

# Systematics and evolution of *Vaccinium* sect. *Cyanococcus* (Ericaceae): progress and prospects

PETER W. FRITSCH<sup>1\*</sup>, ANDREW A. CROWL<sup>2,3</sup>, HAMID ASHRAFI<sup>3</sup>, PAUL S. MANOS<sup>2</sup>

<sup>1</sup>Botanical Research Institute of Texas, Fort Worth, TX 76107

\*pfritsch@brit.org

<sup>2</sup>Department of Biology, Duke University, Durham, NC 27708

<sup>3</sup>Department of Horticultural Science, North Carolina State University, Raleigh, NC 27695

ORCID: PWF 0000-0002-3606-663X; AAC 0000-0002-1745-0687;

HA 0000-0002-2094-2367; PSM 0000-0003-3122-1538

## ABSTRACT

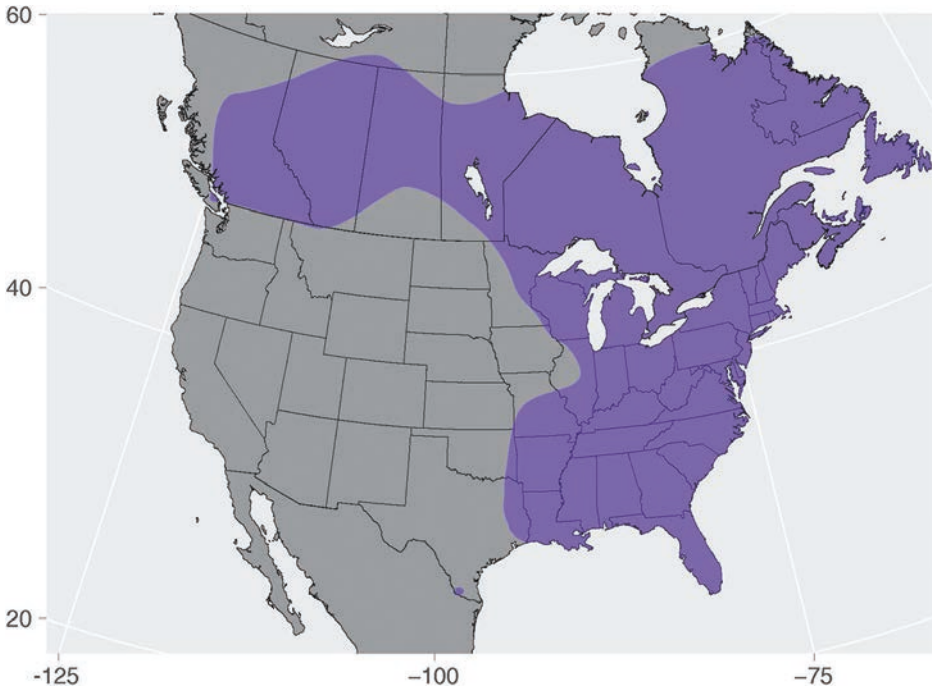
The true blueberries (*Vaccinium* sect. *Cyanococcus*; Ericaceae) comprise a clade of about nine to 24 species distributed mainly in eastern temperate North America, with one species reaching farther west. Despite extensive study, the systematics and evolution of the group are still poorly understood. Limited morphological variation, multiple ploidy levels of uncertain origin, and natural hybridization all contribute to the challenge. Questionable analytical methods, such as the use of phenetics and an overemphasis on crossing experiments, have further impeded progress. Here we review the history of research on the systematics and evolution of *V.* sect. *Cyanococcus* with the aim of clarifying and summarizing hypotheses of species origins and diversification, especially in relation to polyploidy. We also present recent progress from our own work and, on that basis, offer promising lines of investigation with morphological and molecular data. We anticipate that these avenues of research will ultimately clarify patterns of natural species diversity in *V.* sect. *Cyanococcus* with benefits for biodiversity studies, conservation, and crop breeding.

**Key words:** biosystematics, blueberries, hybridization, morphology, North America, polyploidy, systematics, taxonomy

## INTRODUCTION

*Vaccinium* section *Cyanococcus* A. Gray, the true blueberries (Ericaceae: Vaccinioideae Arn.: Vaccinieae Rchb.; henceforth “*Cyanococcus*”), is a polyploid complex of ca. nine to 24 primarily outcrossing diploid (2x), tetraploid (4x), and/or hexaploid (6x) species distributed across much of temperate North America (Camp 1945; Song and Hancock 2011; Vander Kloet 1988; Figure 1). The species of this group are of immense ecological and economic value. They form a ubiquitous component of heathlands and acidophilic plant communities and serve as a critical food source for wildlife. They also comprise the wild progenitors of most of the blueberry cultivars that form the basis of a vibrant and growing multi-billion-dollar agroindustry. Blueberries represent one of only a handful of major crop plants originating in North America.

Although the sectional limits of *Cyanococcus* are clear, the number of accepted species and their boundaries vary dramatically among the key published taxonomic treatments. Considered the current standard, the treatment in the *Flora of North America* (Vander



**Figure 1.** Geographic distribution of *Vaccinium* sect. *Cyanococcus*, endemic to North America.

Kloet 2009) is undoubtedly flawed, yet alternatives produced concomitantly or since cover only limited geographic regions or are largely recapitulations of prior work that is itself questionable. The unresolved taxonomy of *Cyanococcus* has led to confusion and inconsistency in species identification and the application of names by researchers across biological fields. It has also resulted in ambiguity in the names of *Cyanococcus* cultivars. For example, the name *Vaccinium corymbosum* L. has been used liberally for both “Highbush” and “Southern Highbush” cultivars more than one meter tall, in northern and southern climates, respectively, despite differences in genetic composition revealed through artificial hybridization with various species (Galletta and Ballington 1996).

Progress in understanding the systematics and evolution of *Cyanococcus* has thus far been reviewed primarily from the perspective of horticulture and crop breeding (e.g., Galletta and Ballington 1996; Luby et al. 1991; Song and Hancock 2011). Here we provide a review with the focus placed primarily on natural populations. We first address the taxonomic establishment and circumscription of the section and discuss the key morphological characters used in taxonomic treatments. This is followed by an overview of the two seminal monographs of *Cyanococcus*, those of W.H. Camp (1945) and S.P. Vander Kloet (1988), comparing them to each other and to more recent, mainly regional treatments, with the aim of providing clarity on the rich set of hypotheses developed by

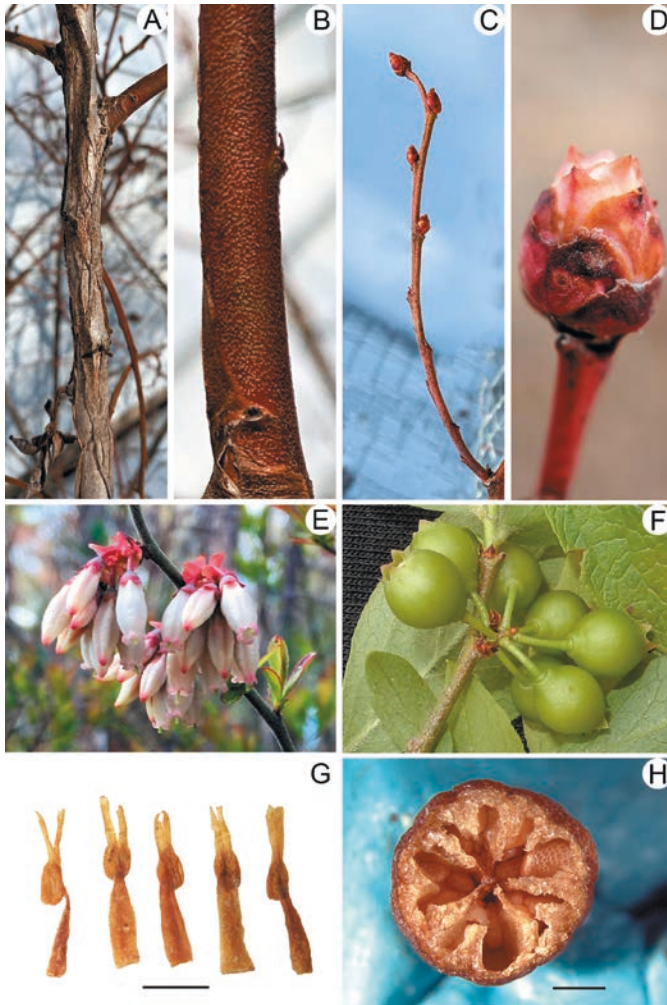
*Cyanococcus* researchers over the last eight decades that can be tested with new data. Relevant evolutionary studies conducted after the work of Vander Kloet are then summarized. We end with our own preliminary findings based on fieldwork and examination of herbarium material, and offer prospects for resolving the systematics and evolution of the group. The terminology of morphological characters and evolutionary concepts used in cited works has been updated and standardized, with changes from the original terms and concepts noted where appropriate. Phylogenetic relationships are addressed at only a cursory level here because the published data are still sparse, are primarily based on phenetics, and do not incorporate ploidy or hybridization into the analyses. We plan to focus more fully on these topics in future publications.

#### ESTABLISHMENT, RANK, POSITION, AND CIRCUMSCRIPTION OF *CYANOCOCCUS*

*Cyanococcus* was first published at the rank of section under *Vaccinium* L. (Gray 1848). It was then raised to the level of subgenus (Klotzsch 1851), and then to genus (Rydborg 1917, although as the illegitimate name *Cyanococcus* (A. Gray) Rydb. non *Cyanococcus* A. Hansgirg publ. 1905; International Plant Names Index 2021). The latter rank was accepted by Small (1933), but Camp (1945) considered it to “probably [merit] the rank of subgenus under *Vaccinium*.” The key treatments that include *Cyanococcus* published since have all recognized it as a section of *Vaccinium* (e.g., Sleumer 1941; Stevens 2004; Vander Kloet 1983, 1988; Vander Kloet and Dickinson 2009). Nonetheless, *Vaccinium* has been considered “wildly polyphyletic” (Stevens 2004), and the available molecular phylogenetic data have supported para- or polyphyly for the genus (Powell and Kron 2002). This has led to proposals that divide *Vaccinium* into various numbers of genera, many corresponding to current sections. In the most extreme of these proposals, *Vaccinium* would be reduced to the single species *V. uliginosum* L. (W.S. Judd, University of Florida, pers. comm.), in which case *Cyanococcus* would be recognized again at the level of genus (although with a new name). However, much greater taxon sampling and improved molecular phylogenetic data will need to be generated and analyzed before seriously contemplating such changes.

*Cyanococcus* fits easily into the tribe Vaccinieae within the Ericaceae by its inferior ovary (Stevens 2004). Other characters that, in combination, place the section in this tribe are abortive pseudoterminal vegetative buds (versus non-abortive), presence of anther tubules (versus absence), and baccate fruits (versus dry; Stevens 2004). The phylogenetic placement of *Cyanococcus* within the Vaccinieae is currently unclear and, as with rank, will require much more extensive sampling within the tribe and improved molecular data than has thus far been applied.

As to circumscription, Gray (1859, 1860) and Sleumer (1941) placed several Japanese and northeastern Asian species in *Cyanococcus*, and Sleumer (1941) erected *Vaccinium* sect. *Pseudocyanococcus* Sleumer for the strongly evergreen species *V. myrsinites* Lam.; these placements have been rejected by subsequent authors. Odell et al. (1989) presented convincing evidence for the current circumscription of the section based on the unique characters within *Vaccinium* of eruptive periderm development, resulting in distinctive elongate diamond-shaped bark patterns and raised branchlet stomata, rendering the branchlets verrucose (Figure 2A, 2B). Several other characters in combination also serve



**Figure 2.** Some diagnostic morphological characters of *Vaccinium* sect. *Cyanococcus*. Species names are tentative, approximating those of Camp (1945). **A.** Eruptive periderm on the bark of a cultivated plant of *V.* sect. *Cyanococcus*. **B.** Raised stomata on the second-year stems of a cultivated plant of *V.* sect. *Cyanococcus*. **C.** Dimorphic buds on a cultivated plant of *V.* sect. *Cyanococcus*. The four larger more distal buds are reproductive buds and the smaller more proximal buds are vegetative. **D.** Flowering bud showing multiple overlapping bud scales. **E.** *V. tenellum*, showing multi-flowered inflorescences. **F.** Infructescence of *V. amoenum*/*V. virgatum* showing the articulation between the pedicel apex and hypanthium, and well-developed calyx lobes. **G.** Stamens of *V. constablaei*. Note the absence of dorsal anther spurs. *Manos CY-442* (BRIT, DUKE). Scale bar = 2 mm. **H.** Cross section of young fruit of *V. amoenum*/*V. virgatum* with upper immature seeds removed showing architecture of pseudo-10-locular ovary. The five septa alternate with five invaginations from the outer wall. *Fritsch 2282* (BRIT, DUKE). Scale bar = 2 mm. Photos: A–C by Anna Becker, used with permission; D, derivative of <https://www.inaturalist.org/observations/41521073> by Nathaniel Sharp licensed under CC BY-NC 4.0; E by A.A. Crowl; F–H by P.W. Fritsch.

to diagnose the section, for example, dimorphic vegetative and flowering buds, flowering buds with more than two overlapping scales, flowers in short racemes, the presence of an articulation between the pedicel apex and hypanthium (ovary), well-developed calyx lobes, the absence of anther spurs, and a pseudo-10-locular ovary (Camp 1945; Vander Kloet 1983; Figure 2C–H). A phylogenetic study based on DNA sequence data from the nuclear ITS and plastid *matK* and *ndhF* regions, which included samples of many sections of *Vaccinium*, supports the monophyly of the section, albeit with only two species of *Cyanococcus* sampled (Powell and Kron 2002). Phylogenomic analysis based on high-throughput DNA sequence data with more species of *Cyanococcus* included also supports the monophyly of the section (Crowl et al. 2022; A.A. Crowl et al. unpubl. data).

