five formations: the Hautrage, Baudour, and Sainte-Barbe Formations to the west and the Saint Pierre and Strepy Formations to the east. The layers of La Louvière and Houdeng-Aimeries from which the ovulate cones were collected belong to the Saint-Pierre Formation. Although the two localities were not precisely dated, the Saint-Pierre Formation is dated to the upper half of the Albian (Yans 2007). Precise dating of these localities will be useful for comparisons among the different fossil assemblages of the Mons Basin.

Sampling

In this study, a sample of 295 complete or incomplete fossil ovulate cones hosted in the RBINS collections was observed. This sampling represents all cones attributed to the species *Pityostrobus andraei* available in the collections. The cone specimens were initially identified as *Pinus andraei* Coemans 1867 by C. Bommer (1892). They were subsequently reinterpreted as *Pityostrobus andraei* (Coemans) Seward 1919 by Alvin (1953). A subsample of 124 of these cones was selected for morphometric analyses because they were either complete or in good condition. The remaining 171 specimens are too fragmented or poorly preserved to be confidently included in one morphotype and to allow for morphometric analysis, as their incompleteness might distort the morphospaces. In addition, eight specimens representing other species of Belgian Pinaceae were incorporated in our analyses to calibrate the amount of morphological variation within *P. andraei* as currently defined. These were *Pityostrobus bommeri* Alvin 1953 (n = 2), *Pityostrobus hautrageanus* Alvin 1960 (n = 1), *Pityostrobus soigniesiensis* Alvin 1960 (n = 1), *Pityostrobus villerotensis* Alvin 1957 (n = 2), *Pseudoaraucaria gibbosa* Alvin 1960 (n = 1), and *Pseudoaraucaria heeri* Alvin 1957 (n = 1).

Imaging and Measurements

First, the selected ovulate cones were classified into several morphotypes based on their general shape and anatomy (see descriptions below). All the specimens selected (n = 132) were individually photographed with a Nikon D90 camera equipped with an AF-S DX NIKKOR 18-105mm f/3.5-5.6G ED VR. The photographs were used for bidimensional analysis using linear morphometry. Ten measurements were taken to characterize cone shape, following the protocols of Gil et al. (2002) and Marcysiak (2004), as follows: (1) cone length, (2) width of the cone base, (3) maximal cone width, (4) cone diameter midway between the base and the maximum diameter, (5) width of the scale apophysis in the middle of the cone, (6) height of the scale apophysis in the middle of the cone, (7) angle of the apophysis, (8) length of the cone base (portion that bears reduced scales), (9) number of scales, and (10) number of gyres along an orthostichy (fig. 2; app. A1; apps. A1-A5 are available online). These last two measurements (9 and 10) are subject to a preservation bias because the base and/or the apex was missing on many cones. Therefore, the number of scales and gyres was, whenever possible, extrapolated by means of the scale bases when they were still present or by doubling the number of scales counted on one side of the cone.

Measurements and observations were performed digitally with ImageJ/Fiji 1.52n software (Schindelin et al. 2012; Schneider et al. 2012). For tridimensional observations and anatomical measurements, two cones were scanned at the micro-computed tomography (CT) facility of the RBINS using an EasyTom 150-kV scanner (RX Solutions, 3D X-Ray tomography systems) fitted with a microfocus X-ray source and a 320 × 420-mm detector. Specimens Pbot-0-0231 and IRSNB_b_6840 of the RBINS collections (app. A1) were scanned in their entirety using a copper filter. Pbot-0-0231 was scanned at a spatial resolution of 0.019 mm, with an electron acceleration energy of 149 kV and a current of 67 μ A. IRSNB_b_6840 was scanned at a spatial resolution of 0.026 mm, with an electron acceleration energy of 143 kV and a current of 70 μ A. Scans were used for anatomical descriptions and measurements that were also performed digitally with the ImageI2 software (Schneider et al. 2012; Rueden et al. 2017). Scans can be consulted on request to the RBINS curator of Palaeontology Collections.

Statistical Analyses

To characterize the morphological variability of *P. andraei*, univariate and multivariate tests were applied to the measurements within the R 3.6.0 environment (R Core Team 2019) using the data matrix obtained with the measurements of the bidimensional analysis (fig. 2). For the univariate analysis, the R package readxl version 1.3.1 was used to import the data (Wickham and Bryan 2019). To compare the different morphotypes described, the variables measured are compared with each other to determine whether populations are significantly different.



