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BRIEF COMMUNICATION

Inferring the evolutionary reduction of corm lobation in *Isoëtes* using Bayesian model-averaged ancestral state reconstruction

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PREMISE OF THE STUDY: Inferring the evolution of characters in *Isoëtes* has been problematic, as these plants are morphologically conservative and yet highly variable and homoplasious within that conserved base morphology. However, molecular phylogenies have given us a valuable tool for testing hypotheses of character evolution within the genus, such as the hypothesis of ongoing morphological reductions.

METHODS: We examined the reduction in lobe number on the underground trunk, or corm, by combining the most recent molecular phylogeny with morphological descriptions gathered from the literature and observations of living specimens. Ancestral character states were inferred using nonstationary evolutionary models, reversible-jump MCMC, and Bayesian model averaging.

KEY RESULTS: Our results support the hypothesis of a directional reduction in lobe number in *lsoëtes*, with the best-supported model of character evolution being one of irreversible reduction. Furthermore, the most probable ancestral corm lobe number of extant *lsoëtes* is three, and a reduction to two lobes has occurred at least six times.

CONCLUSIONS: From our results, we can infer that corm lobation, like many other traits in *lsoëtes*, shows a degree of homoplasy, and yet also shows ongoing evolutionary reduction.

KEY WORDS Bayes factors; evolutionary reduction; Isoetaceae; model selection; morphological evolution; morphological simplification; RevBayes.

Interpreting the morphological evolution of *Isoëtes* L. has troubled botanists for many years. These plants have an outwardly simple, highly conserved body plan, consisting of an apical rosette of linear sporophylls on top of a reduced corm-like trunk (Engelmann, 1882; Pfeiffer, 1922; Cox and Hickey, 1984; Hickey, 1986; Taylor and Hickey, 1992; Budke et al., 2005), yet are highly variable both within (Budke et al., 2005; Liu et al., 2006) and among closely related species (Cox and Hickey, 1984; Hickey, 1986; Taylor and Hickey, 1992; Romero and Real, 2005; Bagella et al., 2011). Even characters once thought to be useful in delimiting natural groups within the genus, such as habitat (Engelmann, 1882) or megaspore morphology (Pfeiffer, 1922), have been found to be labile (Cox and Hickey, 1984; Taylor and Hickey, 1992; Budke et al., 2005; Hickey, 2007; Bagella

et al., 2011). And while some characters, such as the glossopodium (the portion of the ligule internal to the leaf), have shown some diagnostic potential (Sharma and Singh, 1984; Pant et al., 2000; Shaw and Hickey, 2005; Singh et al., 2010; Freund, 2016), actually examining and interpreting these structures requires considerable histological and computational effort, making them ill-suited for field identification. This absence of consistent, dependable characters creates a paradox: the lack of reliable traits impedes the inference of phylogenies or classifications in the genus, but without a phylogeny, examining character evolution is exceptionally difficult.

However, molecular phylogenetics has vastly improved our understanding of *Isoëtes* (Rydin and Wikstrom, 2002; Hoot et al., 2004, 2006; Schuettpelz and Hoot, 2006; Larsén and Rydin, 2016) and has provided evidence for five major clades within the genus: Gondwanan (Clade A), Laurasian (Clade B), Italian (Clade C), Austro-Asian (Clade D), and New World or American (Clade E; Hoot et al., 2006; Larsén and Rydin, 2016). This phylogeny has completely overturned the old morphological and ecological systems of classification and has also provided a framework to begin earnestly studying character evolution in the genus. While characters such as spore ornamentation are highly homoplastic (Cox and Hickey, 1984; Larsén and Rydin, 2016), there are other changes that may be informative, such as serial reduction of the corm (Karrfalt and Eggert, 1977a, b; Pigg, 1992; Grauvogel-Stamm and Lugardon, 2001).