#### MORPHOLOGICAL CHARACTERS USED IN THE TAXONOMY OF *CYANOCOCCUS*

One of the main challenges in resolving the systematics and evolution of *Cyanococcus* has been the relative paucity of distinct morphological characters for use in species delimitation. The characters that have been used to delimit taxa often overlap from taxon to taxon and some are often tightly correlated (e.g., the color of the leaves, pedicels, and hypanthium). The key morphological characters that have generally been used in taxonomic treatments of *Cyanococcus* are as follows. Habit: plant height, shape; extent of clonality (single above-ground stems vs. several stems together or many stems forming colonies). Branchlet: density; color; angularity; presence, arrangement, and length of surface trichomes. Vegetative bud: color; size. Leaf: persistence. Leaf blade: color on abaxial and adaxial surfaces; sheen on the abaxial surface; shape; dimensions; presence, density, and length of simple trichomes on the abaxial surface; presence of stipitate-glandular trichomes on the abaxial surface; presence, orientation, and length of trichomes on midvein on the abaxial surface; presence and regularity of (stipitate-glandular-tipped) serrations on the margin. Flower: density per inflorescence. Pedicel and calyx: color; presence of simple and stipitate-glandular trichomes. Corolla: color; shape; dimensions; presence of stipitate-glandular trichomes on the abaxial surface. Stamens: presence and arrangement of simple trichomes on the filaments; diameter of pollen tetrads. Fruit: color; sheen; dimensions; presence and quantity of surficial wax (“bloom”); seed weight.

Most of these characters are fully retained on herbarium specimens. For those that are not, we implore collectors to supply field information with the specimen. Particularly important are the following: plant clonality and diameter of the clone when present (care should be taken to distinguish among physically overlapping clones as assessed by, for example, consistently different leaf blade shapes); plant height, and the number of basal stems; colors of various organs (branchlets, leaves, calyx, corolla [note any patterns such as lines], fruit); corolla shape, which can be distorted on dried specimens through, for example, the splitting of campanulate or urceolate corollas; the presence or absence, and degree of, wax deposits on the calyx and fruit (“bloom”); fruit sheen (dull or glossy); and the orientation of flowers and fruit. It is also important to indicate plant species associates, especially the presence of other species of *Cyanococcus* growing with the collected plant. Separate collections should be made to document local variation in morphology, or suspected hybrids or introgressants. Collections of the same plant should ideally be made at



different times of the season because flowers and fruits typically occur asynchronously in populations, and flowering often occurs prior to full leaf expansion. Finally, care should be taken to note whether any suckering branchlets have been collected because the leaves on these branchlets can differ from those on normal branchlets in at least shape and dimensions.

#### TAXONOMIC TREATMENTS

Critical taxonomic work on *Cyanococcus* can be divided into three segments roughly corresponding to time intervals. Before this work, the only group-wide treatments of *Cyanococcus* were produced as part of general floristic accounts for all or part of North America (e.g., Britton and Brown 1913; Chapman 1897; Robinson and Fernald 1908; Small 1933). The first two segments comprise the *Cyanococcus*-wide works of W.H. Camp in the 1940s and those of S.P. Vander Kloet mainly in the 1970s through 1990s, respectively. Both of these investigators conducted their research in the overall context of crop improvement, applying data from experimental crossing studies, genetics, and cytology to questions of species delimitation and the evolution of natural populations. These individuals produced a wealth of research that has advanced the knowledge of *Cyanococcus* substantially, although the focus on horticulture appears to have too strongly favored evidence relevant for crop breeding such as crossing information and fruit quality. The third segment consists of treatments of *Vaccinium*, including *Cyanococcus*, for specific U.S. states, written partly in response to those of Camp and/or Vander Kloet.

**W.H. Camp.** Comprehensive taxonomic study of *Cyanococcus* began in earnest with the work of W.H. Camp (Figure 3A) while he was an herbarium curator at the New York Botanical Garden. Although lesser known than some of the other botanists contributing to the evolutionary synthesis such as G.L. Stebbins and E. Anderson, Camp was nonetheless influential in the development and promulgation of biosystematics (i.e., the study of taxonomy and classification based directly on evolutionary principles and processes) as applied to botany (Kleinman 2009). Under the tenets of the biological species concept (Mayr 1942, 1963), the plant biosystematists used data from morphology, experimental crossing studies, genetics, and cytology to assess the reproductive isolation of individuals and populations emphasized for species delimitation, as well as to disentangle genetic from environmental influences on morphology and physiology. Although the theory of phylogenetic classification had yet to be fully developed, biosystematic classifications can be considered phylogenetically based (e.g., species were placed along or terminating lines of descent).

Through prior studies (Camp 1942; Camp and Gilly 1943; Darrow and Camp 1945) but with much new information, Camp (1945) developed the first comprehensive taxonomic treatment of *Cyanococcus*; this was also one of the early taxonomic products of the plant biosystematists. *Cyanococcus* was divided into 24 species, with many earlier species and varietal names placed into synonymy (Table 1; Supplementary Data). Hypotheses were proposed only for the origins of most of the tetraploids and hexaploids; the origins of the diploids and the tetraploid *Vaccinium hirsutum* Buckley were left unspecified. Camp strongly emphasized estimates of ploidy derived from chromosome squashes to delimit species boundaries. No species possesses more than



**Figure 3.** Authors of key taxonomic treatments of *Vaccinium* sect. *Cyanococcus*. **A.** W.H. Camp. **B.** S.P. Vander Kloet. **C.** D.B. Ward. **D.** L.J. Uttal. A, courtesy of the Archives of the New York Botanical Garden; B, courtesy of Melanie Priesnitz, used by permission; C, courtesy of Gordon Ward, used by permission; D, courtesy of Jeannie Uttal Breeden, used by permission.

**Table 1.** Comparison of the two major taxonomic treatments of *Vaccinium* sect. *Cyanococcus*, that of Camp (1945) and Vander Kloet (2009), with their hypotheses for the origin of the tetraploids (4x) and hexaploids (6x) indicated. The terms “highbush” and “lowbush” are used in the sense of Vander Kloet as > 1 m tall versus < 1 m tall, respectively; auto = autopolyploid; allo = allopolyploid. See Supplementary Data for additional notes.

Camp (1945) species	Camp hypotheses	Vander Kloet (2009) species	Vander Kloet hypotheses
<b>“Lowbush”</b>			
<i>V. angustifolium</i> Aiton (2x)	—	<i>V. angustifolium</i> (4x)	Allo of <i>V. boreale</i> and <i>V. pallidum</i>
<i>V. darrowii</i> Camp (2x)	—	<i>V. darrowii</i> (2x)	—
<i>V. myrtilloides</i> Michx. (2x)	—	<i>V. myrtilloides</i> (2x)	—
<i>V. pallidum</i> Aiton (2x)	—	<i>V. pallidum</i> (2x)	—
<i>V. tenellum</i> Aiton (2x)	—	<i>V. tenellum</i> (2x)	—
<i>V. vacillans</i> Kalm ex Torr. (2x)	—	<i>V. pallidum</i> (2x)	—
<i>V. altomontanum</i> Ashe (4x)	Polyphyletic auto of <i>V. vacillans</i>	<i>V. pallidum</i> (4x)	No hypothesis indicated
<i>V. brittonii</i> Porter ex E.P. Bicknell (4x)	Auto of <i>V. angustifolium</i> or sister of <i>V. lamarckii</i>	<i>V. angustifolium</i> (4x)	Allo of <i>V. boreale</i> and <i>V. pallidum</i>
<i>V. hirsutum</i> Buckley (4x)	No hypothesis specified	<i>V. hirsutum</i> (4x)	No hypothesis specified
<i>V. lamarckii</i> Camp (4x)	Auto of <i>V. angustifolium</i>	<i>V. angustifolium</i> (4x)	Allo of <i>V. boreale</i> and <i>V. pallidum</i>
<i>V. myrsinites</i> Lam. (4x)	Segmental allo of <i>V. darrowii</i> × <i>V. tenellum</i>	<i>V. myrsinites</i> (2x, 4x)	4x is allo of <i>V. tenellum</i> × <i>V. darrowii</i>
Not treated	—	<i>V. boreale</i> I.V. Hall & Aalders (2x)	—
<b>“Highbush”</b>			
<i>V. atrococcum</i> A. Heller (2x)	—	<i>V. corymbosum</i> (2x, 4x, 6x)	Complex allo of “lowbush” diploids; a “compilospecies”
<i>V. caesariense</i> Mack. (2x)	—	“	“
<i>V. elliotii</i> Chapm. (2x)	—	“	“
<i>V. arkansanum</i> Ashe (“probably 4x”)	Theoretical auto of <i>V. atrococcum</i>	“	“
<i>V. australe</i> Small (= <i>V. formosum</i> Andrews; 4x)	Auto of <i>V. caesariense</i>	“	“
<i>V. corymbosum</i> L. (4x)	Complex allo of six tetraploids	“	“
<i>V. fuscatum</i> Aiton (“probably 4x”)	Theoretical allo of <i>V. atrococcum</i> × <i>V. darrowii</i>	“	“
<i>V. marianum</i> P. Watson (“probably 4x”)	Theoretical allo of <i>V. atrococcum</i> × <i>V. caesariense</i>	“	“
<i>V. simulatum</i> Small (4x)	Auto of <i>V. pallidum</i>	“	“
<i>V. virgatum</i> Aiton (4x)	Auto of <i>V. tenellum</i>	“	“
<i>V. amoenum</i> Aiton (6x)	Auto of <i>V. tenellum</i> / <i>V. virgatum</i>	“	“
<i>V. ashei</i> J.M. Reade (6x)	Complex allo of five tetraploids	“	“
<i>V. constablaei</i> A. Gray (6x)	Allo of <i>V. altomontanum</i> × <i>V. simulatum</i>	“	“



one ploidy level in the treatment, and species often form ancestor-descendant diploid ( $2x$ , with  $x = 12$ )–tetraploid ( $4x$ )–hexaploid ( $6x$ ) series (the “euploidion” of Camp and Gilly 1943) that either branch or converge with other species or series, thus forming reticulate patterns of relationship (the “polyploid complex” of Grant 1981). Some species were thought to have originated through autopolyploidy and others through allopolyploidy (terms as in, for example, Grant 1981; Spoelhof et al. 2017; Stebbins 1950), and two (the tetraploid *V. corymbosum* and hexaploid *V. ashei* J.M. Reade) were considered to combine parts of the genomes of several ancestral species.

Today the categories “highbush” and “lowbush” are applied to various *Cyanococcus* species in the horticultural literature, usually with the additional category “rabbit-eye” included and “highbush” often divided into “southern highbush” and “northern highbush” (e.g., Galletta and Ballington 1996; Luby et al. 1991; Vander Kloet 1983; Weakley 2020). However, Camp generally avoided the term “lowbush” and applied “highbush” only to those species more than one meter tall that he considered being involved in the evolution of *Vaccinium corymbosum*. An exception is the use of “lowbush” in the general key to species, but it appears that Camp used it merely as an artificial means of identification, with the division at one meter. Camp also referred frequently to “halfhigh” plants, which supposedly resulted from various crosses of “lowbush” and “highbush” types.

Camp ascribed his observations of wide variation within and among the species of *Cyanococcus* mainly to hybridization and introgression, and less commonly to segmental allopolyploidy. Based largely on the observation that viable hybrids are freely produced experimentally from parental species with the same ploidy level (either diploid, tetraploid, or hexaploid; Darrow and Camp 1945), Camp concluded that natural hybridization among the species of *Cyanococcus* is rampant and assumed that morphological similarity between experimentally produced hybrid progeny and wild individuals demonstrated that the latter are hybrids. Camp also thought that at least some of the allopolyploidization events produced “segregative allopolyploids” ( $\pm$  equivalent to “segmental allopolyploids” of, for example, Grant 1981; Mason and Wendel 2020; Stebbins 1950); that is, allopolyploids in which the morphology of progeny tends toward that of one or the other parent through genetic recombination. Individuals with morphology ranging outside the norm of a species and resulting from hybridization, introgression (although not termed as such), and/or segmental allopolyploidization were designated as “elements” or “phases” (e.g., “the ‘simulatooid’ element of *Vaccinium corymbosum*” or “the ‘alto-montanoid’ element of *V. constablaei*” A. Gray). Camp also considered the boundary corresponding to the edge of the ice sheet during the last glacial maximum to have significantly influenced the current geographic distribution of the “highbush” species of *Cyanococcus*, such that distributional information relative to this boundary was included in the “Key to the basic populations” of species, sometimes to the exclusion of morphology. Camp proposed that interbreeding does not swamp species boundaries in *Cyanococcus* because hybrid establishment mainly occurs in habitats modified by humans, whereas in undisturbed areas the species were considered to be more fully separated by habitat and thus tend to breed true.

