

Grass pollen surface ornamentation: a review of morphotypes and taxonomic utility

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Abstract: The classification of grass pollen is a classic problem in palynology. One approach to this problem is to image grass pollen using scanning electron microscopy (SEM) and to construct morphotypes on the basis of surface ornamentation patterns. In this paper, we construct a database of published SEM images of grass pollen surface ornamentation and use this database to review the status and utility of grass pollen morphotypes. Very little SEM data on grass pollen surface ornamentation have been collected. There are more SEM data on the pollen of species from larger subfamilies such as the Pooideae, but there are no SEM data on the pollen of relatively small subfamilies or the early-diverging APP clade of grasses. The available data support the following six morphotypes: the *Hordeum*-type, *Triticum*-type, *Avena*-type, *Setaria*-type, *Pariana*-type and *Stipa*-type. The phylogenetic distribution of these morphotypes cannot be assessed reliably with the available data, but the *Pariana* and *Stipa* types are apparently restricted to the BEP clade. We suggest that fossil grass pollen grains could be classified using these morphotypes and that this could be a means of generating useful evolutionary and palaeoecological information from the grass pollen fossil record.

Keywords: Poaceae; grass; pollen; phylogeny; taxonomy

Supplementary material: Our database of grass pollen surface ornamentation is available at <https://doi.org/10.6084/m9.figshare.c.2134356>

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Grass pollen is remarkably similar throughout Poaceae (the grass family) and is spheroidal with a single pore that is usually, but not always (Skvarla *et al.* 2003), surrounded by an annulus. When viewed using transmitted light microscopy with brightfield illumination, which is the standard microscopy technique in palynology (Sivaguru *et al.* 2012), grass pollen grains have a psilate to faint scabrate surface ornamentation (Wodehouse 1935; Fægri *et al.* 1992; Beug 2004). When viewed using scanning electron microscopy (SEM), however, grass pollen grains are characterized by diverse and intricate surface ornamentation patterns (see Mander & Punyasena (2014, p. 939) for a comparison of light microscopy and SEM images of grass pollen).

Palynologists have made several attempts to use surface ornamentation patterns to classify grass pollen. These include recent efforts to quantify the complexity of these patterns using computational image analysis (Mander *et al.* 2013), and the development of grass pollen morphotypes that describe grass pollen surface ornamentation using descriptive terminology (e.g. Köhler & Lange 1979). However, attempts to classify grass pollen grains using surface ornamentation patterns have been blighted by concerns about the degree to which these patterns reflect grass phylogeny. For example, Kellogg (2015) has pointed out that the morphological groups produced by the computational measures of grass pollen sculptural element size and density developed by Mander *et al.* (2013) do not correlate with phylogeny, and Page (1978, p. 314) has commented that the ‘overall taxonomic significance of sexine pattern in grass pollen is very limited’.

Poaceae is a large plant family and currently contains 11 157 species (GPWG 2012). A recently published molecular phylogeny of grasses (GPWG 2012) provides clarification on the evolutionary relationships of the 13 subfamilies that are contained within Poaceae and, in this paper, we use this to review current understanding of the

phylogenetic distribution and taxonomic utility of grass pollen surface ornamentation. We focus our attention on grass pollen morphotypes (e.g. Köhler & Lange 1979) and aim to investigate the following questions: (1) how much SEM data have been collected on grass pollen surface ornamentation? (2) How are the grass pollen morphotypes distributed among the subfamilies of Poaceae?

Database construction

We searched the literature and constructed a database of published SEM images of grass pollen surface ornamentation. This literature included SEM surveys of taxa from several different subfamilies within Poaceae (Watson & Bell 1975; Page 1978; Köhler & Lange 1979; Perveen & Qaiser 2012; Mander *et al.* 2013), focused studies of a single species (Datta & Chaturvedi 2004) or subfamily (Andersen & Bertelsen 1972; Grant 1972; Peltre *et al.* 1987; Liu *et al.* 2004) and isolated SEM images published as part of other morphological work (Skvarla *et al.* 2003; Sivaguru *et al.* 2012). We also include images from the PalDat pollen image database (<http://paldat.org>), with the understanding that the broader palynological community contributed these images and that their taxonomic provenance is subject to greater uncertainty. Each SEM image in these publications was visually inspected and only those images in which the surface ornamentation was clear and not masked by folding of the exine, poor illumination or deposits on the surface of the pollen grain were included in our database. Each grass taxon in the database was assigned to a Poaceae subfamily using A World-Wide Phylogenetic Classification of Poaceae (available via <http://www.tropicos.org>; published as Soreng (2000, 2001) and Soreng *et al.* (2015)).

