

## Mushrooms nomenclatural novelties no. 14

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**Fibroporiaceae** Audet, fam. nov.

**IF554248**

Basidiomata annual, resupinate or pileate (flabelliform to dimidiate), often widely effused, margin often fimbriate with rhizomorphs. Pore surface white to apricot-orange or lemon yellow to curry buff when fresh, pores circular to angular, dissepiments thin, entire to slightly lacerate. Subiculum soft and cottony, white to apricot-orange. Hyphal system monomitic to dimitic, generative hyphae mostly with clamps and thin-walled, occasionally thick-walled, rarely thick-walled with simple septa, skeletal hyphae dominant in subiculum and rhizomorphs, solid to thick-walled, rarely dichotomously branched, IKI–, CB–.

Cystidia absent, cystidioles present or absent. Basidiospores oblong to broadly ellipsoid, slightly thick-walled, hyaline, smooth, uninucleated, IKI–, CB–. Grows on angiosperm or gymnosperm wood. Causing brown decay. Nuclear behavior normal and mating system tetrapolar heterothallic (Fibroporia).

Holotype : *Fibroporia* Parmasto, Consp. System. Corticiac. (Tartu): 176, 1968.

Other genus accepted in this family: *Pseudofibroporia* Yuan Y. Chen, B.K. Cui & Y.C. Dai, in Chen, Wu, Wang & Cui, Mycol. Progr. 16(5): 528 (2017).

Comment : Only by RPB1, the family Fibroporiaceae based on some species of *Fibroporia* (4) is on another clade than *Adustoporiaceae*, *Amyloporiaceae*, *Lentoporiaceae* and *Rhodoniaceae* in bayesian analyses and also by alignment with unpublished phylogeny. With multi loci *Pseudofibroporia* was suggested as the closest relative of *Fibroporia* from Chen et al. paper 2017. See personal phylogeny below about the support of this new family.

Table 2 from figure 2

Taxon	Phylotree	SSU (18S) Genbank
<i>Adustoporia sinuosa</i>	Asinu6786	AY336786
<i>Amyloporia xantha</i>	Axantha5919	KR605919
<i>Fibroporia albicans</i>	Falb0524	KU550524
<i>Fibroporia bambusae</i>	Fbamb0527	KU550527
<i>Fibroporia bohemica</i>	Fbohe0529	KU550529
<i>Fibroporia ceracea</i>	Fcer0530	KU550530
<i>Fibroporia citrina</i>	Fcit0533	KU550533
<i>Fibroporia radiculosa</i>	Fradi0538	KU550538
<i>Fibroporia vaillantii</i>	Avail8583	AJ488583
<i>Fomitopsis pinicola</i>	Fpinic5858	KR605858
<i>Lentoporia carbonica</i>	Acarb9059	U59059
<i>Pseudofibroporia citrinella</i>	Pcitr0540	KU550540

Table 6 from figure 6

Taxon	Phylotree	Genbank	
		LSU	RPB2
<i>Adustoporia sinuosa</i>	Asinu5070	KC585070	KT895894
<i>Amyloporia carbonica</i>	Acarb7844	AF287844	AY218470
<i>Amyloporia subxantha</i>	Asubx8826	KT968826	KT895897
<i>Amyloporia xantha</i>	Axant2284	EU232284	KP134912
<i>Cerrena unicolor</i>	Cunic5209	KP135209	KP134968
<i>Fibroporia albicans</i>	Falb0485	KU550485	KR610838
<i>Fibroporia bambusae</i>	Fbamb0488	KU550488	KU55054
<i>Fibroporia ceracea</i>	Fcera0490	KU550490	KU550547
<i>Fibroporia citrina</i>	Fcit0492	KU550492	KU550551
<i>Fibroporia gossypium</i>	Agosy2298	EU232298	KU550550
<i>Fibroporia radiculosa</i>	Fradic5167	KC585167	KT895899
<i>Postia rancida</i>	Pranc0999	KX900999	KX901242
<i>Pseudofibroporia citrinella</i>	Pscit0500	KU550500	KU550556
<i>Rhodonía placenta</i>	Pplac2285	KC585223	KT893746