The isoetalians have a long fossil history, spanning >300 million yr (Pigg, 1992; Gensel and Pigg, 2010; Taylor et al., 2009). Due to this long history, and the abundance of fossil representatives, there is a long-standing comparison between the corm of extant Isoëtes and the stigmarian appendages of their extinct arborescent relatives. This comparison is supported by the unique suite of characters that are found only in the isoetalian lycophytes (Stewart, 1947; Karrfalt and Eggert, 1977a, b; Jennings et al., 1983; Pigg, 1992). These features include a unique and independently evolved form of secondary growth that is fundamentally different from the secondary growth of spermatophytes (Scott and Hill, 1900; Stokey, 1909; Pfeiffer, 1922; Stewart, 1947; Karrfalt and Eggert, 1977a; Pigg and Rothwell, 1985; Ash and Pigg, 1991; Pigg, 1992; Gensel and Pigg, 2010). Their rooting structures are also unique and have been compared to a dichotomous branching system, with their rootlets arising from modified leaves (Rothwell and Erwin, 1985). Some fossil isoetalians, such as the arborescent Protostigmaria Jennings (Jennings et al., 1983) or geophytic Nathorstiana Richter (Richter, 1910; Mägdefrau, 1932; Taylor et al., 2009), have subterranean morphology comparable to that of modern Isoëtes (i.e., not producing elongated stigmarian appendages), with the size, elaboration, or arborescent habit of the plants separating them from the contemporary taxa (Stewart, 1947; Karrfalt, 1984a; Pigg, 1992). Protostigmaria, while having a generally comparable morphology to extant Isoëtes, differs by having as many as 13 lobes on its rooting axis. Nathorstiana, while having corms that are quite similar to those of extant Isoëtes, shows high levels of corm lobe variability, like some extant species of Isoëtes, yet little correlation of plant size with lobe number (some of the small plants have four lobes, and some larger plants are unlobed; Richter, 1910; Mägdefrau, 1932; Jennings et al., 1983; Rothwell, 1984; Rothwell and Erwin, 1985; Taylor et al., 2009). Extant Isoëtes, and potentially the extinct forms, also possess a unique mechanism to penetrate into the substrate by laterally displacing the soil and then pulling the corm deeper through a combination of secondary growth and development of new rootlets. While functionally similar to the contractile roots of monocots, this mechanism does not compress the soil below the plant, allowing for deeper substrate penetration (Karrfalt, 1977). Due to these shared features, it has been hypothesized that modern Isoëtes are the result of continuous reduction of the once arborescent isoetalian body plan, with the reduced corm lobe numbers in modern Isoëtes continuing this trend (Stewart, 1947; Karrfalt and Eggert, 1977a, b; Jennings et al., 1983; Pigg, 1992).

These corm lobes are formed by the basal, rootlet-producing meristems (Engelmann, 1882; Stokey, 1909; Osborn, 1922; Pfeiffer, 1922; Bhambie, 1963; Karrfalt and Eggert, 1977a; Budke et al., 2005); extant plants have either a trilobate or bilobate base morphology. This morphology appears early in ontogeny and has been observed in sporelings (F. D. Freund, personal observation). In trilobate species, there are three basal furrows, which run down the lateral face of the corm and join together at the distal end (Fig. 1A–C). These basal furrows divide the corm into three sections where

secondary growth ultimately results in the formation of the triple corm lobes (Stokey, 1909; Osborn, 1922; Bhambie, 1963; F. D. Freund, personal observation). By contrast, the bilobate species have only a single furrow, which runs in a line across the base of the corm, dividing it into two halves (Fig. 1D–F). While the plants do occasionally acquire additional lobes as they age, the base morphology—the minimum number of lobes the plants have before any elaboration—is consistent (Karrfalt and Eggert, 1977a, b). Also, there are other, rarer morphologies, such as the rhizomatous, mat-forming *I. tegetiformans* Rury (Rury, 1978) and the monolobate *I. andicola* (Amstutz) L.D. Gómez (formerly treated as *Stylites andicola* Amstutz; Amstutz, 1957). However, both *I. andicola* and *I. tegetiformans* begin life with a bilobate morphology before secondarily acquiring these alternate states (Rury, 1978; Karrfalt, 1984b; Tryon et al., 1994); their base morphology is bilobate.