Fig. 2 Illustration of the eight measurements applied to the ovulate cones, as follows (following Gil et al. 2002; Marcysiak 2004): (1) cone length, (2) length of the cone base, (3) maximal cone width, (4) cone diameter midway between the base and the maximum diameter, (5) width of the scale apophysis in the middle of the cone, (6) height of the scale apophysis in the middle of the cone, (7) angle of the apophysis, and (8) base length (at the bottom of the cone, where the scales are reduced). s = scale; g = gyre.

We tested the normality of selected variables (1, 3, 5, 6, 9, 10; fig. 2) using the Shapiro-Wilk test (Shapiro and Wilk 1965). A Fisher test was applied to test homoscedasticity (Fisher 2006). For testing homogeneity between means of different variables, a parametric Student's *t*-test was performed on data that follow a normal distribution; the nonparametric Wilcoxon test (Mann-Whitney *U*-test) was performed on the others (Wilcoxon 1945; Mann and Whitney 1947). To compare means for variables with homogeneous variances, we used a permutation Student's *t*-test with the R package RVAideMemoire version 0.9-75 (Hervé 2019; app. A2).

Multivariate analysis was performed to characterize the morphospace occupancy of the samples and to circumscribe the different cone morphotypes. The R packages readxl version 1.3.1, ape version 5.3, devtools version 2.3.0, ggplot2 version 3.3.0, and ggbiplot version 0.55 were used (Vu 2011; Wickham 2016; Paradis and Schliep 2019; Wickham and Bryan 2019; Wickham et al. 2020). The data set was scaled (z transformed), and all individuals with missing data were removed, which produced a matrix of 81 specimens and 12 groups (app. A3). Principal component analysis (PCA) was applied to the resulting data set to circumscribe morphospace occupation. Only seven of the measurements (cone length [1], maximal cone width [3], cone diameter midway between the base and the maximum diameter [4], width of the scale apophysis in the middle of the cone [5], height of the scale apophysis in the middle of the cone [6], scale number [9], and number of gyres along an orthostichy [10]) are included in the PCA because the other measurements have too many missing data to be correctly represented. Morphotypes C, F, and I and the Belgian species P. gibbosa are not represented because they are each represented by only one specimen that is missing some data. Normal data ellipses with a size of 0.68 are used to highlight morphospaces for morphotypes with more than two specimens. A multivariate analysis of variance (MANOVA) was computed to test the differences between individuals of morphotypes A and B included in the PCA (Bray et al. 1985). Other morphotypes were removed from the analysis, as they do not have enough specimens.

Results

Definition of Morphotypes

Description of the 124 selected cones resulted in the separation of eight morphotypes (fig. 3; table 1). Those specimens either were initially figured by Alvin (1953) as *Pityostrobus andraei* (A, B, G, H) or were classified as *P. andraei* by Alvin in the RBINS collections (C–F). We figured the most representative and relevant cones for each morphotype (fig. 3). Of these specimens, 50% and 25% belong to two morphotypes, referred to as A (n = 66) and B (n = 30), respectively. Seven percent belong to six other morphotypes, referred to as C–H (n = 9; fig. 3; table 1). The remaining 15% of the specimens were not assigned to any morphotype because they were too abraded (n = 19; app. A1).

All morphotypes are lignified ovulate cones with bract-scale complexes that are helically arranged. The scales bear two inverted seeds at their base. Seeds are winged.

Morphotype A. Morphotype A is represented by lignified ovulate cones that are at least 2.8–7.4 cm long and 1.3–3.2 cm in diameter and ovoid (n = 66; 50%; figs. 3A, 4A–4C; IRSNB_b_6844). Bract-scale complexes are helically arranged. The ovuliferous scales are 12–20 mm long and 2.6–8 mm wide, oriented at an angle of 20° to the stele axis. Pith is 0.8–1.6 mm in diameter. The vascular cylinder is 150 µm in diameter, slightly dissected, with rare resin canals. Cortex is 0.85–1.4 mm in diameter, with one to five resin canals. Vascular traces to the ovuliferous scale and bract diverge separately from the cone axis; the ovuliferous scale trace is formed from one vascular strand, abaxially concave. Bracts are reduced, 10%–20% of scales' length. There are two seeds per scale, winged, measuring 3.3 mm in length and 1 mm in thickness. Wings measure

two-thirds of the scale length. Sarcotesta and sclerotesta together measure 0.13–0.2 mm.

