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Aculeastrum M. Scholler, U. Braun & Bubner

Based on analysis of ITS sequence data, Scholler et al. (2022), introduced Aculeastrum to accommodate A. americanum (Farl.) M. Scholler & U. Braun (= Pucciniastrum arcticum var. americanum Farl.) as a type species and A. arcticum (Tranzschel) M. Scholler & U. Braun (≡ *P. arcticum* Tranzschel). *Aculeastrum americanum* was originally described as a rust pathogen on leaves of *Rubus occidentalis* from the USA (Farlow 1908), while A. articum was described on leaves of Rubus spp. from North America, Lappland, and Russia (Tranzschel 1895). Aculeastrum with A. americanum as the type was re-collected from leaves of Rubus occidentalis in the USA (Scholler et al.2022). Aculeastrum produces uredinia and telia on Rubus spp. and its aecia on needles of the alternate host, Picea. The genus is characterized by uredinia with four to six conspicuously spiny ostiolar cells, urediniospores with 4–6(–7) strictly bipolar germ pores, and mostly 2–4-celled teliospores without any visible germ pores. Aculeastrum resembles Quasipucciniastrum and Thekopsora and members of all three genera have telial hosts within the *Rosaceae* and have previously been included in the genus *Pucciniastrum* (family *Pucciniaceae*). The three genera differ in morphology, life cycle, symptomology, and host range. Phylogenetic evidence obtained from the analysis of ITS sequence data shows that all three are in a distinct lineage within the family Coleosporiaceae (Scholler et al.2022). Based on phylogenetic inferences from analysis of 28S, 18S and CO3 datasets, Pucciniastrum arcticum var. americanum had been transferred to Thekopsora americana (Farl.) Aime & McTaggart, in Pucciniastraceae by Aime & McTaggart (2020). The evidence of Scholler et al. (2022), clearly places Aculeastrum in Coleosporiaceae (Melampsorineae, Pucciniales, Pucciniomycetes).

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