Here, we explore the evolution of corm lobation across the most recent *Isoëtes* phylogeny (Larsén and Rydin, 2016). If *Isoëtes* has undergone evolutionary reduction in corm lobation number as Karrfalt and Eggart (1977b) hypothesized, we expect the ancestral state for extant *Isoëtes* to be trilobate, and the rate of transition from trilobate to bilobate to be higher than the rate of bilobate to trilobate. To test this hypothesis, we used reversible-jump Markov chain Monte Carlo (MCMC; Green, 1995) to explore multiple nonstationary statistical models of morphological evolution (Klopfstein et al., 2015) and to assess whether there is evidence for directional or even irreversible evolution in corm lobation morphology. Finally, we used simulations to examine our power to detect irreversible evolution on datasets of this size.

METHODS

Corm lobation characterization

Corm lobe numbers were collected from observations of fresh material, herbarium specimens (Appendix 1), and published descriptions (Appendix 2). We characterized lobation from fresh plants for eight species (Isoëtes appalachiana D.F. Brunt. & D.M. Britton [four individuals]; I. bolanderi Engelm. [20+ individuals]; I. eatonii R. Dodge [three individuals]; I. englemannii A. Braun [four individuals]; I. howellii Engelm. [200+ individuals]; I. nuttallii A. Braun ex Engelm. [400+ individuals]; I. occidentalis L.F. Hend. [15+ individuals]; and Isoëtes orcuttii A.A. Eaton [50+ individuals]). Each individual was cleaned of encrusting soil, then examined to determine the number of basal furrows and to assess the symmetry of the corm lobes. Plants of different ages were observed for each species when possible to get a sense of variability in lobation pattern as the plants aged, as well as to identify any unusual morphological outliers such as plants with asymmetric lobation or nonlinear basal furrows that may have occurred due to advanced age or damage to the corm. The I. nuttallii samples include two sporelings, one of which was still connected to the megaspore, which already showed distinctly trilobate arrangement of their rootlets. Taxa in which the plants have a single contiguous furrow and two symmetrical lobes were scored as "bilobate," while species with three linear furrows with symmetrical lobation around the base of the corm were scored as "trilobate" (Fig. 1).

For lobation numbers garnered from the literature, species with a definitive, single reported corm lobation value were assigned that number, while species with a range of reported corm lobe numbers were scored as "undefined" due to the possibility that the description is of a taxon with a range of corm morphologies, such as