Camp’s treatise is at once comprehensive, groundbreaking, and thought-provoking. The work offers a wealth of information, including an abundance of hypotheses on *Cyanococcus* species boundaries and evolution that can be further refined and tested

with new data. Despite the significant advance represented by Camp's treatment, the overall value of the work is nonetheless compromised in several ways, which must be noted for a complete understanding of its implications. First, other than types, herbarium specimens are not cited; this renders the (somewhat abbreviated) species descriptions, discussions, and figures as the only means for the reader to obtain information on species boundaries, infraspecific and interspecific variation, hybridization, and introgression. Furthermore, the few figures in the treatment consist merely of low-resolution photographs of some of the species in the living condition. Second, if one is to assume that the work of Darrow et al. (1944) represents the entirety of the count data on which the species chromosome numbers are based, then the determination of ploidy is apparently based on only one to several counts for each species. Moreover, Darrow et al. (1944) do not clearly distinguish between chromosome counts made from cultivated versus wild-collected material. Third, three of the species are provided with an indication of a ploidy level (all tetraploids) that is merely a prediction based on morphology versus an empirical chromosome count; thus, the hypothesized origin of these species, and even their very existence, must be questioned. Fourth, assertions of natural hybridization and introgression are inadequately documented (i.e., presented without supporting data). Fifth, the evolutionary origin and diversification of many of the species are too often presented as substantiated fact when the evidence was scant at best, including highly speculative movements of ancestral species in response to climatic and glaciation events.

**S.P. Vander Kloet.** The work of Camp was used as the basis for further investigating the taxonomy and evolution of *Cyanococcus* by S.P. Vander Kloet, professor of biology at Acadia University and curator of its E.C. Smith Herbarium (Figure 3B). Like Camp, Vander Kloet relied heavily on experimental crossing studies as a basis for taxonomic conclusions in *Cyanococcus*. In contrast to Camp, he employed morphological phenetic analyses based on the principles of numerical taxonomy (Sneath and Sokal 1973); as such, he rejected the use of phylogenetics in estimating lines of descent and delimiting species boundaries a priori. After contributing numerous publications on the taxonomy and evolution of *Cyanococcus* (Vander Kloet 1976a, 1976b, 1977a, 1977b, 1978a, 1978b, 1980, 1983), Vander Kloet produced a culminating treatment of the group as part of a monograph of *Vaccinium* in North America, recognizing nine species of *Cyanococcus* (Vander Kloet 1988). Like the works of Camp, those of Vander Kloet often suffer from inadequate evidence-based conclusions. This applies particularly to assertions of natural hybridization and introgression but extends to species circumscription as well, because only rarely are herbarium specimens cited. The data from experimental crosses are too often presented anecdotally and backed neither by published data nor cited herbarium vouchers. For example, the published works often lack the data sets on which phenetic analyses are based.

The most prominent element of Vander Kloet's revisionary work (see particularly Vander Kloet 1980) based on the above approaches was the conclusion that all *Cyanococcus* species more than one meter tall ("highbush") form a "compilospecies," that is, a genetically aggressive species that acquires the heredities of closely related sympatric species through hybridization and introgression (Harlan and deWet 1963). Vander Kloet proposed that this compilospecies, *Vaccinium corymbosum*, originated from hybridization

between *V. darrowii* Camp and *V. tenellum* Aiton, and that it subsequently acquired and is still acquiring characters from the other diploid “lowbush” blueberries in his treatment (i.e., *V. boreale* I.V. Hall & Aalders, *V. pallidum* Aiton, and *V. myrtilloides* Michx.), through repeated crossing, introgression, and polyploidization where they contact each other. As such, *V. corymbosum*, now the sole “highbush” species in *Cyanococcus*, was considered a highly variable species both morphologically and genetically, arising through multiple origins and comprising diploids, tetraploids, and hexaploids. Consequently, 12 of Camp’s species were placed in synonymy (Table 1).

As evidence for justifying this concept of *Vaccinium corymbosum*, Vander Kloet cited the clustering of the “highbush” samples in his phenetic analysis relative to the “lowbush” samples, as well as the general lack of internal resolution within the “highbush” cluster. Because the character matrix for this analysis was not provided in the publication, it is difficult to independently assess these conclusions. Vander Kloet considered the findings of significantly fewer stainable pollen tetrads in “highbush” tetraploids versus “lowbush” diploids, and progeny from seed collected in the field from a single open-pollinated shrub that only partially resembled the parent, further evidence of a hybrid origin of *V. corymbosum*. It is unfortunate that of the 56 populations collected, Vander Kloet detailed only one example of the latter observation—a hybrid between plants corresponding to the diploids *V. atrococcum* A. Heller and *V. caesariense* Mack. sensu Camp (1945). The rest of the data are merely summarized by the statement, “Similar results were obtained from the other seed collections acquired and treated in the same way as described above . . .” Vander Kloet’s discussion implies that tetraploid individuals and populations of *V. corymbosum* have been repeatedly produced from the “lowbush” diploids, and hexaploids from tetraploid  $\times$  diploid crosses, although the crossing data from hexaploids were sparse and therefore excluded from the study.

Based on observed habitat and phenological differences between *Vaccinium angustifolium* Aiton and *V. corymbosum*, Vander Kloet (1976a) suggested that the other members of *Cyanococcus*, although still hybridizing in nature, are sufficiently isolated reproductively so as to be regarded as species. Thus, in at least one instance, Vander Kloet used ploidy to distinguish between morphologically similar species (diploid *V. boreale* versus tetraploid *V. angustifolium*; Vander Kloet 1977a). This may be one reason why, in contrast to Camp, Vander Kloet (1977b) asserted that “... the frequency of naturally occurring hybrids in areas of sympatry is much lower than expected from ... experimental data. Indeed, ... hybrids are rare,” which ironically seems to contradict his concept of *V. corymbosum* as a compilespecies formed through hybridization.

In addition to recircumscribing *Vaccinium corymbosum*, Vander Kloet modified the concepts of *V. angustifolium*, *V. pallidum*, and *V. myrsinites* sensu Camp (see Supplementary Data). *Vaccinium angustifolium* is generally equivalent to Camp’s concept of *V. lamarckii* Camp as altered to become the sole species throughout the geographic range encompassing Camp’s *V. brittonii* Porter ex E.P. Bicknell and *V. lamarckii* at the tetraploid level, whereas *V. boreale* is reserved for plants that are morphologically similar to *V. angustifolium* but diploid. Aalders and Hall (1963) showed that the diagnostic characters of *V. brittonii* (i.e., glaucous stems and leaves) are controlled by a single gene and exposure to sunlight (Aalders and Hall 1963), and Vander Kloet (1978a) demonstrated

that the characters defining *V. brittonii* and *V. lamarckii* do not breed true and co-occur extensively without phenological differences (Vander Kloet 1978a).

As reviewed in Luby et al. (1991), Aalders and Hall (1962) documented a positive correlation between leaf stomatal size and ploidy in *Vaccinium angustifolium* and their recently described species *V. boreale*. They measured the lengths of stomata in 12 individuals each of diploids and tetraploids consistent with *V. angustifolium* in morphology, finding a positive correlation between stomatal length and ploidy. They then measured the (presumed) average stomatal length from the type specimen of *V. angustifolium*, finding that it occurred well outside the range of stomatal lengths of the diploids but within the range of the tetraploids (the statistical basis for this conclusion was not presented). Thus, tetraploidy was ascribed to the type of *V. angustifolium*, which justified the continued recognition of their new species *V. boreale* based on its diploid ploidy level and several morphological characters; that is, corolla length (3.0–3.5 mm versus 5–7 mm), pollen tetrad size (35–40  $\mu\text{m}$  versus 40–43  $\mu\text{m}$ ), and a higher degree of branching (Hall and Aalders 1961). Hall and Aalders (1961) stated that no other vegetative characters could be found to distinguish these species with confidence.

Vander Kloet (1988) recognized *Vaccinium boreale* but, in contrast to Hall and Aalders (1961), only used smaller plant height and smaller leaf blade size in his key to distinguish it from *V. angustifolium* with morphology; this is despite the clearly different corolla lengths presented in his descriptions (3–4 mm versus 4–6 mm, respectively). Vander Kloet (1977a) asserted that vegetative characters could be used to distinguish the two species but we consider the data presented too imprecise for properly assessing this claim. In particular, the extent to which the plants used for the morphological measurements were assessed for ploidy is unclear. *Vaccinium boreale* sensu Vander Kloet can be considered equivalent to *V. angustifolium* sensu Camp except for a much more restricted geographic range, occurring only in far eastern Canada south through the northern border area of the northeastern United States (see Supplementary Data). In related research, Vander Kloet (1977a) cited the similarity of progeny to *V. angustifolium* in experimental crosses between *V. boreale* (2x) and *V. pallidum* (2x) in proposing that tetraploid *V. angustifolium* originated from hybridization between these two species, albeit the progeny of the crosses were all diploid.

Vander Kloet (1978b) presented convincing evidence from morphology and crossing experiments for placing *Vaccinium vacillans* Kalm ex Torr. into the synonymy of *V. pallidum*. In sampling the ploidy of individuals from several populations, he found that, out of 119 individuals sampled for chromosome number, 93% were diploid and 7% tetraploid. He thus confirmed the presence of tetraploid individuals in *V. pallidum*-like individuals as found by Camp (1945; no herbarium vouchers or images of chromosome squashes were cited). However, Vander Kloet chose to synonymize the tetraploids under *V. pallidum*, unlike Camp, who apparently recognized them as *V. altomontanum* Ashe (the application of the name *V. altomontanum* is unresolved because its type appears to be missing). Vander Kloet found no morphological characters to distinguish the tetraploids from the diploids and no evidence to indicate that the tetraploids behave ecologically as a distinct species. Although Vander Kloet did not specify a hypothesized origin of the

tetraploids, it appears that he agreed with Camp that they have been produced as polyphyletic autopoloids of the diploids.

The main difference in the concept of *Vaccinium myrsinites* from that of Camp is that both the diploid and tetraploid levels are cited, without explanation, in Vander Kloet (2009), although only the diploid level is cited in Vander Kloet (1983, 1988). Vander Kloet apparently considered the tetraploid level to have originated in the same way as did Camp, that is, from a cross between *V. darrowii* and *V. tenellum*.

Vander Kloet's taxonomic view of *Cyanococcus* is currently considered the standard, having been adopted by the U.S. Department of Agriculture, plant breeders, and many local and regional floras, including the *Flora of North America* (Vander Kloet 2009). A key consequence is that the terms "highbush" and "lowbush" have become firmly established in distinguishing *Cyanococcus* species on the basis of plant height, that is, generally more than one meter tall (*Vaccinium corymbosum* and any segregates) versus less than one meter tall (remaining species; e.g., Weakley 2020). Despite his assertion of widely occurring hybridization among the diploid "lowbush" members of *Cyanococcus* resulting in the polymorphic and polyphyletic *V. corymbosum*, Vander Kloet (1983, 1988) considered the diploids to have largely parapatric geographic distributions, with each of the species flanked by only one or two others.

**Key Treatments for U.S. States.** Just prior to and since the time of Vander Kloet's work, two researchers with field and herbarium experience in *Vaccinium* each produced a taxonomic treatment of *Cyanococcus* for a U.S. state as part of a more extensive treatment of *Vaccinium*. Although constrained by geography, their studies often address more general taxonomic problems in *Cyanococcus* and thus achieve relevance here.

D.B. Ward of the University of Florida (Figure 3C) published a treatment of *Vaccinium* for the state of Florida (Ward 1974). Ward compared his work to that of Camp (1945) and others (Correll and Johnston 1970; Rehder 1940; Small 1933; Vander Kloet had not yet begun to publish on *Vaccinium*). Ward summarized the contrast between his conclusions and those of Camp with the statement that *Cyanococcus* is "... difficult but not in any way an irresolvable tangle of intergrading populations." Nonetheless, Ward seems to have oversimplified Camp's taxonomy by not addressing issues regarding potential variation in ploidy among morphologically similar plants. Ward recognized *V. amoenum* Aiton, *V. ashei*, *V. australe* Small, *V. darrowii*, *V. elliotii* Chapm., *V. fuscatum* Aiton, *V. myrsinites*, and *V. tenellum* for Florida. Differences from Camp's treatment are noted below. Ward did not habitually indicate ploidy for the species entries; ploidy is thus indicated below only when noted in the treatment.

Ward agreed with Camp in distinguishing *Vaccinium amoenum* (6x) from *V. virgatum* Aiton (4x; not occurring in Florida according to Ward), differentiating the two by the larger habit, leaves, and flowers of *V. amoenum*, as well as inflorescences that are limited to no more than two or three successive axils per branchlet (versus small panicle clusters in the axils of many successive leaves per branchlet) and more widely separated from each other.

Ward considered *Vaccinium ashei* sensu Camp to be "... perhaps less of a taxonomic category than it is a philosophical concept," yet Ward recognized it, apparently in



deference to Camp. It was distinguished from *V. amoenum* by leaves with abaxial surfaces that are glabrous or pubescent only along the midvein and that are more sparsely stipitate glandular.

*Vaccinium arkansanum* Ashe and *V. atrococcum* were combined with *V. fuscatum*, with Ward stating that although ploidy differs within this group, on a practical basis (we suppose meaning morphologically) the species cannot be distinguished. Ward differentiated *V. fuscatum* from *V. australe* by pubescent leaves and branchlets and black non-glaucous fruit (versus glabrous leaves and branchlets and blue-glaucous fruit) without commenting on or distinguishing among ploidy levels. Apparently, Ward did not consider *V. fuscatum* to be a hybrid of *V. atrococcum* and *V. darrowii* (unlike Camp), because at the end of the entry for *V. fuscatum* he stated that “Hybrids of *V. fuscatum* with ... *V. darrowii*[i] are rather frequent,” yet *V. atrococcum* was placed in synonymy with *V. fuscatum*.