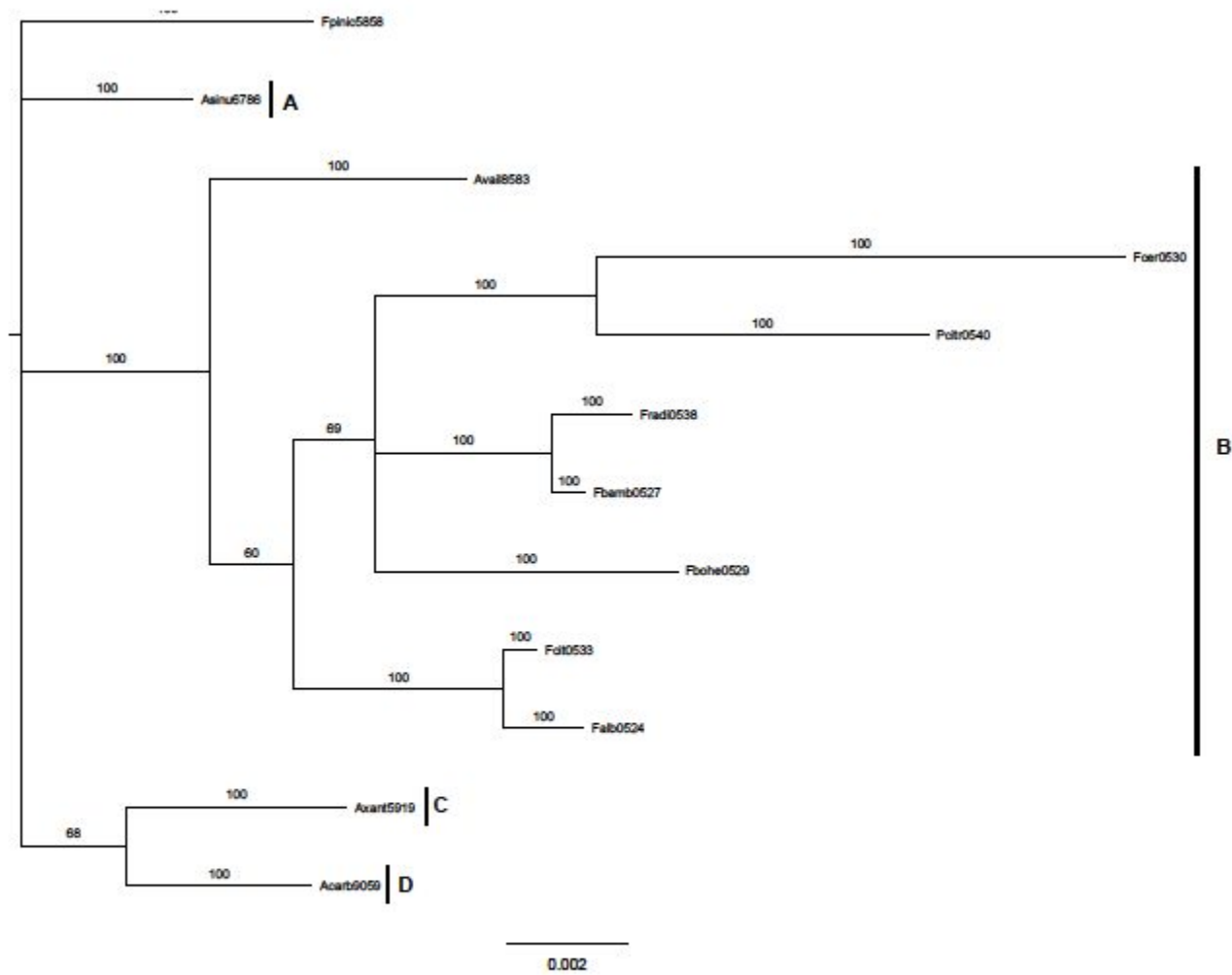


Figure 2: Strict consensus phylogree of small subunit ribosomal RNA gene (18S) calculated by the MrBayes software (Ronquist et al. 2015; 1 000 000 generations). A: *Adustoporiaceae*; B: *Fibroporiaceae*; C: *Amyloporiaceae*; D: *Lentoporiaceae*; outgroup: *Fpinic5858*

