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## A new combination in *Mononeuria* (Caryophyllaceae)

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The nomenclatural change *Mononeuria caroliniana* comb. nov. is proposed on the basis of molecular phylogenetic results analyzing the nuclear internal transcribed spacer.

Keywords: *Minuartia*, North America, taxonomy

### Introduction

A recent molecular phylogenetic study of *Minuartia* Loefling in Linnaeus (1753: 89) *sensu* McNeill (1962) by Dillenberger & Kadereit (2014a) demonstrated that species formerly placed in *Minuartia* were members of 11 different lineages. The 33 North American species (north of Mexico; see Rabeler *et al.* 2005) are now part of *Cherleria* Linnaeus (1753: 425), *Mononeuria* Reichenbach (1841: 205), *Pseudocherleria* Dillenberger & Kadereit (2014b: 451), and *Sabulina* Reichenbach (1832: XXIV). The results by Dillenberger & Kadereit (2014a) was later accepted by several authors (e.g., Iamonico 2014, Hernández-Ledesma *et al.* 2015, Fedorowchuk & Mosyakin 2016, Iamonico 2016, Moore & Dillenberger 2017).

*Mononeuria* was first published by Reichenbach (1841: 118) as a generic synonym to *Alsine* unranked *Uninerviae* Fenzl in Endlicher (1840: 965). Dillenberger & Kadereit (2014a) included all of the species of *Minuartia* sect. *Uninerviae* (Fenzl in Endlicher 1840: 965) Mattfeld (1921: 28) in *Mononeuria*, based on McNeill (1962). To confirm the placement and position of all North American species thought to belong to the genus *Sabulina*, we sequenced several taxa not included in Dillenberger & Kadereit (2014a) and found an unexpected placement of *Sabulina caroliniana* (Walter 1788: 141) Small (1933: 1504) in *Mononeuria*.