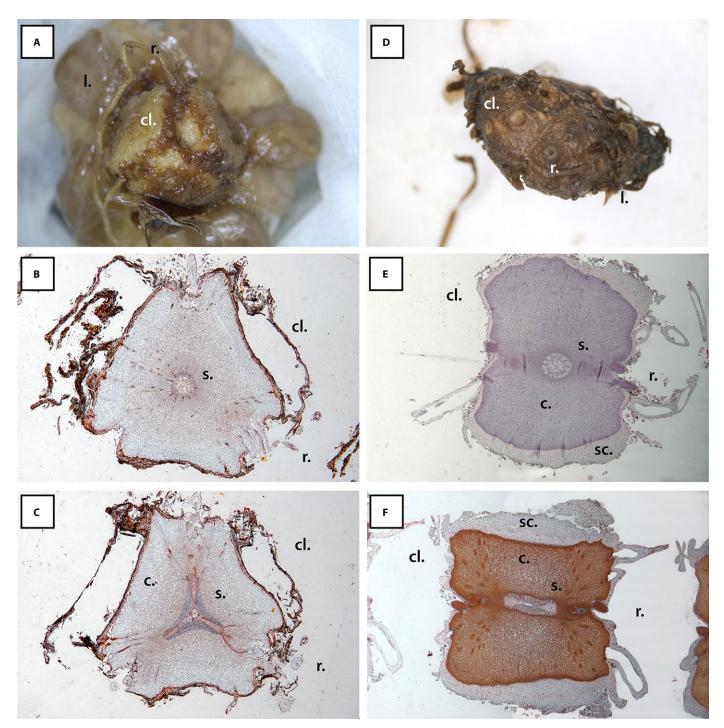


FIGURE 1. Comparative morphology of trilobate and bilobate corms. Trilobed corm: (A) *Isoëtes nuttallii* (F. Freund 266, UC) in proximal view with senesced cortical tissue removed (8.1×, 29 mm field of view); (B–C) transverse sections of *I. nuttallii* (F. Freund 169, RSA796374) below apical rosette (B) and near basal furrows (C) (32.5×, 7.1 mm field of view). Bilobed corm: (D) *I. howellii* (F. Freund 266.1, UC) in proximal view (24×, 9.5 mm field of view); (E–F) cross sections of *I. bolanderi* (F. Freund 10, RSA811643) below apical rosette (E) and near basal furrow (F) (24.5×, 9.3 mm field of view). All images were adjusted in Adobe Photoshop to improve visibility by using the image levels tool and the dodge tool to even out lighting and coloration. For herbarium acronyms, see Index Herbariorum (http://sweetgum.nybg.org/science/ih/). Labels: c. = cortex ground tissue, cl. = corm lobe, r. = rootlets, l. = leaves, s. = stele, and sc. = senesced cortical ground tissue.

I. tuckermanii A. Br. (Karrfalt and Eggert, 1977b; Croft, 1980), or due to the inclusion of multiple cryptic species within a single taxon (Appendix 2).

Additionally, three accessions were coded as "unknown" for lobation due to identification uncertainties: (1) One accession of *I. savatieri* Franchet was collected in Uruguay, outside of the

currently accepted range of the species (Hickey et al., 2003), and we were unable to examine the specimen to determine its identity or lobation. (2) Isoëtes histrix Bory & Durieu is a known species complex (Bagella et al., 2011, 2015), and specimens fall in two different areas of our phylogeny. One is sister to I. setacea Lam., a position that is consistent with morphology and with existing hypotheses of their relationship (Hoot et al., 2006; Bagella et al., 2011; Troia and Greuter, 2014); we treated this accession as correctly identified. The second accession falls phylogenetically distant from the first, and presumably is misidentified; we were unable to ascertain its true identity or morphology, so we coded it as unknown. (3) Isoëtes australis R.O. Williams also shows up in two places in the phylogeny: Clade A and Clade D. The Clade A plant was collected and identified by Dr. Carl Taylor, an authority on Isoëtes, so we treated this accession as correctly identified and coded the Clade D plant as unknown. All character states and literature sources can be found in Appendix 2.

Phylogenetic modeling

Ancestral character state reconstructions were performed on a posterior sample of 15,000 trees from Larsén and Rydin (2016) that we rooted on the bipartition between Clade A and the remainder of the genus (following Larsén and Rydin, 2016).

We employed reversible-jump MCMC (Green, 1995) in RevBayes (Höhna et al., 2016) to explore the space of all five possible continuous-time Markov models of phenotypic character evolution and to infer ancestral states. The reversible-jump MCMC sampled from the five models in proportion to their posterior probability. This approach enabled model-fit comparisons through Bayes factors (Kass and Raftery, 1995) and provided the opportunity to account for model uncertainty by making model-averaged ancestral state and parameter estimates (Madigan and Raftery, 1994; Kass and Raftery, 1995; Huelsenbeck et al., 2004; Freyman and Höhna, 2017). The five models of corm lobation evolution considered were as follows: a model with the rate of lobation gain and loss set to be equal (the 1-rate model); a model where the rates of lobation gain and loss are independent and nonzero (the 2-rate model); two irreversible models where the rate of either lobation gain or loss was fixed to zero; and lastly a model where both rates were fixed to zero. To test for directional evolution, we used nonstationary models of character evolution with root state frequencies that differed from the stationary frequencies of the process (Klopfstein et al., 2015).