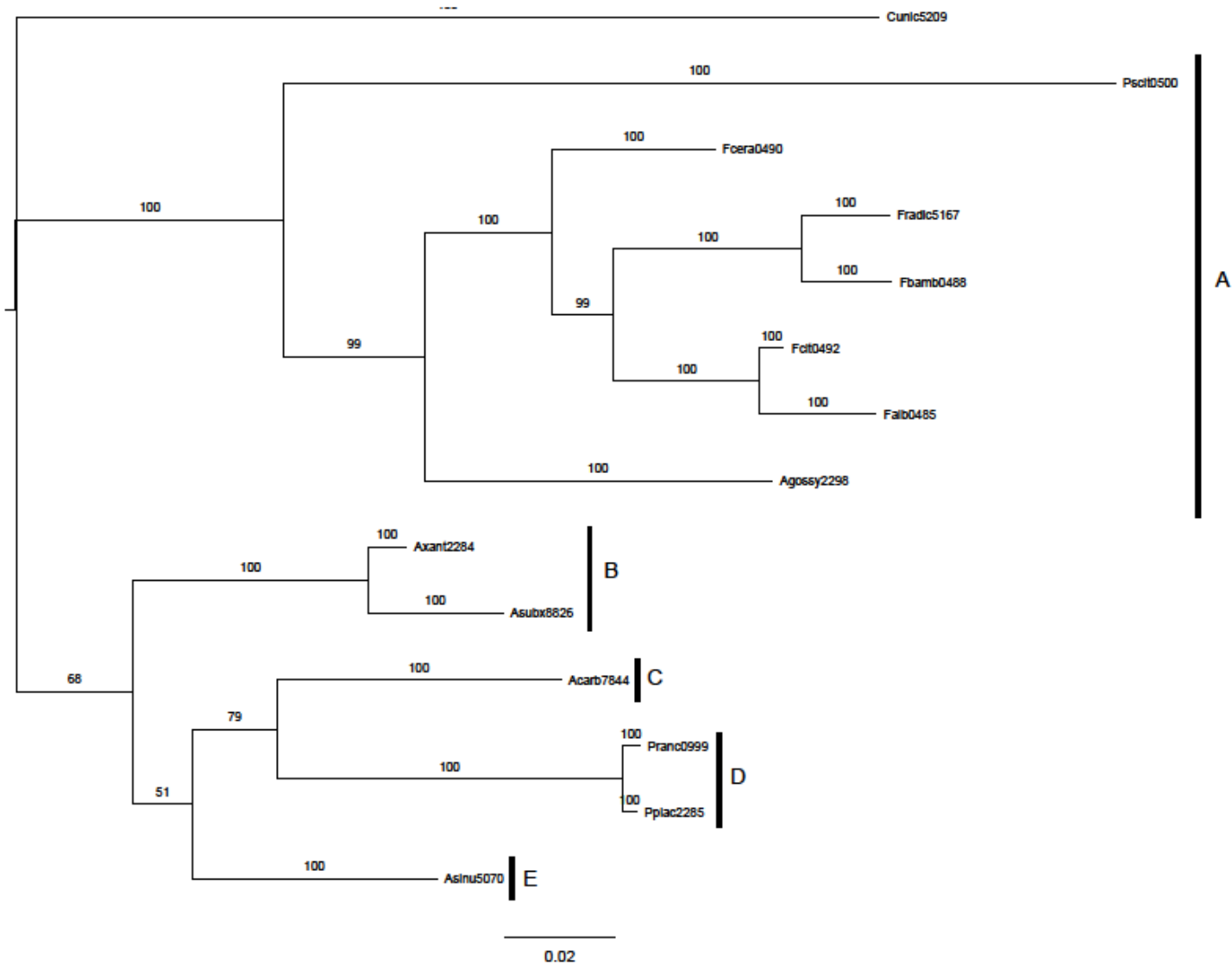


Figure 6: Strict consensus phylogenetic tree of LSU +RPB2 calculated by the MrBayes software (Ronquist et al. 2015; 1 000 000 generations). A: *Fibroporiaceae*; B: *Amyloporiaceae*; C: *Lentoporiaceae*; D: *Rhodoniaceae*; E: *Adustoporiaceae*; outgroup: *Cunic5209*