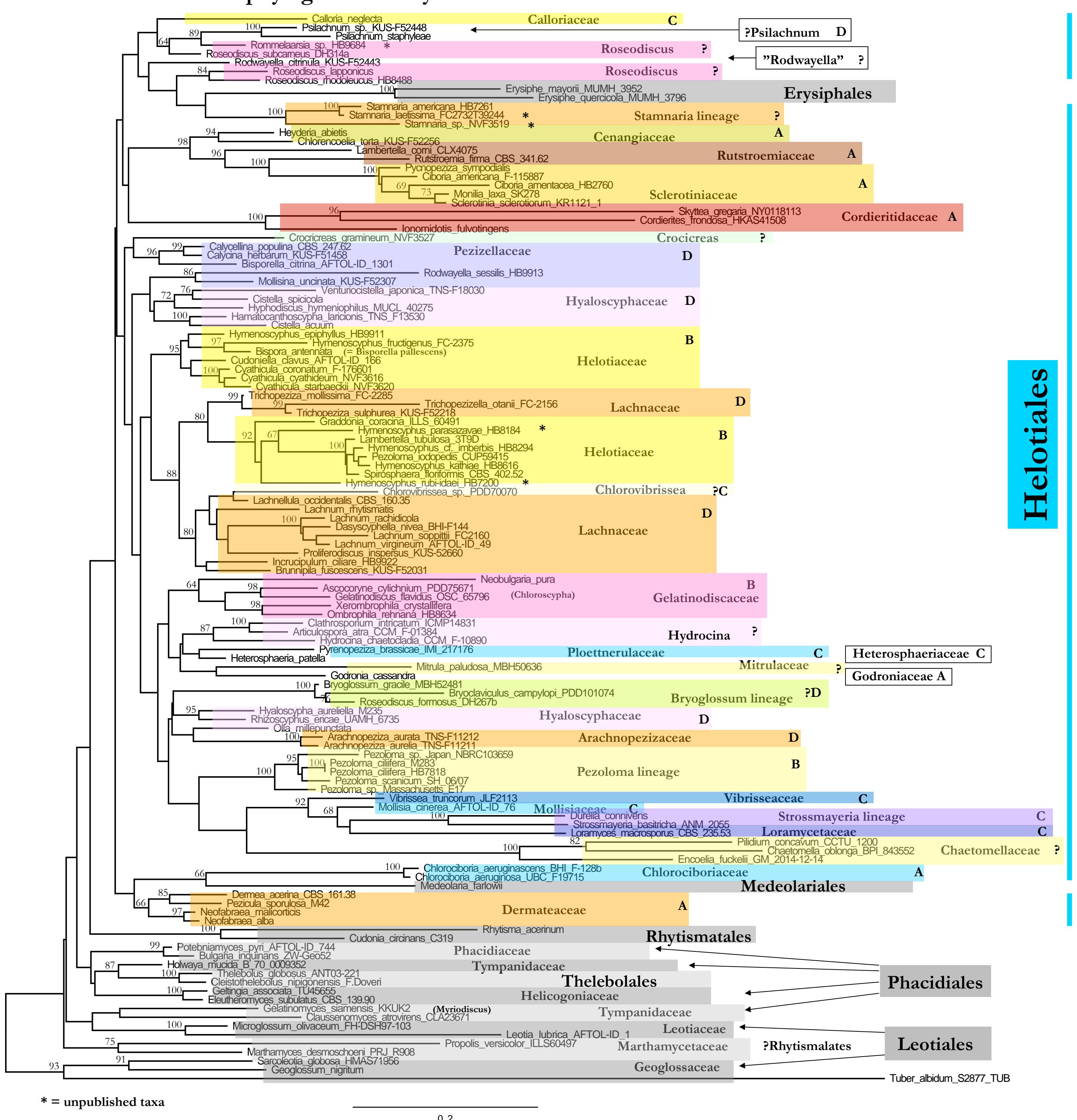


INTRODUCTION – This study arose from a compilation of the non-lichenized discomycetes for the 13th edition of 'A. Engler's Syllabus of Plant Families', in which some new families were established and forgotten ones resurrected. It includes recent, partly unpublished results and tries to combine morphological and molecular data. 24 families and some undescribed lineages are here proposed in the Helotiales *sensu stricto* (excluding Leotiaceae, Phacidiales, and some other orders). Paraphyletic groups are accepted whenever obvious morphological support is noted. A pragmatic approach based mainly on morphology tries to arrange these 24 families in 4 main groups (A-D). Our phylogenetic analysis hardly represents these groups, although it shows high bootstrap support for most of the proposed families. The lack of support at the basal nodes of the tree indicates limitations of our data, which might be

due to taxon sampling errors and/or lack of more gene regions. We consider this presentation as a stimulus for further work, which should include more, also protein-coding, gene regions. The families are mostly well represented with high bootstrap support, and also some family associations show high support. Some unexpected results are worth of mention: (1) *Roseodiscus formosus* does not cluster with the type *R. rhodoleucus*; (2) *Crocireas* requires restriction to the type *C. gramineum*, while the large genus *Cyathula* is unrelated to them; (3) enceloioid turned out to belong in various families, *E. fuscicula* in Chaetomellaceae, *E. fascicularis* in Sclerotiniaceae, *E. tihacea* in Rutstroemiaceae, *E. heteromera* and *E. fimbriata* in Cordieritidaceae, and the type *E. furfuracea* in Cenangiaceae. A large number of helotialean genera are still unsequenced, and even some sequenced ones cannot presently be placed in any of the presently recognized families.