Each of the five models was assigned an equal prior probability using a uniform set-partitioning prior. The root state frequencies were estimated using a flat Dirichlet prior. The rates of corm lobation gain and loss were drawn from an exponential distribution with a mean of one expected character state transition over the tree ($\lambda = \tau/1$, where τ is the length of the tree).

The MCMC was run for 22,000 iterations, where each iteration consisted of 48 MCMC proposals. The 48 proposals were scheduled randomly from six different Metropolis-Hastings moves that updated the sampled tree, root frequencies, and corm lobation gain and loss rate parameters. The first 2000 iterations were discarded as burn-in, and samples were logged every 10 iterations. Convergence of the MCMC was confirmed by ensuring that the effective sample size of all parameters was over 600. The results were summarized and plotted using the RevGadgets R package (https://github.com/revbayes/RevGadgets). The scripts that specify our model, run the analysis, and summarize results are available in the code repository at https://github.com/wf8/isoetes.

Simulations

To test how many observed characters are necessary to reliably infer irreversible evolution on a phylogeny the size of ours, we simulated 10 datasets with each of 1, 5, 10, 50, or 100 characters per tip (for a total of 50 simulations; note that our empirical dataset has a single character per tip). Each dataset was simulated under an irreversible model with the mean rate of corm lobation loss set to the value estimated by the irreversible model using the observed corm lobation data (2.39 changes per unit branch length). We performed the simulations using RevBayes over the maximum a posteriori phylogeny from the same tree distribution used to infer the ancestral states. For each of the 50 simulated datasets, an MCMC analysis was run for 11,000 iterations, with the first 1000 iterations dropped as burn-in. The model used was identical to that used for the observed corm lobation dataset, except that for the simulated datasets we fixed the maximum a posteriori phylogeny instead of integrating over the posterior distribution of trees.

Figure editing

Figure plates of prepared slides and preserved materials were imported into Adobe Photoshop and adjusted using the levels and dodge tools to improve image definition and to reduce uneven color levels resulting from variations in background exposure. No features that were part of the original images were removed.

RESULTS

Model-fit comparisons

The maximum a posteriori model of corm lobation evolution was the tri- to bi- irreversible model (which did not allow transitions from the bilobate to the trilobate state) with a posterior probability of 0.38 (Table 1). This tri- to bi- irreversible model was weakly supported over the 1-rate and 2-rate reversible models (Bayes factor = 1.26 and 1.21, respectively; Kass and Raftery, 1995); however, all three models were strongly supported over the bi- to tri- irreversible model. Since the Bayes factor support for the best-supported model over the next two was negligible, we focus mostly on the model-averaged parameter estimates and ancestral states.

Model-averaged parameter estimates and ancestral states

The model-averaged estimated rate of transition from trilobate to bilobate forms was significantly nonzero (mean = 2.17 changes per unit branch length, 95% HPD interval: 0.015–5.69), whereas the rate of bilobate to trilobate transitions was not significantly nonzero (mean = 0.82, 95% HPD interval: 0.0–3.35; Fig. 2). The model-averaged maximum a posteriori ancestral state of *Isoëtes* was trilobate with a posterior probability of 1.0 (Fig. 3). The ancestral state of the New World clade ("Clade E-2"; Fig. 3) was bilobate with a posterior probability of 0.99. The bilobate morphology arose independently in six places over the phylogeny (Fig. 3). No reversals from bilobate to trilobate were inferred. All species with unknown base corm lobation characters (*I. stevensii* J.R. Croft, *I. habbemensis* Alston, and *I. hallasanensis* H.K. Choi, Ch. Kim & J. Jung) were derived from a trilobate most recent common ancestor, with a posterior probability near 1.0.