Morphotype B. Morphotype B is represented by lignified ovulate cones that are 7.2–14 cm long and 1.6–3.0 cm in diameter and elongated and cylindrical (n = 30; 25%; figs. 3B, 4D–4F; IRSNB_b_6840). Bract-scale complexes are helically arranged. The ovuliferous scales are 20 mm long and 6.8–8.5 mm wide, oriented at an angle of 14° to the stele axis. Pith is 1.8–2.1 mm in diameter; resin canals are absent. The vascular cylinder is continuous, slightly dissected at the scale bases, and 0.3–0.4 mm in thickness. Cortex is 0.9–2 mm in diameter, with 15 resin canals. Scale trace is flat at its divergence from the axis and then becomes cylindrical. Apophysis is rhomboid in shape with a terminal umbo. Seeds are ovate in shape and measure, on average, 4.5 mm in length and 1.5 mm in thickness. Sarcotesta and sclerotesta together are 0.15–0.19 mm wide.

Morphotype C. Morphotype C is represented by a lignified ovulate cone that is 3.3 cm long and 1.8 cm wide and ovoid cylindrical in shape (n = 1; 0.8%; fig. 3C; Pbot-06058-0117). Bract-scale complexes are helically arranged. Ovuliferous scales are 12 mm long and 10 mm wide. Apophysis is flabellate, radially striated, thin, without an umbo. Apophyses are 2.4 mm long and 10 mm in diameter. Scales have two inverted seeds at their base. Seeds are winged and are 2–3.2 mm long and 1.6–1.8 mm in thickness. The sarcotesta and sclerotesta are 0.59 mm thick.

Morphotype D. Morphotype D is represented by lignified ovulate cones that are 4.2–4.3 cm long and 2–2.2 cm wide and ovoid in shape (n = 2; 1.6%; fig. 3D; Pbot-06058-92). Bract-scale complexes are helically arranged. Ovuliferous scales are 14–15 mm long, 6–9 mm wide, and 0.3–0.8 mm thick. Scales are thickened into a rhomboid apophysis and have no umbo. Apophyses are 7.8–9.4 mm long and 5.4–5.8 mm high. Scales have two inverted seeds at their base. Seeds are winged and are 5–6 mm long and 8–9 mm in thickness. The sarcotesta and sclerotesta are 0.16–0.21 mm thick.

Morphotype E. Morphotype E is represented by lignified ovulate cones that are 4.4 cm long and 1.8–2 cm in diameter and elongated and cylindrical in shape (n = 3; 2.4%; fig. 3*E*; Pbot-0-0232). Bract-scale complexes are helically arranged. Scales are thickened into an apophysis, rhomboid and mucronate. Apophyses are 6–7 mm long and 8–9 mm wide. A terminal umbo is present and follows the shape of the apophysis.

Morphotype F. Morphotype F is represented by a lignified ovulate cone that is 6 cm long and 3.1 cm wide and ovoid and cylindrical in shape (n = 1; 0.8%; fig. 3F; Pbot-0-0160). Bract-scale complexes are helically arranged. Ovuliferous scales are 20 mm long and 8–10 mm wide. Reduced bracts are visible. Scales are distally thickened and slightly folded abaxially, forming a rhomboid-mucronate apophysis. Apophyses are 6.3 mm long, 10 mm wide, and 1.5 mm thick. Scales have two inverted seeds at their base. Seeds are winged. The sarcotesta and sclerotesta are 0.16 mm thick.

Morphotype G. Morphotype G is represented by a lignified ovulate cone that is 5.9 cm long and 2.1 cm wide and cylindrical in shape (n = 1; 0.8%; fig. 3G; IRSNB_b_6855). Bract-scale complexes are helically arranged. Ovuliferous scales are 16 mm long, 6–9 mm wide, and 0.5–1 mm thick. Scales are distally thick-ened, forming a rounded apophysis. Apophyses are 7.7 mm long and 11 mm wide and have a terminal umbo folded abaxially. Scales have two inverted seeds at their base. Seeds are winged