There is essentially no difference in Ward’s concept of *Vaccinium tenellum* from that of Camp, although Ward and Lyrene (2007) provided evidence to show that *V. tenellum* does not occur in Florida. They stated that prior herbarium specimen documentation of the species from the state was based on misidentifications of *Gaylussacia dumosa* (Andrews) A. Gray.

L.J. Uttal of Virginia Polytechnic Institute and State University (Virginia Tech; Figure 3D) published a treatment of *Vaccinium* for the state of Virginia. The treatment was produced partly to counter the concept of Vander Kloet’s circumscription of the single highly variable species of “highbush” blueberry *V. corymbosum* (Uttal 1987), although Uttal readily adopted other species of Vander Kloet and their circumscriptions. Uttal agreed with Ward (1974) in segregating *V. elliotii*, *V. virgatum* (although thought not to occur in Virginia), and other species recognized by Camp (1945) from *V. corymbosum*. Uttal’s work was based entirely on morphology and for the most part did not include quantitative data. Below we summarize and comment on the species of *Cyanococcus* in Virginia sensu Uttal:

There are essentially no differences between Uttal’s and Camp’s concepts of *Vaccinium caesariense* (2x), *V. elliotii* (2x), *V. myrtilloides* (2x), *V. simulatum* Small (4x), and *V. tenellum* (2x).

*Vaccinium angustifolium* (4x). Uttal adopted Vander Kloet’s concept of this species.

*Vaccinium corymbosum* (4x). This species sensu Uttal best fits the concept of Camp except that *V. constablaei* (6x) is placed in synonymy. Uttal considered *V. constablaei* not to occur in Virginia, but considered it unreliably distinguishable from *V. corymbosum* with morphology.

*Vaccinium formosum* Andrews (4x). Uttal (1986) considered *V. formosum* to be an earlier name and thus the correct name for Camp’s *V. australe*.

*Vaccinium fuscatum* (2x, 4x). Uttal differed from Vander Kloet in recognizing this species as distinct from *V. corymbosum*. However, he also diverged from Camp (and apparently agreed with Ward [1974]) in adopting it as a species versus a hybrid between *V. atrococcum* (which he placed in the synonymy of *V. fuscatum*) and *V. darrowii*. He based this decision on the type material of *V. atrococcum*, which he asserted

could not have been collected any farther south than South Carolina, whereas Camp's concept of *V. fuscatum* applies exclusively to plants from southern Georgia and Florida. According to Uttal, if the latter plants referred to by Camp merit recognition then they require a new name. Ironically, however, Uttal mentioned that the small leaves of the type of *V. holophyllum* (Small) Uphof suggest a hybrid between *V. fuscatum* and *V. darrowii*, which would in fact match Camp's concept of *V. fuscatum* as (an allotetraploid) hybrid between *V. atrococcum* and *V. darrowii*.

*Vaccinium*  $\times$  *marianum* P. Watson. Uttal employed a nothospecies name for this hybrid while leaving other recognized hybrids unnamed, apparently because Uttal considered morphological intermediates of *V. formosum* and *V. fuscatum* to be the most common hybrids in the Virginia flora. Whereas Camp specified *V. marianum* as an allotetraploid of diploid *V. caesariense* (2x) and *V. atrococcum* (2x), Uttal clarified neither the ploidy level(s) at which hybridization occurred, nor the resulting ploidy level post-hybridization. Because Uttal considered *V. fuscatum* to comprise both 2x and 4x individuals, this aspect of *V.  $\times$ marianum* remains unresolved in his treatment.

*Vaccinium pallidum*. There is essentially no difference in the concept in this species from that of Vander Kloet, with both diploids and tetraploids included. *Vaccinium altomontanum*, apparently sensu Camp (1945; 4x), was placed in synonymy, as was *V. vacillans*.

**Other treatments.** Taxonomic studies of *Cyanococcus* published after Uttal's treatment provide additional insights but are limited in scope. Three principal regional treatments of *Cyanococcus* focusing on the southeastern U.S. largely reflect the concepts of one or more of the prior global or state treatments above. The treatment of Godfrey (1988) for northern Florida and adjacent Georgia and Alabama followed "with not a little reluctance" Vander Kloet's (1988) treatment except that *Vaccinium elliottii* was considered distinct enough morphologically to warrant separate species recognition. The treatment of Luteyn et al. (1996) for the whole of the southeastern U.S. entirely reflects Vander Kloet's treatment, undoubtedly because, according to the Acknowledgments section, Vander Kloet was responsible for the treatment of *Vaccinium* in that work. The treatment of Weakley (2020) for the southeastern U.S. comprises elements of Camp, Vander Kloet, and Uttal, while also according with Ward's observation that hybrids are not nearly as frequent as asserted by Camp and Vander Kloet. Weakley's treatment recognizes 15 species, including *V. angustifolium*, *V. caesariense*, *V. corymbosum*, *V. darrowii*, *V. elliottii*, *V. fuscatum*, *V. formosum*, *V. hirsutum*, *V. myrsinites*, *V. myrtilloides*, *V. pallidum*, *V. tenellum*, and *V. virgatum*. The treatment deviates from prior treatments in also recognizing *V. altomontanum* and *V. simulatum*, while placing *V. constablaei* in synonymy with *V. corymbosum*. The differences among treatments generally in the latter regard, at least in part, likely result from the missing type of *V. altomontanum* together with misapplications of one or more of these names, but careful study will be needed to address this problem.

In other work, Luby et al. (1991) and Galletta and Ballington (1996) stated that sufficient morphological and ecological separation exists to recognize *Vaccinium elliottii*, and possibly *V. constablaei* and *V. simulatum*, all sensu Camp (1945), while leaving the rest

of *V. corymbosum* sensu Vander Kloet (1983, 1988) intact. Luby et al. (1991) thought it best to consider the state of knowledge of the taxonomy of *Cyanococcus* (at the time) preliminary, urging further studies with multiple approaches. Various blueberry breeders have considered *V. fuscatum* (defined as diploid, in contrast to Camp 1945) and *V. ashei* (6x but “more properly denoted *V. virgatum*”) to be sufficiently distinct from *V. corymbosum* sensu Vander Kloet to justify their recognition (Song and Hancock 2011).

#### EVOLUTIONARY STUDIES ON *CYANOCOCCUS* AFTER VANDER KLOET

Various studies investigating the nature of polyploidy and species relationships in *Cyanococcus* have been conducted since the time of Vander Kloet. These are addressed in turn below.

**Ploidy.** The original observation of Camp and others that *Vaccinium* species of the same level of ploidy can generally be crossed successfully under artificial conditions has been corroborated (with some exceptions; see Ballington and Galletta [1978]). Because of the strong triploid and pentaploid block in *Cyanococcus*, crosses among species of differing ploidy are only rarely successful (Hancock 1998; Lyrene et al. 2003; Ortiz et al. 1999). Nonetheless, the rare exceptions, thought to result from the production of unreduced gametes at low frequency (Ortiz et al. 1992a, 1992b), are exploited in blueberry breeding in the development of new cultivars (Hancock 1998). This suggests a means by which ploidy might increase in natural populations.

Allozyme and random amplified polymorphic DNA studies of progeny from tetraploid *Cyanococcus* cultivars found a clear pattern of tetrasomic inheritance, which is expected in autopolyploids (Krebs and Hancock 1989; Qu and Hancock 1997; Qu et al. 1998). Based on reanalyzed data from Aalders and Hall (1963), Hokanson and Hancock (1993) suggested that *Vaccinium angustifolium* shows tetrasomic inheritance and is thus an autotetraploid, contrary to the view of Vander Kloet (1977a), who considered it an allopolyploid hybrid of the diploid parents *V. boreale* and *V. pallidum*. On the basis of the results from Hokanson and Hancock (1993) and other less definitive studies, all polyploidy in *Cyanococcus* has been proposed to be derived through autopolyploidization (Lyrene et al. 2003; Soltis et al. 2007).

We emphasize, however, that variation in the definition of auto- and allopolyploidy affects the interpretation of these conclusions. In a taxonomic or mode-of-origin definition, autopolyploids arise from within a single species, whereas allopolyploids arise from interspecific hybridization. Conversely, in a genetic or cytological definition, autopolyploids display multivalent nonpreferential pairing of chromosomes during meiosis and allopolyploids display bivalent pairing (Doyle and Egan 2010; Ramsey and Schemske 2002). These two definitions will often lead to the same conclusion but sometimes a taxon may be classified as an autopolyploid under one definition and an allopolyploid under the other. In groups with highly conserved chromosome evolution, for example, hybridization between highly divergent taxa (i.e., clearly distinct species) might still result in the nonpreferential pairing of chromosomes. Conversely, divergence at individual loci could impair the ability to detect tetrasomic inheritance. Further confusing the issue is that there is no explicit criterion as to the threshold of genetic divergence above

which two entities should be considered distinct species, versus populations of a single species, not to mention the related issues involving the application of various species concepts. Depending on the group under study and research goals, both definitions can nonetheless be useful. We advocate that studies explicitly state the definition used when deciding to label a taxon as an auto- or allopolyploid. Regardless, whether autopolyploidy is the sole or primary mode of polyploidization, versus allopolyploidy as proposed by earlier authors, has not yet been assessed in natural populations of *Cyanococcus*.

Hummer et al. (2015) used U.S. Department of Agriculture germplasm stock, including many of the original species in Vander Kloet's living collection and other living material, to measure ploidy levels with flow cytometry in samples of many *Vaccinium* species. The ploidy levels in the species of *Cyanococcus* largely matched those reported for the same species in prior work.

Poster et al. (2017) achieved a critical breakthrough in understanding the prevalence and significance of ploidal variation in natural populations of *Cyanococcus*. By carefully documenting diploid and tetraploid individuals of *Vaccinium corymbosum* (apparently sensu Vander Kloet) in sympatry in New Jersey with flow cytometry, they found ploidy to be significantly correlated with both flower size and phenology, with the tetraploids having larger flowers and a peak flowering period approximately one week later than the diploids. The significant difference in flower phenology suggests an isolating mechanism between the diploids and tetraploids, although the overlap also indicates that individuals of different ploidy levels can potentially crossbreed. The strong triploid block in *Cyanococcus* would presumably result in hybrid infertility, but unreduced gametes in diploids could result in unidirectional gene flow into the tetraploid phase. Whether morphological characters beyond flower size correlate with the two ploidal segments of the population remained unaddressed in the study.

**Stomatal density and size.** Stomatal density and size on the abaxial surfaces of leaves have been correlated with ploidy in many plant groups (e.g., Beck et al. 2003; Padoan et al. 2013; Sax and Sax 1937; Tan and Dunn 1973). When patterns are detected, stomatal size is most often positively correlated with ploidy, and stomatal density is negatively correlated. In some studies, these stomatal characters have consistently predicted the level of ploidy.

Darrow et al. (1944) were the first to suggest that stomatal size and/or density could be used to estimate ploidy in *Vaccinium*. They initially considered the method to show promise for determining ploidy in the genus, but later concluded that the method was likely confounded by the complex lines of descent common in the genus. Nonetheless, as noted earlier, Aalders and Hall (1962) used stomatal length to estimate ploidy for the nomenclatural type of *V. angustifolium*. Subsequently, Chavez and Lyrene (2009) and Dweikat and Lyrene (1991) found that stomatal guard cells (and pollen tetrad diameter) were significantly longer in colchicine-derived tetraploid leaf tissue of *V. elliotii* and *V. darrowii* than in the untreated diploids from the same genetic line.

**Species relationships.** Bruederle and Vorsa (1994) conducted an allozyme study with 25 natural populations of diploid *Cyanococcus*. Based on phenetic analysis with genetic identity values and the unrooted pair group method with arithmetic mean (UPGMA) algorithm combined with an assessment of morphological distinctness, they delimited seven

diploid species: *Vaccinium boreale*, *V. corymbosum*, *V. darrowii*, *V. elliotii*, *V. myrtilloides*, *V. pallidum*, and *V. tenellum*. They anecdotally mentioned that all of their samples of *V. pallidum* from Arkansas were found to be tetraploid and thus excluded from the study. They recovered a phenogram with the pattern ((((*V. atrococcum*, *V. caesariense*), (*V. darrowii*, *V. tenellum*))), (*V. elliotii*, *V. vacillans*)), (*V. boreale*, *V. myrtilloides*)), with the samples of *V. myrtilloides* not all grouping together. Based on genetic similarity and (anecdotally presented) overlapping variation in fruit color and leaf vestiture within populations, they treated *V. atrococcum* and *V. caesariense* as conspecific with the caveat that only three populations were sampled. Both species were referred to the more inclusive but variable concept of *V. corymbosum* sensu Vander Kloet (1988) with *V. elliotii* excluded. *Vaccinium elliotii* was considered a species distinct from *V. corymbosum* by habit, leaf and branchlet morphology, flowering phenology, and ecological distribution (but again without presenting data). Based on their field observations of morphology, they anecdotally considered hybridization in *Cyanococcus* to be “widespread,” although we note that the allozyme data seem not to have uncovered instances of hybridization.