Each SEM image in this database was grouped into one of the six morphotypes shown in Figure 1. Four of these, the *Hordeum*-type, *Triticum*-type, *Avena*-type and *Setaria*-type, are morphotypes that

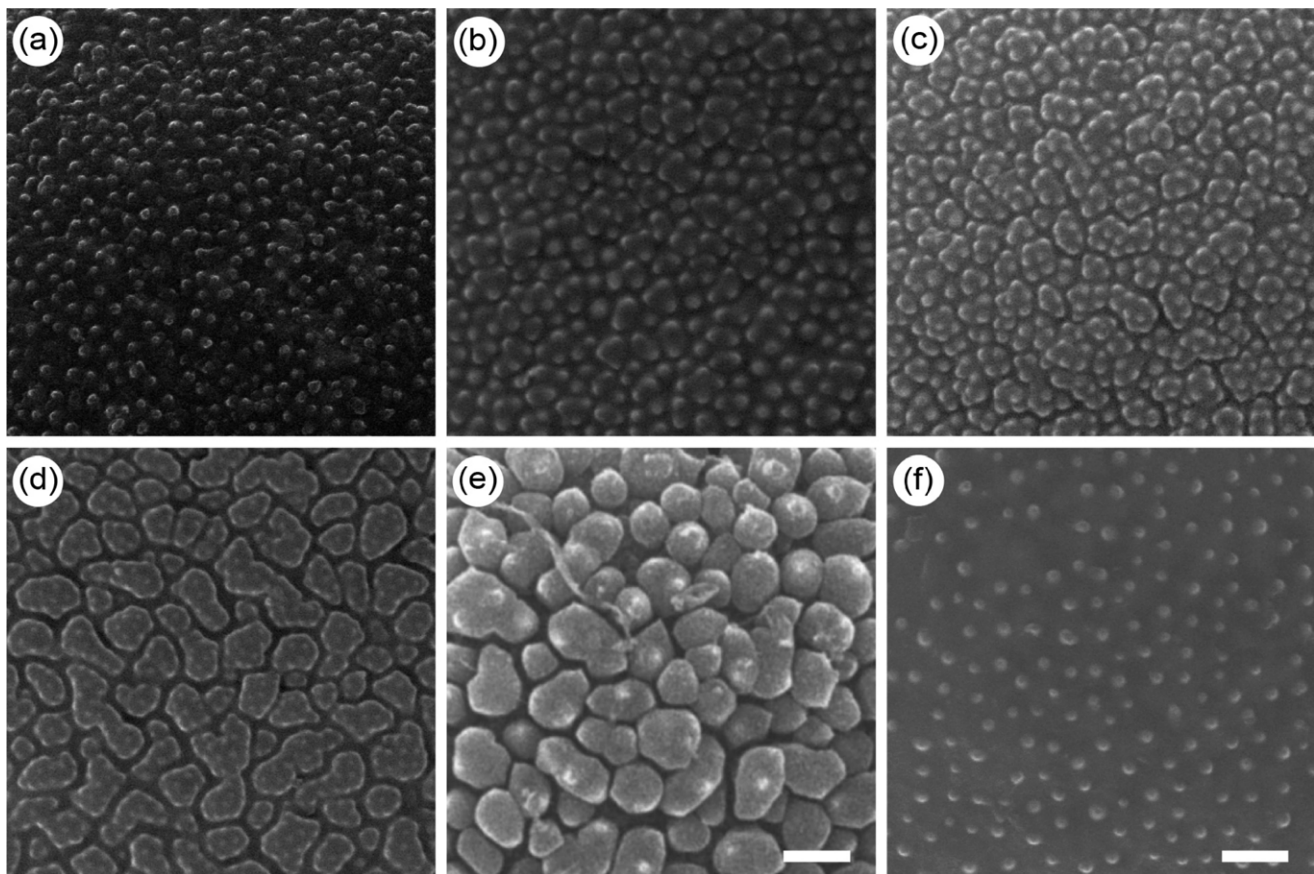


Fig. 1. Thumbnails showing examples of (a) the *Hordeum*-type, (b) *Triticum*-type, (c) *Avena*-type, (d) *Setaria*-type, (e) *Pariana*-type and (f) *Stipa*-type. Taxa shown in these thumbnails are as follows: (a) *Agropyron repens*, (b) *Triodia basedowii*, (c) *Oryza sativa*, (d) *Oplismenus hirtellus*, (e) *Pariana argentea* and (f) *Stipa tenuifolia*. (a)–(d) and (f) Same scale; scale bar 1 μ m. (e) Reproduced with permission from Skvarla *et al.* (2003, fig. 2); scale bar 1 μ m.