Fig. 3 Eight morphotypes separated within the sample (n = 124) of *Pityostrobus andraei* species. A, Morphotype A, IRSNB_b_6844; 50%. B, Morphotype B, IRSNB_b_6840; 25%. C, Morphotype C, Pbot-06058-0117; interpreted as close to *Pityostrobus villerotensis*. D, Morphotype D, Pbot-06058-92. E, Morphotype E, Pbot-0-0232; submature stage of morphotype B. F, Morphotype F, Pbot-0-0160. G, Morphotype G, IRSNB_b_6855. H, Morphotype H, IRSNB_b_6843. Scale bars = 3 cm. u = umbo.

and are 4.8 mm long and 1.9–2.2 mm in diameter. The sarcotesta and sclerotesta are 0.1 mm thick.

Morphotype H. Morphotype H is represented by a lignified ovulate cone that is 5.5 cm long and 2 cm wide and cylindrical

in shape (n = 1; 0.8%; fig. 3*H*; IRSNB_b_6843). Bract-scale complexes are helically arranged. Scales are distally thickened, forming an acute apophysis. Apophyses are 9 mm long and 9.2 mm wide and have a terminal umbo that is slightly folded

			General features			Cone axis			
Morphotype	e n	Sample percentage (%)	Cone shape	Cone length (mm)	Cone width (mm)	Pith diameter (mm)	Vascular cylinder diameter (mm)	Cor diam (mi	tex eter n)
А	66	50	Ovoid	28-74	13-32	.8-1.6	.15	.85-1.4	
В	30	25	Elongated and cylindrical	72–140	16-30	1.8–2.1	.3–.4	.9–2	
С	1	.8	Ovoid and cylindrical	33	18	na	na	na	
D	2	1.6	Ovoid	42	20	1.7-2	na	na	
E	3	2.4	Elongated and cylindrical	44	18–20	na	na	na	
F	1	.8	Ovoid and cylindrical	60	31	na	na	na	
G	1	.8	Cylindrical	59	21	na	na	na	
Н	1	.8	Cylindrical	55	20	na	na	na	
-	Ovuliferous scales				Apophysis		Seeds		
	Resin canals in the cortex	Length (mm)	Width (mm)	Umbo absence/presence and position	Length (mm)	Width (mm)	Sarcotesta and sclerotesta thickness (mm)	Length (mm)	Width (mm)
А	1-5	12-20	2.6-8	Not present	3.6-10	5-13	.132	3.3	1
В	15	20	6.8-8.5	Terminal	8.4	7.6	.1519	4.5	1.5
С	na	na	10	Not present	2.4	10	.59	2-3.2	1.6-1.8
D	na	na	6–9	Not present	7.8-9.4	5.4-5.8	.1621	5-6	8-9
E	na	na	na	Terminal	6-7	8–9	na	na	na
F	na	20	8-10	Not present	6.3	10	.16	na	na
G	na	16	6–9	Terminal	7.7	11	.1	4.8	1.9–2.2
Н	na	na	9.2	Terminal	9	9.2	na	na	na

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Comparison of the Main Characteristics of the Eight Morphotypes

Note. The table is based on 586 measurements in millimeters. na = not applicable.

abaxially. Scales have two inverted seeds at their base. Seeds are winged.

Morphometric Analyses

Our data set scores 10 variables for 132 specimens. Although the data set has a fairly low proportion of missing data (32%; app. A1), the completeness threshold for our multivariate analyses required a data set reduction to seven variables for 81 specimens (0% missing data).

Because of this, only morphotypes A and B are amenable to statistical comparison. We thus here focus on the characterization and the comparison of their respective variabilities. Morphotypes C–H are excluded from these analyses as they are represented by only one to three specimens each (0.8%–2.4% of the sample). Half of these specimens have missing data because of distortion or poor preservation.

Univariate Analysis

The results of this analysis are presented in five boxplots comparing the cones in terms of length (variable 1), the cone length-to-width ratio (1, 2), the apophysis shape ratio (width-to-height; 5, 6), the number of scales (9), and the number of gyres (10), along with the associated *P* values (fig. 5; app. A4).

These characters were chosen because they are the most representative variables describing cone morphology with the least missing data.