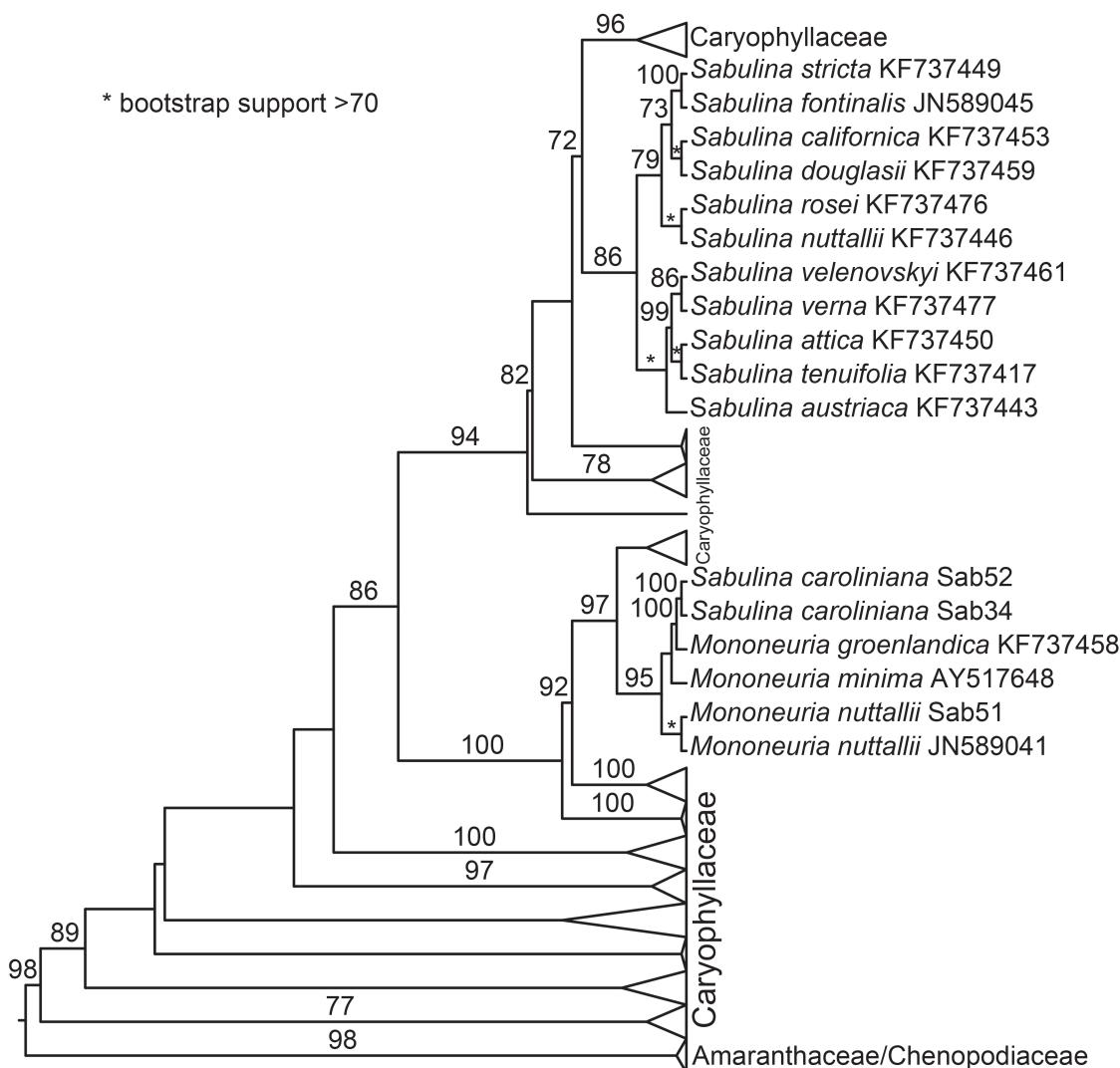
### Material & Methods

Nuclear internal transcribed spacer of two individuals of *Sabulina caroliniana* and one individual of *Mononeuria nuttallii* (Torrey & A. Gray 1838: 183) Dillenberger & Kadereit (2014a: 84) were sequenced according to the methods described by Legler & Dillenberger (2017). Material was obtained from three specimens at OSC, i.e. *S. caroliniana* Sab34: USA. Georgia: Richmond County, 08 May 1992, *Moore* 1464 (OSC 181464, isovouchers at CM, GA), *M. nuttallii* Sab51: USA. Texas: Brazos County, 08 April 1992, *Wipff & Jones* 2244 (OSC 203770, isovoucher at UTEP), and *S. caroliniana* Sab52: USA. North Carolina: Richmond County, 05 May 1967, *Bozeman & Logue* 9124 (OSC 127623, isovouchers at CONV, CUH, DSC, FUGR, HTTU, KE, MISS, MUR, NBYC, NCU, PEMB, UBC, UNCC, USCH, USMS, VSC). The new sequences were included in a larger dataset of previously published sequences representing the entire Caryophyllaceae (128 samples and three outgroup samples; modified data set of Dillenberger & Kadereit 2014a, see TreeBASE to download the alignment). Sequences were aligned with MUSCLE v.3.8.31 (Edgar 2004) implemented in seaview v.4.3.0 (Gouy *et al.* 2010). Phylogenetic analysis was carried out using RAxML v.8.0.26 (Stamatakis *et al.* 2008) with the GTR+Γ substitution model and the fast bootstrap algorithm with automatic halt based on the autoMRE criterion. Sequences were submitted to GenBank (MG839540–MG839542); alignment and phylogenetic tree can be downloaded from TreeBASE (<http://purl.org/phylo/treebase/phylows/study/TB2:S22236>).

Isovouchers were located via consulting SEINet (<http://swbiodiversity.org/seinet/>) and type images were examined via JSTOR Plants ([plants.jstor.org](http://plants.jstor.org)). Herbarium codes follow Thiers (2018+).

## Results

The samples of *Sabulina caroliniana* are nested with high bootstrap support within *Mononeuria* (Fig. 1) being well-supported as sister to *M. groenlandica* (Retzius 1795: 107) Dillenberger & Kadereit (2014a: 84). The phylogeny unambiguously shows that *S. caroliniana* is not a member of *Sabulina* (Fig. 1).



**FIGURE 1.** Maximum likelihood tree obtained with RAxML. All clades collapsed except for *Mononeuria* and *Sabulina*. Bootstrap support values above branches only shown if  $\geq 70$ .

## Discussion

*Sabulina caroliniana* had usually been considered unrelated to any of the species now placed in *Mononeuria*. Mattfeld (1922) recognized *S. caroliniana* in sect. *Sclerophylla* Mattf. (1922: 22), where he placed each of the three species into its own series, recognizing the heterogeneous nature of the section. Small (1933) proposed to transfer *S. caroliniana* to *Sabulina* along with the five members of sect. *Uninerviae* known at that time, but segregating *S. caroliniana* and *S. stricta* (Michaux 1803: 274) Small ex Rydberg (1932: 320) from the others in an unranked group based on leaves being linear-subulate in those species. Weber (1989) published the new genus *Minuopsis* Weber (1989: 426), based on Mattfeld's sect. *Sclerophylla*. Fig. 1 shows that the three members of Mattfeld's section (and later Weber's genus), i.e. *Sabulina stricta*, *Sabulina nuttallii*, and *Sabulina caroliniana*, although all perennials with woody taproots, are not closely related.

*Mononeuria caroliniana* is found along the coastal plain of eastern North America, occupying sandy areas and oak or pine woodlands from New York south to Florida. It is most closely related to *Mononeuria groenlandica*, a mat-forming perennial ranging from Tennessee north to Greenland (and one isolated report in Brazil); in the southeastern United States, it is found on granite outcrops at higher elevations (Rabeler *et al.* 2005).

## Taxonomic Treatment

*Mononeuria caroliniana* (Walter) Dillenb. & Rabeler, *comb. nov.*

Basionym:—*Arenaria caroliniana* Walter (1788: 141).

≡ *Minuartia caroliniana* (Walter) Mattf. (1921: 28) ≡ *Sabulina caroliniana* (Walter) Small (1933: 1504).

Neotype (designated by Ward 2007):—USA. Georgia: Tattnall County, 3 mi W of Reidsville, 03 April 1948, *A. J. Cronquist* 4932 (GH non vidi, isoneotypes, FLAS51686!, MO non vidi, NY01185415!, PH00090974!, US00955677!).

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