have been established during previous work (Köhler & Lange 1979). The *Hordeum*-type 'is characterised by single detached spinules generally not grouped in [areolae]' (Köhler & Lange 1979, p. 134) (Fig. 1a). The *Triticum*-type 'is characterised by small [areolae] bearing only one to three, rarely more, spinules' (Köhler & Lange 1979, p. 137) (Fig. 1b). The *Avena*-type 'has larger irregularly polygonal or elongate [areolae] studded with several, [1–10] spinules' (Köhler & Lange 1979, p. 137) (Fig. 1c). The *Setaria*-type 'is characterised by extensive field-like [areolae] of irregularly polygonal outlines. Their bulging surface is studded with very small pointed spinules' (Köhler & Lange 1979, p. 139) (Fig. 1d). In these descriptions, the term 'areolae' replaces 'insulae', following Punt *et al.* (2007).

To these four established morphotypes we add the *Pariana*-type and the *Stipa*-type. The *Pariana*-type is characterized by 'fused and spinulose areolate elements grouped into irregular islands interspersed with globular/granular exine particles approximately 0.25 μ m in diameter' (Skvarla *et al.* 2003, p. 925) (Fig. 1e), while the *Stipa*-type is characterized by simple scabrate ornamentation that lacks areolae and has sparsely distributed granula (Fig. 1f). Our database is summarized in Table 1.

SEM data and the taxonomic utility of grass pollen morphotypes

In our database, there are SEM data on the surface ornamentation of the pollen of 99 species of grass. This represents 0.89% of the 11 157 grass species that are currently recognized (GPWG 2012) and there are SEM data on pollen surface ornamentation for just six of the 13 subfamilies within Poaceae (Table 1). There is a positive relationship between the species richness of each subfamily and the number of

species with an SEM record of their surface ornamentation (Fig. 2). Larger subfamilies have the most species with an SEM record of their pollen surface ornamentation, whereas smaller subfamilies have received considerably less attention (Fig. 2).

The Poaceae subfamilies are traditionally grouped into three clades (Table 1). There are no SEM data on the pollen of the early diverging APP clade of grasses, which consists of three small subfamilies and contains a total of 28 species (Fig. 3). In the BEP clade, which contains 5423 species within three subfamilies that each exclusively use the C3 photosynthetic pathway (Christin *et al.* 2013), there are SEM data on the surface ornamentation of one species within the Ehrhartoideae subfamily, six species within the Bambusoideae subfamily and 53 species within the Pooideae subfamily (Fig. 3). In the PACMAD clade, which contains a total of 5706 species and includes those that have evolved C4 photosynthesis (Christin *et al.* 2013), there are SEM data on the surface ornamentation of 22 species within the Panicoideae subfamily, one species within the Arundinoideae subfamily and 16 species within the Chloridoideae subfamily (Fig. 3). However, there are no SEM data on the pollen of the six species contained in the *Centropodia* clade or the other three subfamilies within the PACMAD clade (Fig. 3).

There are not enough SEM data to reliably assess the phylogenetic distribution of grass pollen surface ornamentation patterns. However, the available information indicates that the *Avena*-, *Hordeum*-, *Setaria*- and *Triticum*-types all occur in both the BEP and PACMAD clades (Fig. 3), and this shows that at present these four morphotypes cannot be used to reliably distinguish major groups of grasses (Watson & Bell 1975; Page 1978; Kellogg 2015). It is intriguing, however, that the *Pariana*- and *Stipa*-types are restricted to the BEP clade, and that the *Pariana*-type only occurs in the Bambusoideae subfamily (Fig. 3). Although the data are

Table 1. A summary of our database showing the morphotypes contained within each subfamily, together with an example taxon and supporting reference for each case