ML phylogenetic analysis based on ITS+LSU rDNA



* = unpublished taxa

0.2

Table of selected characters of families and lineages here recognized within the Helotiales.

Lineage	inc.s.	devel.	stipe	erump.	stroma	exc.	hairs	cryst.	ionom.	par.	VBs	amyl.	ring	croz.	sheath	lipid	asex.	cdg.	lifest.	desicc.
A	Sclerotiniaceous & enceloioid taxa																			
	Sclerotiniaceae	Me(C)	st	er/su	+/-	p/(g)	-/(h)	-/(+)	-	c	+/-	B/r	S	+/-	(1')	h	s/(p)	hyg	xer	
	Sclerotiniaceae	Pr(Me)	st	er/su	+/-	p/(g)	-/(h)	+/-	-	c	+/-	B/r	S	+/-	(1')	h	s/(p)	hyg		
	Cenangiaceae	Me(C)s	st	er	-	g	-	-	c	+/-	B/r	S	+/-	(1')	h	s/(p)	hyg			
	Cordieritidaceae	Me	st/s	er/su	-/+	g/(i)	-h	-	c	-	-	-	-	-	-	-	p	s/p	xer/hyg	
	Chlorociboriaceae	Pr	st	er/su	-	g/i	-h	-	c	-	B/r	C	+/-	-	-	-	s	hyg		
	Dermateaceae	Pr/(Me)	st	er	-	g	-	-	c	-	B/r	C	+/-	-	-	-	p	s/p	xer	
	Godroniaceae	Me	s/(st)	er	-	g/p	-	+/-	c	-	B/r	C	+/-	-	-	-	p	p	xer	
	Gelatinodiscaceae	Pr	s/st	su	-	p/g	-	+/-	-	c	+/-	B/r	C	+/-	-	-	s/(p)	hyg/xer		
	Hyaloscyphaceae	Pr	s/st	su	-	p/g	-	+/-	-	c	+/-	B/r	C	+/-	-	-	s/(p)	hyg		
B	Helotaceous taxa																			
	Discinella-Pezoloma	Pr	s	su	-	p	-t	-	c	-	B/r	C	+/-	-	-	-	h	s	hyg	
	Helotiaceae	Pr	s/t/s	er/su	-/+	p/g	-/(h)	-/(+)	-	c	+/-	B(r)	H/C	+/-	+/-	-	h/s	h/p	hyg(xer)	
	Roesleriaceae	Maz	st	su?	-	p/-	-	-	c	? -	-	-	-	-	-	1	-	s/p	hyg	
	Callosporaceae	Me?	s	er	-	g	-	-	c	-	B(r)	C	+/-	-	-	1-3	h/s	p	xer/hyg	
	Ploettnerulaceae	Me	s	er	-	g	-	-	c	-	B(r)	C	+/-	-	-	1-2	a/p	p	xer	
	Mollisiaceae s.i.	Pr	s	su(er)	-	g	-h	-	c	+/-	B(r)	C	+/-	-	-	1-3	h/s	p	hyg	
	Hylomycetaceae	Me/Ci	s?	-	g	-	-	c	? -	-	-	-	-	-	?	h	h/p	s	hyg	
	Vibrissaceae	Pr	s/st	su	-	g	-	-	c	+/-	B/r	V	+/-	-	-	1-3	h	h/p	s	hyg
	Strossmayeria	Pr	s	su	-	p	-	-	c	+/-	b/-	C	+/-	+/-	(amyl)	1-3	h	h/p	s	Xer
C	Mollisiaceous taxa																			
	Bryoglossum	Cs	st	?	-	p	-	-	c	-	B/-	T	+/-	-	?	1	-	p	Hyg	
	Lachnaceae	Pr	s/st	su	-	p/g	h	+/-	-	I(c)	+/-	B/r/-	C	+/-	-	-	1-3	p	s/(p)	hyg(xer)
	Arachnopezizaceae	Pr	s	sub	-	g	h	-	c	-	B	C	+/-	-	(app)	1-3	-	h	Hyg	
	Hyaloscyphaceae	Pr/(C)	s/st	su	-	p/g	h/-	-	c	-	B/r/-	C	+/-	-	(*)	1-3	h/p/e	p	hyg(xer)	
	Pezizellaceae	Pr	s/st	su	-	p/g	h/-	-	c	-	B/r/-	C	+/-	-	(*)	1-3	h/p/s	p	hyg(xer)	
D	Hyaloscyphaceous taxa																			
	Development: Cor = corticioid; Cs = davate-stipitate; Pr/Me = apothecialoid; opening in the pro- (Pr) or mesohymenial (Me) phase; Cl = cleistothelial; Maz = mazacodium. Stipe = s: sessile; st: stipitate. Erumpent: er = erumpent; su = superficial. Stroma: dark stromatous host tissue present (+) or absent (-). Excipulum: g = globose; p = prismatic; i = intricate. Hairs: h = hairs; t = teeth. Crystals: octahedral crystals present (+) or absent (-). Ionomidotic: KOH releasing pigment into medium (ionomidotic reaction), present (+) or absent (-). Paraphysis shape: c = cylindrical to clavate; l = lanceolate. VBs: refractive vacuoles in paraphyses/excipular cells present (+) or absent (-). Amyloid: iodine reaction of apical ring (Lugol). B = cyanobiotic (blue); R = hemiamyloid (red, types RB/RR, - = inamyloid). Ring: apical ring type. S = Sclerotinia-type; C = Calycina-type; Pe = Pezizella-type; Pl = Hymenoscyphus-type; V = Vibrissa-type. Croziers: = arcuate at the ascus base present (+) or absent (-). Sheath around ascospores, either gelatinous or separating after discharge (ap = terminal gelatinous appendages, amyl = amyloid sheath). Lipid: relative lipid content of ascospores, 1 = low, 2 = medium, 3 = high. Asexual state: h = hymenocarpous; pe = penicillate; s = symmetral; sporochidial; a = acervular; p = pycnidial. Conidiogenesis: h = holoblastic; p = phialic; a = arachnid. Lifesyle: s ± saprobic; p = ± parasitic (or mycoherbivore). Desiccation tolerance: xer = desiccation-tolerant (xeric substrate); hyg = desiccation-sensitive (hygric substrate).																			