Crowl et al. (2022) conducted an analysis of the diploid representatives of *Cyanococcus* with target-enrichment genomic DNA sequence data, using flow cytometry to verify the ploidy of the samples. The analysis recovered monophyly for most morphospecies. Phylogenetic estimates were, overall, nearly identical to the phenogram of Bruederle and Vorsa (1994). *Vaccinium caesariense* was found to be nested within *V. fuscatum* (= *V. atrococcum* sensu Camp), and the two samples of *V. boreale* did not group together. The sister placement of *V. elliotii* varied between the *V. caesariense* + *V. fuscatum* clade and *V. pallidum*, depending on the analysis. Network analysis suggested a deep-level diploid hybrid speciation event between *V. elliotii* and the most recent common ancestor of the *V. boreale* + *V. myrtilloides* clade.

Rowland et al. (2022) conducted phenetic UPGMA and neighbor-joining analyses of 50 mainly cultivated accessions of *Cyanococcus* with 249 expressed sequence tag-polymerase chain reaction markers. The analysis recovered a pattern of diploid relationships similar to the two studies above, but with *V. pallidum* sister to [*V. boreale* + *V. myrtilloides*]; *V. elliotii* was sister to [*V. caesariense* + *V. fuscatum* (probably = *V. atrococcum* sensu Camp)]. They found that tetraploid *Vaccinium corymbosum* grouped most closely with the diploids *V. caesariense* and *V. fuscatum*. Tetraploid *V. angustifolium* grouped with the diploids *V. boreale* and *V. myrtilloides*, and hexaploid *V. virgatum* grouped most closely with *V. tenellum*. They considered the data to support the recognition of *V. elliotii* as a distinct species. The implications of the work for natural populations are limited by the lack of a clearly stated taxonomic framework and the use of cultivated material unspecified as to origin.

#### CURRENT STATUS, NEW DATA, AND FUTURE DIRECTIONS

We feel that progress in understanding the systematics and evolution of *Cyanococcus* over the past 80 years has been slower than might be expected, even after Luby et al. (1991) urged more comprehensive work on the group. Addressing the taxonomic problems in *Cyanococcus* from a primarily horticultural perspective has been a recurring motif. Given the tremendous economic potential of the group for blueberry breeding, we

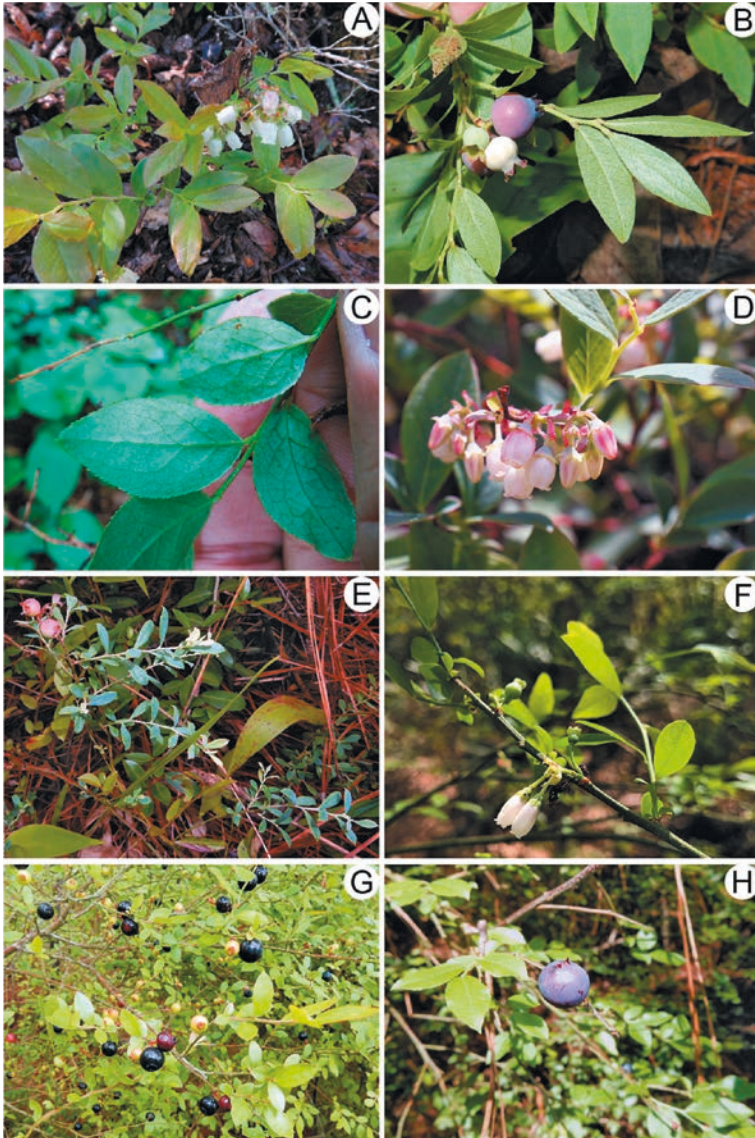


consider this understandable. However, such a perspective has often resulted in misleading extrapolation of studies based on cultivated material of unknown or imprecisely documented sources to natural variation and the limits of species. The problem is exacerbated by the inconsistent application of both scientific and common species names to cultivated samples and cultivars, the latter of which in any case may or may not correspond to species in nature. It is also compounded by the relative lack in many studies of samples vouchered with herbarium specimens, which limits the interpretation and impact of such studies. Over-reliance on crossing experiments to assess the origins of species and polyploids has also hindered the field. The morphological similarity of experimentally generated progeny to any particular species, which has been taken as *de facto* evidence of the origin of that species, may be mere coincidence and, in any case, has not been adequately documented with data in the instances proposed. Finally, assessment of ploidy has relied on far too few samples with which to base chromosome numbers, which often have been assumed too freely to define an entire species over its geographic range.

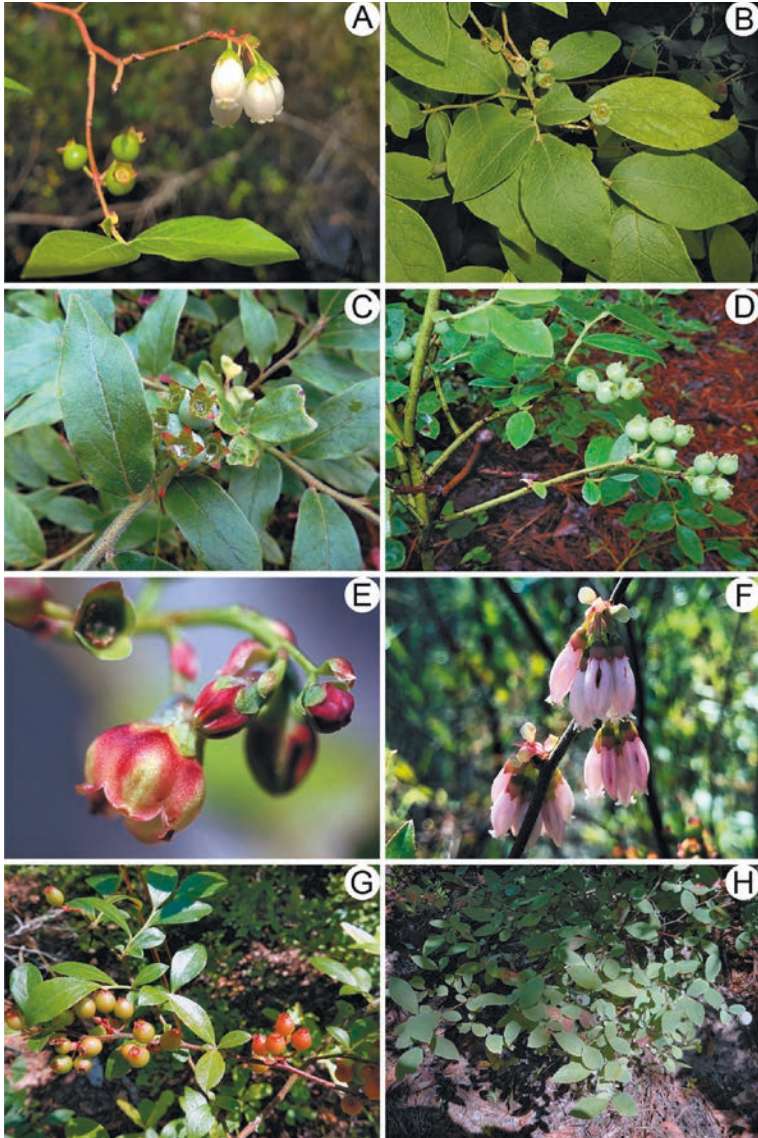
Our assessment of the current state of *Cyanococcus* systematics differs little from that of Luby et al. (1991), now over 30 years old. We feel that the treatment of Vander Kloet (2009), generally considered the current standard, to be flawed by an over-aggregation of species. Such “lumping” risks masking taxonomically significant diversity, obscures the origins of the cultivated blueberries, and leads to confusion in the field and in breeding programs. Conversely, the treatment of Camp (1945) likely has over-divided the section, as seems clear, for example, in the case of *Vaccinium pallidum* versus *V. vacillans*.

In this context, we have undertaken a multi-year collaborative project to address the longstanding issues in the systematics and evolution of *Cyanococcus* with new perspectives and data sources. We are integrating field- and herbarium-based research on samples collected from natural populations across the geographic range of *Cyanococcus* with data from DNA sequencing for phylogenetic analysis and flow cytometry to estimate ploidy. The sequence data will be used to assemble phylogenomic datasets from target enrichment and whole-genome data to infer evolutionary relationships, and a data pipeline will be designed to test hypotheses regarding the origins and mode of polyploidization and identification of diploid progenitors. Based on the sum of data, a comprehensive taxonomic revision of *Cyanococcus* will be produced. We ask the following questions to address the systematics and evolution of *Cyanococcus*: How many species are there, and what are their morphological definitions, geographic boundaries, and habitat distinctions? How is diversity generated through polyploidization? How evolutionarily labile is polyploidy—does it arise sparingly or frequently from diploidy? What is the prevalence of autopolyploidy versus allopolyploidy and their relative roles, if any, in species formation? Can morphology be used to help assess species, and is it correlated with ploidy? How prevalent is mixed ploidy within a locality (“population”)? Is hybridization rampant in the group, as initially thought, and what is its role in the group’s evolution?

Although we have only recently initiated our project, already approximately 3000 herbarium specimens of *Cyanococcus* have been examined from the combined BRIT-SMU, -VDB, and -NLU herbaria, and 550 leaf samples have been collected from across much of the range of the section for DNA sequence and flow cytometry analysis.

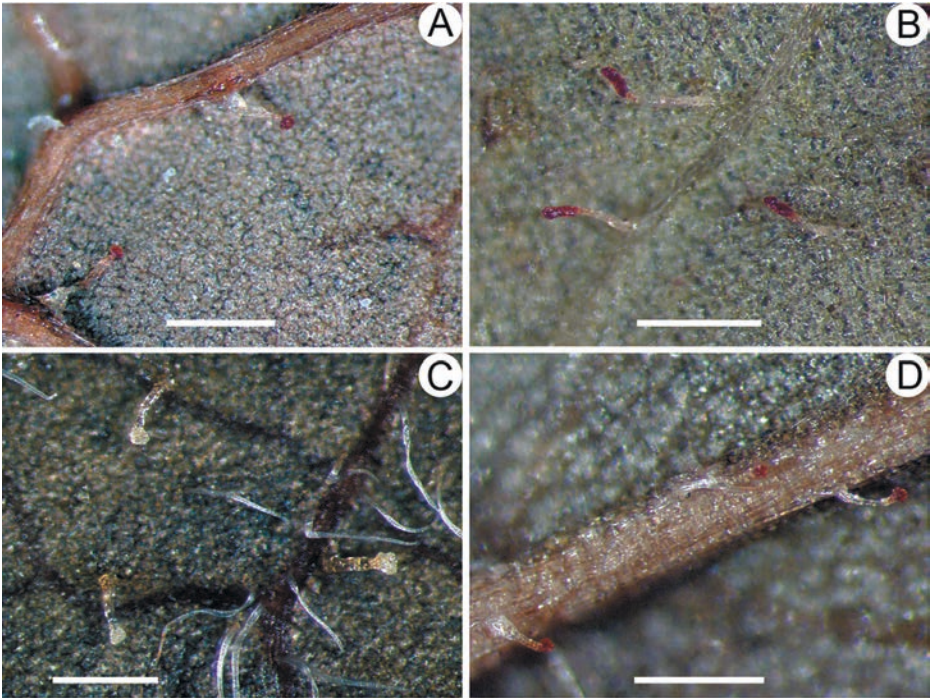


**Figure 4.** Field images of *Vaccinium* sect. *Cyanococcus*. Species names are tentative, approximating those of Camp (1945). **A.** Flowering plant of *V. angustifolium*. Cultivated, University of California Botanical Garden 78.0093 (origin: Nova Scotia, Canada). **B.** Fruiting plant of *V. angustifolium*. *Fritsch* 2371 (BRIT, DUKE). **C.** Pallid-leaved form of *V. angustifolium*. *Fritsch* 2373 (BRIT, DUKE). **D.** Inflorescence of *V. constablaei*. *Manos* CY-439 (BRIT, DUKE). **E.** Fruiting plant of *V. darrowii*. *Fritsch* 2310 (BRIT, DUKE). **F.** Flowering branchlet of *V. elliotii*. Note lack of exserted style. *Crowl* CY-368 (BRIT). **G.** *V. elliotii* with glossy black fruits. *Fritsch* 2295 (BRIT, DUKE). **H.** *V. elliotii* with glaucous fruit. *Fritsch* 2333 (BRIT, DUKE). Photos: A–C, E, G, H by P.W. Fritsch; D by P.S. Manos; F by A.A. Crowl.