Clade	Subfamily	Ornamentation types	Example taxon	Reference
PACMAD	Panicoideae	Avena	<i>Saccharum strictum</i>	Page (1978)
		Hordeum	<i>Zea mays</i>	Watson & Bell (1975)
		Setaria	<i>Tripsacum dactyloides</i>	Grant (1972)
		Triticum	<i>Panicum miliaceum</i>	Köhler & Lange (1979)
	Micrairoideae	No data		
	Arundinoideae	Avena	<i>Phragmites australis</i>	Watson & Bell (1975)
	Danthonioideae	No data		
	<i>Centropodia</i> clade	No data		
	Chloridoideae	Hordeum	<i>Spartina townsendii</i>	Liu <i>et al.</i> (2004)
		Setaria	<i>Spartina pectinata</i>	Page (1978)
		Triticum	<i>Triodia basedowii</i>	Mander <i>et al.</i> (2013)
BEP	Aristidoideae	No data		
	Ehrhartoideae	Avena	<i>Oryza sativa</i>	Köhler & Lange (1979)
	Bambusoideae	Hordeum	<i>Arundinaria alpina</i>	Page (1978)
		Pariana	<i>Pariana argentea</i>	Skvarla <i>et al.</i> (2003)
		Setaria	<i>Olyra micrantha</i>	Page (1978)
		Triticum	<i>Fargesia nitida</i>	http://www.paldat.org
		Stipa	<i>Diandrolyra bicolor</i>	Page (1978)
	Pooideae	Avena	<i>Avena sativa</i>	Köhler & Lange (1979)
		Hordeum	<i>Bromus inermis</i>	Köhler & Lange (1979)
		Setaria	<i>Poa compressa</i>	Skvarla <i>et al.</i> (2003)
		Stipa	<i>Stipa tenuifolia</i>	Mander <i>et al.</i> (2013)
Triticum		<i>Triticum aestivum</i>	Köhler & Lange (1979)	
APP	Puelioideae	No data		
	Pharoideae	No data		
	Anomochlooideae	No data		

extremely limited (only 6 of the 1462 species contained within the Bambusoideae subfamily are present in our database, for example (Fig. 3)), this raises the possibility that a simple morphotyping approach based on SEM images of surface ornamentation might be used to distinguish the BEP clade and the Bambusoideae subfamily in the fossil record and in samples of modern material.

Future work

In general, very little SEM data on grass pollen surface ornamentation have been acquired (e.g. Fig. 2). The pollen of cereal grasses such as

rye (*Secale cereale*), wheat (*Triticum* spp.) and barley (*Hordeum vulgare*) have received considerable morphological attention in the context of archaeobotanical studies of crop domestication (e.g. Köhler & Lange 1979), but the pollen morphology of smaller subfamilies, such as the Micrairoideae and the Danthonioideae, is unknown (see Fig. 3). This lack of data is problematic for several reasons. From an evolutionary perspective, it prevents phylogenetic analyses of the mechanisms that have generated the extant diversity of grass pollen form. For example, it is unclear whether the occurrence of the *Triticum*-type in the Panicoideae, Chloridoideae and Pooideae (Fig. 3) reflects multiple character state gains or the loss, reversal or regaining of characters (e.g. Cronk 2009). In this context, data on the pollen morphology of the early diverging APP clade of grasses would be useful in order to generate comparisons between more primitive and more derived subfamilies. As noted by Kellogg (2015), the lack of data also hinders efforts to examine whether aspects of grass pollen morphology, such as surface ornamentation, have any clear functional significance or whether they correlate with other aspects of biology, such as environment or mode of pollination (e.g. Soderstrom & Calderón 1971; Skvarla *et al.* 2003).

The lack of SEM data on grass pollen surface ornamentation also represents an obstacle to the classification of fossil grass pollen in terms of extant taxa (e.g. Lee *et al.* 2004). This is because workers are currently unable to compare fossil specimens to a database or monograph that displays the full range of natural morphological variation of extant taxa (e.g. Birks & Birks 1980; Mander & Punyasena 2014), and the morphological boundaries of grass pollen morphotypes are somewhat unclear as a result (Mander *et al.* 2013). Additionally, the commonness and rarity of each morphotype is unclear. For example, in the context of the entire Poaceae phylogeny, the *Setaria*-type occurs 38 times while the *Pariana*-type occurs only twice (Fig. 3). However, there are not enough data at present to reliably indicate whether certain morphotypes are common or rare in each subfamily and this information might help to constrain the affinities of fossil specimens.