COMMENTS ON SOME CHARACTERS

Asci structure: Striking correlations between apical ring types and molecular data disclosed various misplaced taxa. E.g., species previously placed in *Torrentiella* (Rutstroemiaceae, *Sclerotinia*-type) had to be transferred to the unrelated new genus *Hymenorrendiella* (Helotiaceae, *Hymenoscyphus*-type). Or *Bisporella citrina* and allies (*Calycina*-type) were transferred to *Calycina* (Pezizellaceae), whilst the type *B. pallens* (teleomorph of *Bisporella antennata*, *Hymenoscyphus*-type) clustered in Helotiaceae.

Vacuoles: Presence vs. absence of refractive vacuoles (= VBs, living state!) correlates with phylogenetic lineages, permitting separation at the family level:

VBs present

Mollisiaceae
Pezizellaceae
Cytidiaceae (Helotiaceae)
Crocires (incertae sedis),
Cenangiaceae (major part)

VBs absent

Ploettnerulaceae
Hyaloscyphaceae
Cyathula (Helotiaceae)
Cordieritidaceae

GROUP C - Mollisiaceous taxa

subgroup I - VBs usually absent

Fam. Heterosphaeriaceae Rehm - 1 genus (7): *Heterosphaeria* (= *Heteropelta*)
Hysteropezizella lineage - 5-7 genera (42): e.g. *Cephaea*, *Coronularia*, *Hysteropeziza*, *Hysteropezella*

Fam. Callariaceae Marchand - c. 15 genera (c. 110): e.g. *Callaria* (= *Cylindrocallia*), *Diploncia*, *Duberia*, *Eupropoda*, *Laetaria*, *Naufragia*, *Naevia*, *Naeviopsis*, *Ploettnera*

Fam. Drepanopezizaceae Bat. & H. Maia - c. 10 genera (c. 52): e.g. *Blumeriella* (= *Phloeospora*), *Diplocarpon* (= *Marssonina*), *Drepanopeziza* (= *Gloeosporidella*), *Leptotrichila* (= *Sporomyces*), *Pseudopeziza*, *Sporopila* (= *Melanodiscus*)

Fam. Ploettnerulaceae Kirschst. (= Pyrenopezizaceae Ruhland ex Welen.) - c. 7 genera (725): e.g. *Dennisiodes*, *Lasiomollisia*, *Oculumacula* (= *Hegdardia*), *Pirottaea* (= *Ploettnerula*), *Pyrenopeziza* (= *Cadophora*)

subgroup II - VBs present

Fam. Mollisiaceae Rehm - c. 11 genera (210): e.g. *Mollisia* (= *Belonopsis*, *Cystodendron*, *Hagnulia*, *Phaeomollisia*, *Phiaedrophala*, *Tapesia*), *Nimbomollisia*

Fam. Loramycetaceae Dennis ex Digby & Goos - 2 genera (4): *Loramycetes*, *Oblectodiscus*

Fam. Vibrissaceae Locq. ex Korf - 3-4 genera (c. 35): *Leucovibrissa*, *Pocillum*, *Vibrissa* (= *Anuringa*)

Fam. Strossmayeria lineage - 3 genera (235): *Durella*, *Gorgoniceps*, *Strossmayeria* (= *Pseudospirodes*)

GROUP D - Hyaloscyphaceous taxa

Fam. Arachnopezizaceae Hosoya, J.G. Han & Baral - 5 genera (c. 30): *Arachnopeziza*, *Arachnoscypha*, *Austropeziza*, *Eriopeziza*, *Paranopeziza*