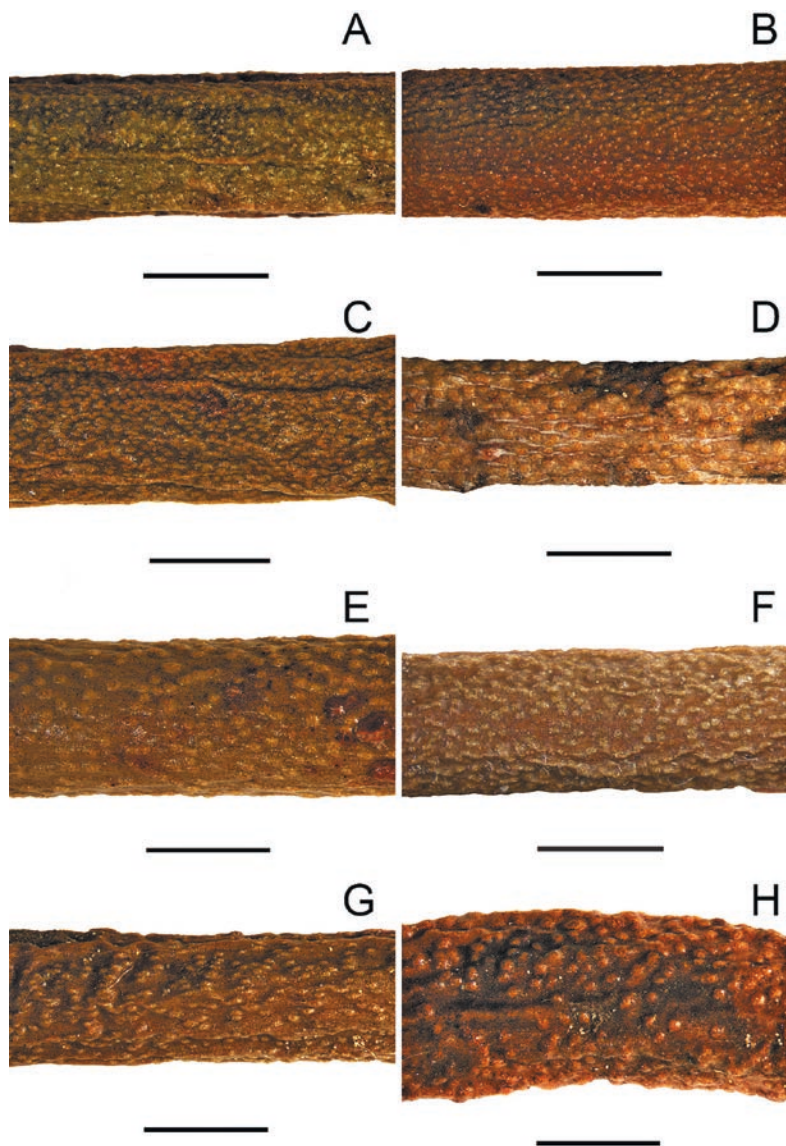
**Figure 5.** Field images of *Vaccinium* sect. *Cyanococcus*. **A.** Flowering branchlet of *V. atrocaecum*/*V. fuscatum*. Crowl CY-375 (BRIT). **B.** Fruiting branchlet of *V. atrocaecum*/*V. fuscatum*. Fritsch 2351 (BRIT, DUKE). **C.** Fruiting branchlet of *V. myrtilloides*. Fritsch 2241 (BRIT, DUKE). **D.** Fruiting branchlets of *V. pallidum* (including *V. vacillans*). Fritsch 2354 (BRIT, DUKE). **E.** Inflorescence of *V. simulatum*. Manos CY-397 (BRIT, DUKE). **F.** Inflorescences of *V. amoenum*/*V. virgatum*. Crowl. CY-374 (BRIT). **G.** Fruiting branchlet of *V. amoenum*/*V. virgatum*. Fritsch 2270 (BRIT, DUKE). **H.** Likely “rabbit-eye” escape from cultivation. Fritsch 2271 (BRIT, DUKE). Species names are tentative, approximating those of Camp (1945). Photos: A, F by A.A. Crowl; B–D, G, H, by P.W. Fritsch; E by P.S. Manos.





**Figure 6.** Stipitate-glandular trichomes on the abaxial surface of the leaf blades in *Vaccinium* sect. *Cyanococcus*. Species names are tentative, approximating those of Camp (1945). **A.** *V. atrococcum/fuscatum*, secondary veins. *Whitehouse 3159* (BRIT). **B.** *V. myrsinites*, secondary veins and surface. Note the elongated heads relative to the globose heads in the other subfigures. *Crowl 188* (BRIT). **C.** *V. amoenum/virgatum*, tertiary veins and surface. *Fritsch 2263* (BRIT, DUKE). **D.** *V. elliottii*, base of midvein. *Fritsch 2280* (BRIT, DUKE). Scale bars = 300  $\mu$ m.

Herbarium vouchers and photographs of the living plants in the field have been taken for most of the molecular and flow cytometry samples. Sampling multiple individuals in sympatry from each locality is a key distinguishing feature of our strategy. The aim is to identify morphological characters within and among localities that correlate with ploidy and phylogenetic data to discern evolutionary units (i.e., species). Although it is too early in the project to draw firm conclusions, particularly regarding our DNA sequence and flow cytometry data, we feel confident enough about the most salient of our observations on morphology in the field and herbarium specimen examination to report preliminarily on them here, accompanied by images of some of the species variation observed in the field and herbarium material (Figures 4–7). We caution that the names used below should be considered tentative, subject to change based on the critical examination of types and the results of further morphological examination of other herbarium material, ploidy assessment, and phylogenomics.



**Figure 7.** Raised stomata on second-year stems of *Vaccinium* sect. *Cyanococcus*. The size and density of the stomata are thought to be positively and negatively correlated with ploidy, respectively. Species names approximate those sensu Camp (1945) except where noted. **A.** *V. atrocoecum* (2x). *Manos* CY-070 (BRIT). **B.** *V. atrocoecum* (2x). *Manos* CY-326 (BRIT, DUKE). **C.** *V. pallidum* including *V. vacillans* (2x). *Manos* CY-052 (BRIT, DUKE). **D.** *V. australe* (4x; = *V. formosum*). *Crowl* CY-235 (BRIT). **E.** *V. arkansanum* (4x). *Manos* CY-155 (BRIT). **F.** *V. simulatum* (4x). *Manos* CY-059 (BRIT, DUKE). **G.** *V. corymbosum* (4x). *Shaw* CY-333 (BRIT, DUKE). **H.** *V. constablaei* (6x). *Manos* CY-028 (BRIT). Species names are tentative, approximating those of Camp (1945). Scale bars = 2 mm.



1. Consistent with the observations of Bruederle and Vorsa (1994), Camp (1945), Godfrey (1988), Luby et al. (1991), Uttal (1987), and Ward (1974), *Vaccinium elliotii* appears to be easily separated from the rest of Vander Kloet's "highbush blueberry" concept (Figures 4F–H and 6D). In places where *V. amoenum*/*V. virgatum* and other "highbush" blueberries occur (see below), plants with the morphology of *V. elliotii* sensu Camp appear to be distinct. One character in particular appears to easily separate *V. elliotii* from the remaining species of *Cyanococcus*, that is, a style that is included within the corolla (versus exserted; Figure 4F). It is ironic and rather inexplicable that Vander Kloet (1998) noted the existence of this character in a paper re-emphasizing the supposed indistinctness of *V. elliotii* from other "highbush" blueberries. Lyrene (1994), apparently independently of Vander Kloet, observed it in five cultivated selections from the wild identified as *V. elliotii*. Whether the short style of *V. elliotii* is a consequence of selection for or against increased selfing, or related in some way to dicliny, has not been adequately tested. Other characters in combination that can serve to distinguish *V. elliotii* are relatively small leaves that are usually not stipitate-glandular abaxially and a fruit with a persistent calyx of relatively narrow width.
2. In contrast to Vander Kloet (1983, 1988) and consistent with Ward (1974) and Uttal (1987), the abaxially stipitate-glandular-leaved plants corresponding to the Coastal Plain *Vaccinium amoenum*/*V. virgatum* sensu Camp (1945) appear to be distinct from the rest of Vander Kloet's (1983, 1988) "highbush blueberry" concept. These plants grow together with other typically non-stipitate-glandular species and are easily distinguished in the field by this character, along with a combination of usually serrate leaf blade margins and (when in flower) strongly exserted styles (Figures 5F, 5G and 6C). The correlation of their ploidy (*V. virgatum* [4x] and/or *V. amoenum* [6x]) with morphology and geography, as predicted by Camp (1945) and Ward (1974), has yet to be assessed. Also evident is that these plants can vary from less than to more than one meter tall and thus do not follow the "highbush" pattern that the treatment of Vander Kloet implies. Our preliminary findings suggest that variation in height is not categorical (which might otherwise imply a link with ploidy) but instead continuous. Another line of investigation within the stipitate-glandular-leaved group should center around the taxonomic distinctness and evolutionary history of *V. ashei*. Our preliminary field and herbarium studies, and those of B.A. Sorrie (pers. comm.) suggest that there is a naturally occurring entity in the Gulf Coastal Plain whose morphology is consistent with the type material and description of *V. ashei*. Like *V. amoenum*/*V. virgatum*, this entity appears to be distinct from the rest of the "highbush blueberry" concept.
3. A third species often growing with *Vaccinium amoenum*/*V. virgatum* and *V. elliotii* in areas where no other species of *Cyanococcus* are thought to occur (e.g., east Texas) corresponds to some version of *V. atrococcum* or *V. arkansanum* sensu Camp (1945), depending on ploidy, or *V. fuscatum* sensu Uttal (1987; Figure 5A, 5B). When these plants occur together with those of *V. amoenum*/*V. virgatum* and *V. elliotii*, we find a dry to wet microhabitat trend, with *V. amoenum*/*V. virgatum* in the driest habitats and *V. atrococcum*/*V. arkansanum* in the wettest (but at most seepy), with *V. elliotii* found in intermediate habitats.

4. The stipitate glands on the abaxial leaf surfaces of *Vaccinium tenellum* and *V. myrsinites* differ from those of all other species of *Cyanococcus* in which such glands occur by having elongate versus subglobose heads (Figure 6). This is easily observed in herbarium material at 64 $\times$  and is in fact depicted in Vander Kloet (1983; Figures 15–17), yet neither Vander Kloet nor others have apparently noted this distinction. The hypothesis in which *V. amoenum*/*V. virgatum* is derived from *V. tenellum* through autopolyploidy, at least on the assumption of homology of these glands alone, should thus be questioned, although their shared presence could still indicate close relationship. Conversely, the morphological similarity of these glands in *V. tenellum* and *V. myrsinites* supports the allopolyploid derivation of *V. myrsinites* from *V. darrowii*  $\times$  *V. tenellum* as first proposed by Camp (1945).
5. The synonymy of *Vaccinium vacillans* under *V. pallidum* (Figure 5D) as proposed by Vander Kloet (1978b, 1988) and supported by Uttal (1987) seems well justified on the basis of morphology, with the few ill-defined characters used by Camp such as margin serration appearing to be taxonomically trivial. Whether *V. altomontanum* can be considered a synonym as well, as in Vander Kloet (1988) and Uttal (1987), will depend on resolution of typology, ploidy assessment, and delimitation of its geographic range (i.e., whether it is an Appalachian endemic or, as Camp suggested, extending as far as Ohio and Arkansas).
6. The treatment of *Vaccinium lamareckii* and *V. brittonii* as synonyms of *V. angustifolium* (4x), as distinct from *V. boreale* (2x), appears to be well documented in the literature and justified on morphology (Aalders and Hall 1962, 1963; Hall and Aalders 1961; Vander Kloet 1977a, 1978a; Figure 4A–C). More study is needed to confirm the difference in corolla size between *V. angustifolium* and *V. boreale*.
7. Vander Kloet (1980) observed variation in the presence and quantity of fruit “bloom” (glaucescence) for the F<sub>1</sub> progeny of two wild-collected individuals of “highbush” blueberries corresponding most closely to *Vaccinium atrococcum* of Camp, and also for the F<sub>1</sub> progeny of wild-collected *V. angustifolium* (Vander Kloet 1978a). We have observed variation in this character within each of *V. atrococcum*, *V. elliottii*, and *V. amoenum*/*V. virgatum* in Arkansas and Louisiana (e.g., cf. Figure 4G, 4H). This character will require careful evaluation as to its taxonomic utility. We suspect that it has been overemphasized in taxonomic treatments because of the focus on fruit qualities in blueberry breeding programs. This is likely true also for fruit flavor and degree of sweetness.
8. Assertions of hybridization in *Cyanococcus* have invariably suffered from inadequate documentation in the literature. Claims are based either on anecdotal morphological observations (i.e., those without precise backing data) or vaguely defined similarity of natural variants or species to experimentally crossed hybrids. More rigorous documentation and analyses based on morphological and genetic data are required before the prevalence of natural hybridization in *Cyanococcus* can be assessed appropriately. Nonetheless, we agree with others (Uttal 1987; Vander Kloet 1977b; Ward 1974; Weakley 2020) that the prevalence of hybridization and introgression in natural populations of *Cyanococcus*, at least as asserted by Camp (1945), appears to have been

overstated. In our field work, we have thus far only infrequently encountered plants that exhibit some form of intermediacy between other species that occur in the same area. Moreover, in at least some of these cases, we speculate that such intermediates represent simple infraspecific variation of morphological characters at one ploidy level having nothing to do with species reticulation.