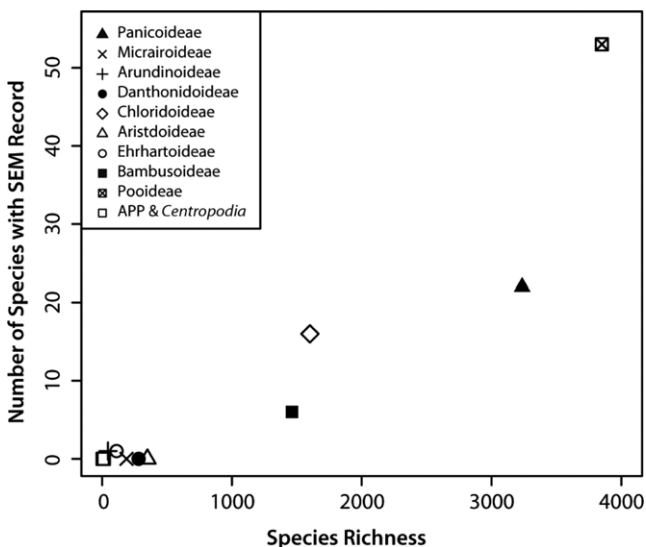


Fig. 2. Scatterplot showing the relationship between the species richness of each subfamily within Poaceae and the number of species with an SEM record of their surface ornamentation. The APP clade of grasses and the *Centropodia* clade are all represented by a single open square.

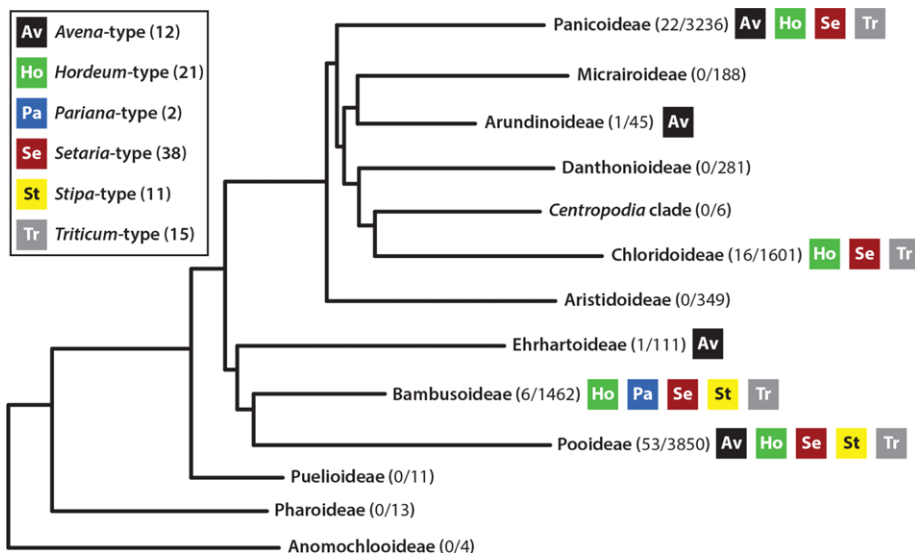


Fig. 3. Phylogenetic tree showing the distribution of grass pollen morphotypes among the subfamilies of Poaceae. Numbers in brackets next to each subfamily show the number of SEM images in our database followed by the number of species within the subfamily. For example, the Panicoideae contains a total of 3236 species and in our database there are published SEM images of the pollen surface ornamentation of 22 species from this subfamily. Numbers in brackets next to each morphotype in the key show the abundance of each morphotype in our database. For example, there are 12 species that are characterized by the *Avena*-type and 21 species that are characterized by the *Hordeum*-type. Poaceae phylogeny and subfamily species richness from GPWG (2012).

We remain encouraged by the possibility that grass pollen can play a role in future palaeoecological studies of grasslands. We do note, though, that the spread of morphotypes across the Poaceae phylogeny suggests that this role is likely to be restricted to the generation of raw diversity counts, which could be used to track diversification history and compositional change through time. Such diversity counts could be calibrated by data from fossil phytoliths, which have high taxonomic resolution and allow workers to assess the presence or absence of important subfamilies (Piperno 2006; Strömberg 2011). Owing to the subtlety of grass pollen morphology and, in particular, the surface ornamentation patterns that we have focused on here, we feel that computational approaches (Mander *et al.* 2013) offer the best chance of generating consistently reproducible classifications (e.g. Mander & Punyasena 2014). However, the morphotyping approach to grass pollen classification that we have reviewed in this paper could be used to generate broad groupings of morphologically similar grass pollen grains in fossil samples, which could then be classified to finer taxonomic levels using computational methods. Morphotypes generated by human analysts could also be used as a yardstick against which to compare computational classifications (e.g. Mander *et al.* 2014) and, of course, also represent an archive of ancient biodiversity in their own right.

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