Continuous variables measured for morphotypes A and B are distributed on significantly different intervals (see *P* values; fig. 5). Cone length differs significantly between morphotypes A and B (fig. 5.1; Student's *t*-test: P < 0.0001). The cone length-to-width ratio is also significantly different between the two morphotypes (fig. 5.2; Student's *t*-test: P < 0.0001). The difference in apophysis size is also significant (fig. 5.3; Student's *t*-test: P < 0.0001). The apophysis shape ratio of morphotype A is much more variable than that of morphotype B (from 0.89 to 2.05 vs. from 0.83 to 1.5, respectively; P < 0.01; app. A2). The number of scales (fig. 5.4; permutation Student's *t*-test: P = 0.002) and number of gyres (fig. 5.5; Wilcoxon test: P < 0.0001) are significantly different as well: morphotype B has more scales and gyres than morphotype A.

Morphotypes C–H were not included in this univariate analysis. The low number of individuals did not allow us to perform statistical tests. However, the visualization of the individuals in the boxplots allowed us to see that morphotypes D–H have values close to those of morphotypes A and B (app. A5). Only morphotype C shows an extreme value for the apophysis shape ratio. This is explained by its scales, which are very

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Fig. 4 Internal views of morphotype A (*A*–*C*; Pbot-0-0231; spatial resolution of 0.019 mm, 149 kV, 67 μ A) and morphotype B (*D*–*F*; IRSNB_b_6840; spatial resolution of 0.026 mm, 143 kV, 70 μ A). *A*, *B*, *E*, and *F* are transverse sections; *C* and *D* are longitudinal sections. *A*, Vascular cylinder with rare resin canals, cortex with one to five resin canals, and two seeds per scale. *B*, Slightly dissected vascular cylinder; the ovuliferous scale trace is formed from one vascular strand. *C*, Ovuliferous scale is oriented at an angle of 20° to the stele axis, vascular traces to the ovuliferous scale and bract diverge separately from the cone axis, and bracts are reduced, 10%–20% of scales' length. *D*, Ovuliferous scales are oriented at an angle of 14° to the stele axis. *E*, Resin canals are absent from the pith, the cortex has 15 resin canals, and the scale trace is flat on derivation of the axis and then becomes cylindrical. *F*, Vascular cylinder is continuous and slightly dissected at the scale bases. Scale bars = 1 cm. a = axis; ap = apophysis; b = bract; bsc = bract-scale complex; c = cortex; ir = interseminal ridge; it = internal tissue; m = micropyle; p = pith; rc = resin canal; s = seed; sc = scale; sct = scale trace; ss = sarcotesta and sclerotesta; u = umbo; vc = vascular cylinder; vs = vascular strand; vtb = bract vascular trace; vts = scale vascular trace; w = wing; xd = xylem discontinuity.

large and flat compared with those of the other morphotypes (fig. 3).

Multivariate Analysis

The measurements were analyzed by means of a PCA to observe the distribution of the different morphotypes in different morphospaces (fig. 6; app. A4). Morphotypes A and B are included in the analysis. The MANOVA shows that the specimens of morphotypes A and B included in the multivariate analysis are significantly different (P < 0.0001).

Components 1 and 2 represent 53.2% and 19.8% of the total measured variance, respectively. Component 1 is mainly influenced by the length and the middiameter (-0.42 and 0.4). respectively). After that, apophysis width (-0.375), apophysis length (-0.385), apophysis height (-0.345), number of scales (-0.381), and number of gyres (-0.332) all equally influence component 1. Component 2 is mostly influenced by the number of gyres and cone width (0.585 and -0.457, respectively). Morphotypes A and B occupy two distinct morphospace regions that are only slightly overlapping. The specimens attributed to morphotype A are more dispersed than those attributed to morphotype B. Morphotypes C and F are not represented in the PCA because they were removed in response to the completeness threshold. Morphotypes E and H are located in the same morphospace region as morphotype B, while morphotypes D, G, and H are located in the same region as morphotype A. The Pseudoaraucaria heeri specimen is also located in the same

morphospace as morphotype A. The specimens of *Pityostrobus* bommeri, *Pityostrobus hautrageanus*, *Pityostrobus soigniesiensis*, and *Pityostrobus villerotensis* are isolated apart from the morphotype A and B morphospaces. The nine unidentified specimens are all located within or in proximity to the morphospace of morphotype A.