One example worthy of further study is the presence or absence of subglobose-headed stipitate glands on the abaxial surface of leaves. Such glands can occur in species (sensu Camp) other than those considered to be defined by the presence of this character (*Vaccinium amoenum*/*V. virgatum*, *V. myrsinites*, and *V. tenellum*). They are clearly present occasionally in specimens of *V. atrococcum* and *V. elliottii* (Figure 6A, 6D). In these cases, the stipes are usually longer than those seen in *V. amoenum*/*V. virgatum*; in *V. atrococcum* they occur most frequently on the leaves of sterile branchlets. Camp (1945) considered the presence of such glands in plants that are otherwise morphologically consistent with *V. elliottii* evidence of hybridization of *V. elliottii* and *V. tenellum*. However, we have only observed subglobose-headed glands in *V. elliottii*, not elongate-headed glands, which might be expected in at least some of such hybrids. Rather than the result of hybridization with a glandular species, the presence or absence of such glands likely represents mere variation within these species. The same variation in the presence or absence of stipitate glands on the leaves abaxially is evident in species of *Vaccinium* in other sections, for example, *V. arboreum* Marshall in *V. sect. Batodendron* (Nutt.) A. Gray, and *V. stamineum* of *V. sect. Polycodium* Rehder; it would thus not be surprising if this type of variation occurs in species of *Cyanococcus* as well.

Another confounding factor potentially leading to an overestimation in the prevalence of natural hybridization in *Cyanococcus* may be the escape of cultivars from plantings into natural environments. Such escapes have already been anecdotally noted by Camp (1945) and Ballington et al. (1982) and have been postulated in northern Florida (P.M. Lyrene, pers. comm.). In a forest in Smith County, Texas, an area considered out of range for native *Cyanococcus* species, we observed a single 2.4 m tall shrub with pale leaves and calyces and sparsely scattered stipitate glands on the leaves abaxially (*Fritsch* 2259 [BRIT, DUKE]). In a clearing on the same property, we observed cultivated blueberry plants (*Fritsch* 2260 [BRIT]) with the same (vegetative only) characters as the plant in the forest, suggesting that the forest plant is progeny from the cultivated plants. In the Watson Native Plant Preserve in Tyler County, Texas, we observed a single plant with the same morphology as the plant above with larger fruits than those from the typical plants of *Vaccinium amoenum*/*V. virgatum* growing in the same preserve (6 × 10 mm versus 4–5 × 5–6 mm in *sicco*) and of clearly different appearance (e.g., glaucous versus green leaves and calyces; *Fritsch* 2271 [BRIT, DUKE] and *Fritsch* 2270 [BRIT, DUKE], respectively; Figure 5H). Blueberry farms are scattered throughout Tyler County. Thus it seems clear that escapes from cultivation do occur at least sporadically, but the extent to which such plants have been conflated with naturally occurring hybridization has not been previously considered. We hypothesize that escape from cultivation into natural habitats of “rabbit-eye,” “northern highbush,” and other cultivars, which

today are often complex hybrids developed through crop breeding programs (Song and Hancock 2011), at least in part explain Camp's observations in wild *Cyanococcus* (e.g., those concerning *V. ashei* and *V. corymbosum* sensu Camp). Assessing the occurrence and prevalence of the intermixing of such escapes with natural populations will require the application of DNA sequence and flow cytometry data.

9. Despite the promise shown by studies for the use of stomatal density and size as an indicator of ploidy in *Cyanococcus* (Aalders and Hall 1962; Chavez and Lyrene 2009; Darrow et al. 1944; Dweikat and Lyrene 1991), apparently none since Aalders and Hall (1962) have assessed these characters as a proxy for estimating ploidy in natural populations of the group. In this regard, it may be fortuitous that in *Cyanococcus* the verrucose stems that define the section in *Vaccinium* result from raised stomata, where each stomate occurs atop a wart-like projection. Stem stomatal size and density can be roughly estimated with a stereomicroscope at 64 $\times$ . We have used second-year stem morphology with other characters (e.g., stem color, corolla color and shape, and plant height) and flow cytometry to consistently distinguish individuals of *V. pallidum* (2x), *V. simulatum* (4x), and *V. constablaei* (6x; all sensu Camp 1945), all growing together in the Southern Appalachian Mountains, from each other (Figure 7). The relationship also appears to exist among *V. atrococcum* (2x), *V. corymbosum* (4x), and *V. formosum* (4x), all sensu Camp (Figure 7). This correlation, if substantiated on more rigorous investigation and particularly if the variation is found to be discontinuous and tied to ploidy, would greatly aid in the identification of *Cyanococcus* species in the Appalachians and elsewhere. Studies on this topic would be most impactful if pollen tetrad diameter were also included.
10. *Vaccinium simulatum*, endemic to the Southern and Central Appalachians, appears to be separable by the combination of height more than 1 m, broadly urceolate to campanulate corollas, and the stomate size and density associated with tetraploid plants (Figure 5E). Luby et al. (1991) also suggested distinct species status for *V. simulatum* but proposed more highly variable leaf morphology than indicated in Camp (1945). Separation of *V. simulatum* and *V. corymbosum* must be more carefully assessed (see below).
11. Once *Vaccinium amoenum*/*V. ashei*/*V. virgatum*, *V. elliottii*, and *V. simulatum* and are excluded from the concept of the "highbush" blueberry, the remaining variation in the group encompasses Camp's *V. caesariense* and *V. atrococcum* (or *V. fuscatum*, depending on currently unresolved nomenclature) at the diploid level, and *V. arkansanum*, *V. corymbosum*, *V. formosum* (= *V. australe*), and *V. marianum* at the tetraploid level. We present the following working hypothesis to explain the morphological variation among the entities in this "residual highbush" group for testing with DNA sequence and flow cytometry data. (A similar hypothesis was proposed by Luby et al. [1991] in more abbreviated form.) We propose that diploid *V. atrococcum* and *V. caesariense* are two ends of a continuous spectrum of leaf pubescence (densely pubescent versus glabrous, respectively) where intermediates merely reflect morphological variation within a single diploid species. Similarly, tetraploid *V. arkansanum* and *V. formosum* comprise a single species with the same patterns of morphological variation (the



named intermediate tetraploid *V. marianum* sensu Camp would be a synonym). This hypothesis would be consistent with the variation in progeny observed anecdotally by Vander Kloet (1980) and in the field by Ballington et al. (1980, 1982), the overall genetic similarity in diploid *V. atrococcum* and *V. caesariense* observed in the allozyme study of Bruederle and Vorsa (1994), and the study of Poster et al. (2017) documenting diploids and tetraploids in a natural population of *V. corymbosum* sensu Vander Kloet. The tetraploids could have originated from a diploid progenitor once or multiple times. The latter would render the tetraploids polyphyletic, and even then, they might cross among themselves locally or panmictically.

The remaining element of the “residual highbush” group to be considered is *Vaccinium corymbosum*, the name Camp applied to the widespread “highbush” blueberry in wetlands throughout the northeastern and north-central United States, and southeastern Canada. In origin and evolution, this entity may be the most enigmatic of those in the entire *Cyanococcus* clade. It could stand by itself as a tetraploid of these northern forests (whether or not, as Camp asserted, its southern boundary corresponds to that of the maximal extent of the last ice sheet). Alternatively, as proposed by Camp (1945), it could intergrade with various other species such as *V. simulatum*, if both are found to be consistently tetraploid, and possibly even with tetra-ploid *V. arkansanum* sensu Camp, in which case it would extend across the entire eastern United States. It may even have been derived polyphyletically from diploid ancestors, then resembling Vander Kloet’s compilospecies concept but with a narrower set of ancestors and/or descendants.

We hope and anticipate that this and the other hypotheses put forward in the literature and here will be addressed in the coming years through estimates of ploidy from flow cytometry, phylogenetic relationships from high-throughput DNA sequence data, and intensive study of morphology, all densely sampled from throughout the geographic range of *Cyanococcus*. With these data in hand, we are optimistic that the complex evolution of this ecologically and economically important group of plants can finally be disentangled, with benefits for biodiversity studies, conservation, and crop breeding.

#### ACKNOWLEDGMENTS

We thank the U.S. National Science Foundation for financial support of this work (DEB-2038213 to Duke and NCSU, and DEB-2038217 to BRIT), and two reviewers for helpful comments on the manuscript.

#### LITERATURE CITED

- AALDERS, L.E. AND I.V. HALL. 1962. New evidence on the cytotaxonomy of *Vaccinium* species as revealed by stomatal measurements from herbarium specimens. *Nature* 196:694. DOI:10.1038/196694a0
- AND ———. 1963. The inheritance and taxonomic significance of the “nigrum” factor in the common lowbush blueberry, *Vaccinium angustifolium*. *Canadian Journal of Genetics & Cytology* 5:115–118. DOI:10.1139/g63-019
- BALLINGTON, J.R. AND G.J. GALLETTA. 1978. Comparative crossability of four diploid *Vaccinium* species. *Journal of the American Society for Horticultural Science* 103:554–560. DOI:10.21273/JASHS.103.4.554

- , W.B. KIRKMAN, D.V. BARKLEY, AND A.F. HUYLER. 1980. *Vaccinium* Germplasm Collections, North Carolina and South Carolina 1978 and 1979. Horticultural Crops Research Series No. 51, North Carolina State University, Raleigh, NC.
- , W.H. GENSEL, Y.M. ISENBERG, AND C.A. WALKER, JR. 1982. *Vaccinium* Germplasm Collections, 1980–1982. Horticultural Crops Research Series No. 60, North Carolina State University, Raleigh, NC.
- BECK, S.L., R.W. DUNLOP, AND A. FOSSEY. 2003. Stomatal length and frequency as a measure of ploidy level in black wattle, *Acacia mearnsii* (de Wild). Botanical Journal of the Linnean Society 141:177–181. DOI:10.1046/j.1095-8339.2003.00132.x
- BRITTON, N.L. AND A. BROWN. 1913. Illustrated Flora of the Northern United States, Canada and the British Possessions, 2nd ed., vol. 3. Charles Scribner's Sons, New York, NY.
- BRUEDERLE, L.P. AND N. VORSA. 1994. Genetic differentiation of diploid blueberry, *Vaccinium* sect. *Cyanococcus* (Ericaceae). Systematic Botany 19:337–349. DOI:10.2307/2419760
- CAMP, W.H. 1942. On the structure of populations in the genus *Vaccinium*. Brittonia 4:189–204. DOI:10.2307/2804713
- . 1945. The North American blueberries with notes on other groups of Vacciniaceae. Brittonia 5:203–275. DOI:10.2307/2804880
- AND C.L. GILLY. 1943. The structure and origin of species. Brittonia 4:323–385. DOI:10.2307/2804896
- CHAPMAN, A.W. 1897. Flora of the Southern United States, 3rd ed. Cambridge Botanical Supply Company, Cambridge, MA.
- CHAVEZ, D.J. AND P.M. LYRENE. 2009. Production and identification of colchicine-derived tetraploid *Vaccinium darrowii* and its use in breeding. Journal of the American Society for Horticultural Science 134:356–363. DOI:10.21273/JASHS.134.3.356
- CORRELL, D.S. AND M.C. JOHNSTON. 1970. Manual of the Vascular Plants of Texas. University of Texas at Dallas, Dallas, TX.
- CROWL, A.A., P.W. FRITSCH, G.P. TILEY, N.P. LYNCH, T.G. RANNEY, H. ASHRAFI, AND P.S. MANOS. 2022. A first complete phylogenomic hypothesis for diploid blueberries (*Vaccinium* section *Cyanococcus*). American Journal of Botany 109:1596–1606. DOI:10.1002/ajb2.16065
- DARROW, G.M. AND W.H. CAMP. 1945. *Vaccinium* hybrids and the development of new horticultural material. Bulletin of the Torrey Botanical Club 72:1–21. DOI:10.2307/2481260
- , H.E. FISCHER, AND H. DERMEN. 1944. Chromosome numbers in *Vaccinium* and related groups. Bulletin of the Torrey Botanical Club 71:498–506. DOI:10.2307/2481242
- DOYLE, J.J. AND A.N. EGAN. 2010. Dating the origins of polyploid events. New Phytologist 186:73–85. DOI:10.1111/j.1469-8137.2009.03118.x
- DWEIKAT, I.M. AND P.M. LYRENE. 1991. Induced tetraploidy in a *Vaccinium elliottii* facilitates crossing with cultivated highbush blueberry. Journal of the American Society for Horticultural Science 116:1063–1066. DOI:10.21273/JASHS.116.6.1063
- GALLETTA, G.J. AND J.R. BALLINGTON. 1996. Blueberries, cranberries and lingonberries, pp. 1–108. In: Janick, J. and J.N. Moore, eds., Fruit Breeding, vol. 2, Vine and Small Fruits. John Wiley & Sons, New York, NY.
- GODFREY, R.K. 1988. Trees, Shrubs, and Woody Vines of Northern Florida and Adjacent Georgia and Alabama. University of Georgia Press, Athens, GA.
- GRANT, V. 1981. Plant Speciation, 2nd ed. Columbia University Press, New York, NY.
- GRAY, A. 1848. Chloris boreali-Americana. Memoirs of the American Academy of Arts & Sciences, n.s., 3:1–56. DOI:10.2307/25058141
- . 1859. Diagnostic characters of new species of phaenogamous plants collected in Japan by Charles Wright. Memoirs of the American Academy of Arts & Sciences, n.s., 6:377–452. DOI:10.2307/25057953
- . 1860. Remarks on the botany of Japan, in its relations to that of North America and of other parts of the northern temperate zone. Edinburgh New Philosophical Journal, n.s., 11:159–163.

