

*Pseudoacremonium sacchari*





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## *Pseudoacremonium* Crous, *gen. nov.*

*Etymology.* Name derived from pseudo- (false-, in Greek) and the genus name *Acremonium*.

*Mycelium* consisting of hyaline, septate, branched hyphae, developing brown zones with solitary brown, globose chlamydo-spores. *Conidiophores* erect, subcylindrical, rarely branched, 1–3-septate, smooth, but with age brown, verruculose and warty. *Conidiogenous cells* terminal, integrated (Type III phialides sensu Mostert et al. 2006), either elongate-ampulliform or navicular, tapering towards apex, with prominent periclinal thick-

ening and inconspicuous collarette, hyaline, smooth, becoming brown and verruculose to warty. *Conidia* in unbranched chains, ellipsoid, aseptate, smooth, hyaline, thin-walled, becoming brown, verruculose, guttulate, with scar at each end, somewhat darkened and refractive; older conidia brown and verruculose, appearing flaky when mounted in lactic acid or water.

*Type species.* *Pseudoacremonium sacchari*.  
MycoBank MB808925.

## *Pseudoacremonium sacchari* Crous, *sp. nov.*

*Etymology.* Named after the host genus from which it was isolated, *Saccharum*.

*Mycelium* consisting of hyaline, septate, branched, 2–3 µm diam hyphae, developing brown zones with solitary brown, globose, 5–7 µm diam chlamydo-spores. *Conidiophores* erect, subcylindrical, rarely branched, 1–3-septate, up to 80 µm tall, 3–3.5 µm diam, hyaline, smooth, but with age brown, verruculose and warty. *Conidiogenous cells* terminal, integrated (Type III phialides sensu Mostert et al. 2006), either elongate-ampulliform or navicular, tapering towards apex, 25–30 × 2.5–4 µm, apex 1–1.5 µm diam, with prominent periclinal thickening and inconspicuous collarette, hyaline, smooth, becoming brown and verruculose to warty. *Conidia* in unbranched chains, ellipsoid, aseptate, smooth, hyaline, thin-walled, becoming brown, verruculose, guttulate, with scar at each end, 0.5–1 µm diam, somewhat darkened, and refractive, (6–)7–8(–9) × 2.5(–3) µm; older conidia brown and verruculose, appearing flaky when mounted in lactic acid or water.

*Culture characteristics* — Colonies reaching 7 mm diam after 2 wk at 22 °C, with sparse aerial mycelium and even, smooth margins. On PDA surface and reverse dirty white. On OA surface pale olivaceous-grey. On MEA surface cinnamon, reverse cinnamon with patches of vinaceous, with a diffuse brick pigment in agar.

*Typus.* VIETNAM, Dong Nai Province, Cat Tien National Park, Nam Cat Tien Sector, on leaves of *Saccharum spontaneum* (*Poaceae*), in association with *Pyricularia contorta*, 5 Mar. 2013, A. Alexandrova & O. Deshcherevskaya (holotype CBS H-21704, culture ex-type CPC 23155 = CBS 137990; ITS sequence GenBank KJ869144, LSU sequence GenBank KJ869201, MycoBank MB808926).

*Notes* — The acremonium-like morphology and pigmented conidia are reminiscent of the genus *Pseudogliomastix*, which clusters outside the *Hypocreales* (Rehner & Samuels 1995), and is thus unavailable to accommodate this taxon, for which the new genus *Pseudoacremonium* is established. *Pseudoacremonium* is characterised by forming chlamydo-spore, and having mycelium that turns brown with age, and unbranched conidial chains that become brown (encrusted, flaky) with slightly darkened scars at each end of the conidia.

*ITS.* Based on a megablast search of NCBI's GenBank nucleotide database, the closest hits using the ITS sequence are *Acremonium cerealis* (GenBank AB540571; Identities = 560/633 (88 %), Gaps = 31/633 (4 %)), *Acremonium persicinum* (GenBank AB540575; Identities = 555/634 (88 %), Gaps = 30/634 (4 %)) and *Stromatonectria caraganae* (GenBank HQ112288; Identities = 694/799 (87 %), Gaps = 41/799 (5 %)).

*LSU.* Based on a megablast search of NCBI's GenBank nucleotide database, the closest hits using the LSU sequence are *Lasionectria mantuana* (GenBank GQ505994; Identities = 826/843 (98 %), no gaps), *Acremonium cerealis* (GenBank HQ232014; Identities = 815/832 (98 %), Gaps = 1/832 (0 %)) and *Hydropisphaera erubescens* (GenBank AY545726; Identities = 841/859 (98 %), Gaps = 2/859 (0 %)).

*Colour illustrations.* Cat Tien National Park, Vietnam; conidiophores and conidia in culture. Scale bars = 10 µm.