Discussion

Species Delimitation in Pityostrobus

Our results allow us to reconsider the species concept of *Pityostrobus andraei* in light of the amount of morphological disparity that it encompasses. When Coemans (1867) and then Alvin (1953) first described the specimens, they assigned the 303 specimens of the RBINS collections to *Pinus andraei* Coemans 1866 and then to *Pityostrobus andraei* (Coemans) Seward 1919, respectively. Among these specimens, we were able to differentiate eight morphotypes, of which morphotypes A and B are the most abundant.

The results of the univariate statistical analysis strongly support the hypothesis that morphotypes A and B (fig. 5) represent two distinct species. The use of continuous characters allows us to significantly differentiate the two morphotypes (see P values; fig. 5). Measurements were made of the external morphology of the cones (fig. 2; app. A1), although these external features are not clearly defined as diagnostic at the species level in existing descriptions (Miller 1976). Here, they allow us to quantify and to render the descriptions mutually consistent. Cone



Fig. 5 Boxplots representing morphometric comparison between morphotypes A and B, including specimens with missing data. Morphotype A: n = 66; morphotype B: n = 30. 1, Cone length- 2, Cone length-to-width ratio. 3, Apophysis shape ratio. 4, Number of scales. 5, Number of gyres.

length and width, apophysis width and height, and number of scales and gyres are all significantly different between the two morphotypes (fig. 5).

Morphotype C looks very similar to the species *Pityostrobus villerotensis* in ovoid-cylindrical shape; thin, flabellate, radially striated apophysis; lack of an umbo; and the sclerotesta and sarcotesta measuring 0.59 mm in diameter. However, *P. villerotensis* was found in a different locality (Villerot) than the cone included in our analyses; Villerot, like the La Louvière and Houdeng-Aimeries localities, has not been dated yet (fig. 1). On the other hand, the Villerot locality is located farther west in the basin and should be older than La Louvière (Yans et al. 2010). Morphotype C could be younger than but morphologically close to *P. villerotensis*.

The PCA shows that the combination of cone measurements that we used enables the assessment of intraspecific variation and clearly shows that morphotypes A and B occupy different morphological spaces. There is no gradual transition from the space occupied by one of the morphospaces to the other, and the overlap between the two morphospaces is minute.

Although morphotypes A and B both include ovulate cones with scales that thicken into an apophysis at the apex and bear two inverted seeds adaxially at their base, figures 5 and 6 highlight the many differences between these two morphotypes. Additional differences can be listed as follows: (1) the bract, which is clearly visible in morphotype A, is not visible in morphotype B, (2) the scale base is thinner in morphotype A, (3) the secondary xylem of the axis is more dissected in morphotype A, (4) the shape of the scale trace is flat in morphotype B and abaxially concave in morphotype A, and (5) cortical resin canals are more numerous in morphotype B.

Coemans (1867) and Alvin (1953) were both aware of its great variability when they described *P. andraei*. Alvin (1953) justified the inclusion of the morphotypes defined here in the same species based on their consistent association in the two localities. As some cones were a little abraded on the surface, he interpreted those that we include in morphotype A as representing either an immature stage or very abraded specimens of *P. andraei* (morphotype B). Here, we show that although some cones of morphotype A were abraded, most of them are well preserved (both internal and external structures). Furthermore, both morphotypes comprise mature or submature specimens, as they are completely lignified, have reduced bracts, and bear mature seeds.

In our opinion, the results of the morphometric analysis, combined with the other morphological and anatomical differences that we highlighted above (cone shape, scale shape, umbo and bract presence, and scale trace shape; see table 1), should be



Fig. 6 Morphospace occupation using the first two axes of the principal components (PC) analysis (n = 81) with superimposed 68% confidence ellipses for each group. Multivariate analysis of variance (var.), P < 0.0001. Length; width; mid_diameter = cone diameter midway between the base and the maximum diameter; l_apophysis = width of the scale apophysis; h_apophysis = height of the scale apophysis; scale_nb = scale number; gyres_nb = number of gyres along an orthostichy. Groups: A = morphotype A (n = 48); B = morphotype B (n = 14); b = *Pityostrobus bommeri* (n = 1); D = morphotype D (n = 1); E = morphotype E (n = 1); G = morphotype G (n = 1); H = morphotype H (n = 1); Ph = *Pityostrobus hautrageanus* (n = 1); Psh = *Pseudoaraucaria heeri* (n = 1); s = *Pityostrobus soigniesiensis* (n = 1); v = *Pityostrobus villerotensis* (n = 2); ? = undetermined morphotype (n = 9).

enough to justify splitting *P. andraei* into at least two distinct species (corresponding to our morphotypes A and B). To redefine the species concept of *P. andraei* in the absence of a clearly defined holotype, isotype, isosyntype, or syntype, a lectotype must be selected from among the illustrated specimens (Turland et al. 2018, chap. II, art. 9.3). This formal taxonomic decision will be the subject of a separate upcoming article. As these Belgian species are included in recent studies on the evolution of Pinaceae (Gernandt et al. 2016; Smith et al. 2017), these new systematic data will probably bring new information to the phylogenies.