- HALL, I.V. AND L.E. AALDERS. 1961. Cytotaxonomy of low-bush blueberries in eastern Canada. *American Journal of Botany* 48:199–201. DOI:10.1002/j.1537-2197.1961.tb11625.x
- HANCOCK, J.F. 1998. Using southern blueberry species in northern highbush breeding, pp. 91–94. *In*: Cline, W.O. and J.R. Ballington, eds., *Proceedings of the 8th North American Blueberry Research and Extension Workers Conference*, Wilmington, North Carolina. Published by North Carolina State University, Raleigh, NC.
- HARLAN, J.R. AND J.M.J. DEWET. 1963. The compilospecies concept. *Evolution* 17:497–501. DOI:10.1111/j.1558-5646.1963.tb03307.x
- HOKANSON, K. AND J. HANCOCK. 1993. The common lowbush blueberry, *Vaccinium angustifolium* Aiton, may be an autopolyploid. *Canadian Journal of Plant Science* 73:889–891. DOI:10.4141/cjps93-116
- HUMMER, K.E., N.V. BASSIL, H.P. RODRÍGUEZ ARMENTA, AND J.W. OLMSTEAD. 2015. *Vaccinium* species ploidy assessment. *Acta Horticulturae* 1101:199–204. DOI:10.17660/ActaHortic.2015.1101.30
- INTERNATIONAL PLANT NAMES INDEX. 2021. The Royal Botanic Gardens, Kew, Harvard University Herbaria and Libraries and Australian National Herbarium. Website (<http://www.ipni.org>). Accessed Dec 2021.
- KLEINMAN, K. 2009. Biosystematics and the origin of species. Edgar Anderson, W. H. Camp, and the evolutionary synthesis, pp. 73–91. *In*: Cain, J. and M. Ruse, eds., *Descended from Darwin: Insights into the History of Evolutionary Studies 1900–1970*. American Philosophical Society, Philadelphia, PA.
- KLOTZSCH, J.F. 1851. Studien über die natürliche klasse Bicornes Linné. *Linnaea* 24:1–88.
- KREBS, S.L. AND J.F. HANCOCK. 1989. Tetrasomic inheritance of isoenzyme markers in the high-bush blueberry, *Vaccinium corymbosum* L. *Heredity* 63:11–18. DOI:10.1038/hdy.1989.70
- LUBY, J.J., J.R. BALLINGTON, A.D. DRAPER, K. PLISKA, AND M.E. AUSTIN. 1991. Blueberries and cranberries (*Vaccinium*), pp. 391–456. *In*: Moore, J.N. and J.R. Ballington, eds., *Genetic Resources of Temperate Fruit and Nut Crops*. International Society for Horticultural Science, Wageningen, Netherlands.
- LUTEYN, J.L., W.S. JUDD, S.P. VANDER KLOET, L.J. DORR, G.D. WALLACE, K.A. KRON, P.F. STEVENS, AND S.E. CLEMANTS. 1996. Ericaceae of the southeastern United States. *Castanea* 61:101–144.
- LYRENE, P.M. 1994. Variation within and among blueberry taxa in flower size and shape. *Journal of the American Society for Horticultural Science* 119:1039–1042. DOI:10.21273/JASHS.119.5.1039
- , N. VORSA, AND J.R. BALLINGTON. 2003. Polyploidy and sexual polyploidization in the genus *Vaccinium*. *Euphytica* 133:27–36. DOI:10.1023/A:1025608408727
- MASON, A.S. AND J.F. WENDEL. 2020. Homoeologous exchanges, segmental allopolyploidy, and polyploid genome evolution. *Frontiers in Genetics* 11:1014. DOI:doi.org/10.3389/fgene.2020.01014
- MAYR, E. 1942. *Systematics and the Origin of Species*. Columbia University Press, New York, NY.
- . 1963. *Animal Species and Evolution*. Harvard University Press, Cambridge, MA.
- ODELL, A.E., S.P. VANDER KLOET, AND R.E. NEWELL. 1989. Stem anatomy of *Vaccinium* section *Cyanococcus* and related taxa. *Canadian Journal of Botany* 67:2328–2334. DOI:10.1139/b89-298
- ORTIZ, R., L.P. BRUEDERLE, T. LAVERTY, AND N. VORSA. 1992a. The origin of polyploids via  $2n$  gametes in *Vaccinium* section *Cyanococcus*. *Euphytica* 61:241–246. DOI:10.1007/BF00039664
- , N. VORSA, L.P. BRUEDERLE, AND T. LAVERTY. 1992b. Occurrence of unreduced pollen in diploid blueberry species, *Vaccinium* sect. *Cyanococcus*. *Theoretical & Applied Genetics* 85:55–60. DOI:10.1007/BF00223844
- , ———, ———, AND ———. 1999. Pollen viability in natural populations of three North American diploid species of blueberry (*Vaccinium* section *Cyanococcus*). *Scientia Horticulturae* 80:39–48. DOI:10.1016/S0304-4238(98)00220-9
- PADOAN, D., A. MOSSAD, B. CHIANCONE, M.A. GERMANA, AND P.S.S.V. KHAN. 2013. Ploidy levels in *Citrus clementine* affects leaf morphology, stomatal density and water content. *Theoretical & Experimental Plant Physiology* 25:283–290.

- POSTER, L.S., S.N. HANDEL, AND P.E. SMOUSE. 2017. Corolla size and temporal displacement of flowering times among sympatric diploid and tetraploid highbush blueberry (*Vaccinium corymbosum*). *Botany* 95:395–404. DOI:10.1139/cjb-2016-013
- POWELL, E.A. AND K.A. KRON. 2002. Hawaiian blueberries and their relatives—a phylogenetic analysis of *Vaccinium* sections *Macropelma*, *Myrtillus*, and *Hemimyrtillus* (Ericaceae). *Systematic Botany* 27:768–779. DOI:10.1043/0363-6445-27.4.768
- QU, L. AND J.F. HANCOCK. 1997. RAPD-based genetic linkage map of blueberry derived from an interspecific cross between diploid *Vaccinium darrowi* [sic] and tetraploid *V. corymbosum*. *Journal of the American Society for Horticultural Science* 122:69–73.
- , ———, AND J.H. WHALLON. 1998. Evolution in an autopolyploid group displaying predominantly bivalent pairing at meiosis: genomic similarity of diploid *Vaccinium darrowi* [sic] and autotetraploid *V. corymbosum* (Ericaceae). *American Journal of Botany* 85:698–703. DOI:10.2307/2446540
- RAMSEY, J. AND D.W. SCHEMSKE. 2002. Neopolyploidy in flowering plants. *Annual Review of Ecology & Systematics* 33:589–639. DOI:10.1146/annurev.ecolsys.33.010802.150437
- REHDER, A. 1940. *Manual of Cultivated Trees and Shrubs*, 2nd ed. Macmillan Co., New York, NY.
- ROBINSON, B.L. AND M.L. FERNALD. 1908. *Gray's New Manual of Botany*, 7th ed. American Book Company, New York, NY.
- ROWLAND, L.J., E.L. OGDEN, AND J.R. BALLINGTON. 2022. Relationships among blueberry species within the section *Cyanococcus* of the *Vaccinium* genus based on EST-PCR markers. *Canadian Journal of Plant Science* 102:744–748. DOI:10.1139/cjps-2021-0221
- RYDBERG, P.A. 1917. *Flora of the Rocky Mountains and Adjacent Plains*. Steinman & Foltz, Lancaster, PA.
- SAX, K. AND H.J. SAX. 1937. Stomata size and distribution in diploid and polyploid plants. *Journal of the Arnold Arboretum* 18:164–172.
- SLEUMER, H. 1941. Vaccinioideen studien. *Botanische Jahrbücher für Systematik, Pflanzengeschichte und Pflanzengeographie* 71:375–510.
- SMALL, J.K. 1933. *Manual of the Southeastern Flora*. University of North Carolina Press, Chapel Hill, NC.
- SNEATH, P.H. AND R.R. SOKAL. 1973. *Numerical Taxonomy*. W.H. Freeman & Co., San Francisco, CA.
- SOLTIS, D.E., P.S. SOLTIS, D.W. SCHEMSKE, J.F. HANCOCK, J.N. THOMPSON, B.C. HUSBAND, AND W.S. JUDD. 2007. Autopolyploidy in angiosperms: have we grossly underestimated the number of species? *Taxon* 56:13–30. DOI:10.2307/25065732
- SONG, G.-Q. AND J.F. HANCOCK. 2011. *Vaccinium*, pp. 197–221. In: Kole, C. ed., *Wild Crop Relatives: Genomic and Breeding Resources, Temperate Fruits*. Springer-Verlag, Berlin, Germany.
- SPOELHOF, J.P., P.S. SOLTIS, AND D.E. SOLTIS. 2017. Pure polyploidy: closing the gaps in autopolyploid research. *Journal of Systematics & Evolution* 55:340–352. DOI:10.1111/jse.12253
- STEBBINS, G.L. 1950. *Variation and Evolution in Plants*. Columbia University Press, New York, NY.
- STEVENS, P.F. 2004. Ericaceae, pp. 145–194. In: Kubitzki, K., ed., *The Families and Genera of Vascular Plants*, vol. 6. Springer-Verlag, Berlin, Germany.
- TAN, G.-Y. AND G.M. DUNN. 1973. Relationship of stomatal length and frequency and pollen-grain diameter to ploidy level in *Bromus inermis* Leyss. *Crop Science* 13:332–334. DOI:10.2135/cropsci1973.0011183X001300030014x
- UTTAL, L.J. 1986. An older name for *Vaccinium australe* Small. *Castanea* 51:221–224.
- . 1987. The genus *Vaccinium* L. (Ericaceae) in Virginia. *Castanea* 52:231–255.
- VANDER KLOET, S.P. 1976a. Nomenclature, taxonomy, and biosystematics of *Vaccinium* section *Cyanococcus* (the blueberries) in North America. I. Natural barriers to gene exchange between *Vaccinium angustifolium* Ait. and *Vaccinium corymbosum* L. *Rhodora* 78:503–515.
- . 1976b. A novel approach to sampling *Vaccinium* populations. *Canadian Journal of Botany* 54:669–671. DOI:10.1139/b76-070



- . 1977a. The taxonomic status of *Vaccinium boreale*. Canadian Journal of Botany 55:281–288. DOI:10.1139/b77-038
- . 1977b. Potential and actual gene exchange among three sympatric species of *Vaccinium* § *Cyanococcus* in Highlands County, Florida. Canadian Journal of Botany 55:2668–2672. DOI:10.1139/b77-304
- . 1978a. Systematics, distribution, and nomenclature of the polymorphic *Vaccinium angustifolium*. Rhodora 80:358–376.
- . 1978b. The taxonomic status of *Vaccinium pallidum*, the hillside blueberries including *Vaccinium vacillans*. Canadian Journal of Botany 56:1559–1574. DOI:10.1139/b78-185
- . 1980. The taxonomy of the highbush blueberry, *Vaccinium corymbosum*. Canadian Journal of Botany 58:1187–1201. DOI:10.1139/b80-148
- . 1983. The taxonomy of *Vaccinium* § *Cyanococcus*: a summation. Canadian Journal of Botany 61:256–266. DOI:10.1139/b83-029
- . 1988. The Genus *Vaccinium* in North America. Research Branch Agriculture Canada Publ. 1828, Ottawa, Ontario, Canada.
- . 1998. The taxonomic status of *Vaccinium elliottii*, once more with feeling!, pp. 17–23. In: Cline W.O. and J.R. Ballington, eds., Proceedings of the 8th North American Blueberry Research and Extension Workers Conference, Wilmington, North Carolina. Published by North Carolina State University, Raleigh, NC.
- . 2009. *Vaccinium*, pp. 515–530. In: Flora of North America Editorial Committee, eds., Flora of North America North of Mexico, vol. 8. Magnoliophyta: Paeoniaceae to Ericaceae. Oxford University Press, New York, NY.
- AND T.A. DICKINSON. 2009. A subgeneric classification of the genus *Vaccinium* and the metamorphosis of *V.* section *Bracteata* Nakai: more terrestrial and less epiphytic in habit, more continental and less insular in distribution. Journal of Plant Research 122:253–268. DOI:10.1007/s10265-008-0211-7
- WARD, D.B. 1974. Contributions to the flora of Florida: 6. *Vaccinium*. Castanea 39:191–205.
- AND P.M. LYRENE. 2007. *Vaccinium tenellum* is not a native Florida blueberry. Castanea 72:45–46. DOI:10.2179/0008-7475(2007)72[45:SN]2.0.CO;2
- WEAKLEY, A.S. 2020. Flora of the southeastern United States. University of North Carolina Herbarium, North Carolina Botanical Garden, Chapel Hill, NC.