Morphometry-Informed Taxonomic Diversity among Fossil Cones

Pinaceous cones are structures that are rarely preserved compared with the vegetative parts of conifers. They are also more likely to disarticulate before burial. This explains why most Cretaceous ovulate cone descriptions are based on isolated or incomplete specimens (Gernandt et al. 2016; table 1). It is also common for external features of the cone to be difficult to observe (e.g., cones encased in nodules; Falder et al. 1998; Rothwell et al. 2012). This is compounded by the fact that many cone morphospecies were described more than 50 years ago (Alvin 1953, 1957, 1960), when species concepts were somewhat different from those employed nowadays. As a result, the taxonomy of fossil Pinaceae is at times inconsistent (Miller 1976; Ryberg et al. 2012; Smith et al. 2017), and this distorts our understanding of the assembly of their diversity throughout the Mesozoic and the Cenozoic. Yet conifer reproductive structures are an ideal organ to study in a comparative framework because they possess many diagnostic characters, which are commonly employed to define fossil and extant species (Farjon 2005).

Our approach is aimed at homogenizing descriptions and taxonomy within a thorough comparative framework and constitutes a first step in the use of morphometrics to study the diversity of pinaceous cones. This type of approach was already tested on extant Pinaceae species in three studies on cone morphology (Gil et al. 2002; Marcysiak 2004; Vetrova 2013). On the other hand, studies of fossil species have already been conducted with other plant groups using morphometric data (Roth-Nebelsick et al. 2000; Benca et al. 2014) but not with Pinaceae species.

The next step will be to apply this methodology to all the different morphotaxa of Belgian fossil ovulate cones from the Cretaceous that are represented by enough specimens to assess their intra- and interspecific variability (Herrera et al. 2015, 2017). Eventually, this will provide important data to complement the studies already carried out on the disparity of conifer cones (Leslie 2011; Smith et al. 2017). The analysis of continuous characters opens the way for accurate disparity studies through the use of linear and geometric morphometrics, which has rarely been used, up until now, with plants (Oyston et al. 2016).

Conclusions

Descriptions of pinaceous morphotaxa are subject to several inherent biases: quality of preservation, number of fossilized specimens, and subjectivity of identification, depending on the author. A combination of descriptions and quantitative analyses is applied to homogenize the identifications. The aim is to establish a methodology for reproducible analyses of individuals of the same family.

By applying morphometrics to a large sample of ovulate cones (n = 132), we were able to clearly distinguish two different morphotypes included in the species *Pityostrobus andraei* (A and B). The use of descriptions combined with statistical analyses demonstrates that morphotypes A and B belong to two distinct populations. We quantify the cone morphology (length, width,

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apophysis shape, number of cone scales and gyres). Statistical analyses demonstrate that these features are significantly different between morphotypes A and B (fig. 5). The PCA shows that the combination of measurements of multiple features of the cone morphology enables the assessment of interspecies variation. The PCA clearly shows that morphotypes A and B occupy different morphospace regions. The main differences are cone shape, scale shape, the length of the bract in morphotype A, the number of resin canals in the cortex, and the sclerotesta and sarcotesta thickness.

This work shows that the specific diversity of the ovulate cones found in the Belgian Lower Cretaceous (Wealden facies) needs to be reevaluated, as the species concept of *P. andraei* initially described by Alvin (1953) includes clearly distinct morphotypes that should be elevated to morphospecies. These morphotypes make it possible to reassess the number of known species in Belgium and consequently Pinaceae diversity during the Early Cretaceous.

Future work should clarify systematics problems for fossil species currently classified within the genera *Pinus* and *Pityostrobus*. Combining this with a review of the Belgian fossil species will allow further characterization of the diversity of the Pinaceae during a key period of their evolution.

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