TABLE 1. Comparisons of models of corm lobation evolution.

Model	Model posterior probability	Bi- to tri- irreversible	Tri- to bi- irreversible	1-rate	2-rate
Bi- to tri- irreversible	0.0	-	<1	<1	<1
Tri- to bi- irreversible	0.38	>1000	-	1.26	1.21
1-rate	0.30	>1000	<1	-	<1
2-rate	0.32	>1000	<1	1.04	-

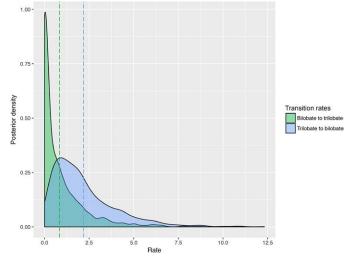


FIGURE 2. Model-averaged posterior densities of transition rates between corm lobation states. Mean values are represented by dashed lines. The rate of transition from trilobate to bilobate forms was significantly nonzero (mean = 2.17, 95% HPD interval: 0.015–5.69), whereas the rate of bilobate to trilobate transitions was not significantly nonzero (mean = 0.82, 95% HPD interval: 0.0–3.35). Transition rates are reported in changes per unit branch length.

Simulations

For simulated datasets with a single character, the true irreversible model was at best weakly supported over the other models; the mean posterior probability of the true irreversible model was 0.37 (range: 0.22–0.44). As the number of characters increased, the posterior probability of the true model increased (Fig. 4). With 100 characters the support for the true model was strong; the mean posterior probability was 0.85 (range: 0.80–0.92).

DISCUSSION

Our analyses support the hypothesis of directional reduction in lobe number over time in *Isoëtes*, with the best-supported model being one of irreversible evolutionary reduction. Additionally, even when incorporating model and phylogenetic uncertainty and allowing for reversals, the model-averaged estimate of the transition rate from trilobate to bilobate was much higher than the estimate of the transition rate from bilobate to trilobate (the latter rate was not significantly nonzero). Furthermore, we found strong support for the ancestral state for all extant *Isoëtes* being trilobate, indicating that there have been multiple convergent reductions of the corms to bilobate. These bilobate forms are nested deeply within clades of trilobate plants. These results support the hypothesis that crown *Isoëtes* has continued a reduction in corm morphology from the larger arborescent lycophytes. Additionally, while the bilobate form has emerged multiple times throughout the phylogeny, it is the dominant morphology in only one major clade: the "American clade" (Clade E-2, *sensu* Larsén and Rydin, 2016; Fig. 3). In this clade, nearly all species have a bilobate morphology.

The simulations demonstrate that the relatively weak support for the irreversible reduction model over the reversible models is likely due to the inherent limitation in statistical power of a single observed morphological character over a phylogeny of this size. Repeating this analysis with a larger, more densely sampled phylogeny, or one that incorporates fossil data, might find stronger support for the irreversible model of corm morphology evolution. Nevertheless, it is in cases like this, where no single model is decisively supported over others, that reversible-jump MCMC and Bayesian model-averaging demonstrate their utility for testing phylogenetic hypotheses of character evolution (Huelsenbeck et al., 2004; Freyman and Höhna, 2017).

While the bilobate morphology dominates only Clade E-2, it does occur in several other areas of the phylogeny (Fig. 2 and Appendix 2). In areas where Clade E-2 co-occurs with others, such as South America and the west coast of the United States, corm lobation is useful in determining to which clade a plant belongs. However, outside the range of Clade E-2, other bilobate taxa occur as single species nested within larger trilobate clades. As such, using corm lobation outside the Americas to assign plants with unknown phylogenetic placement is not advisable, since they may represent other independent evolutions of the character state.

When assessing corm lobation numbers, it is important to determine whether the corm lobe numbers are the base lobe numbers or additional lobes that have developed as the plant ages (Stokey, 1909; Karrfalt and Eggert, 1977a, b; F. D. Freund, personal observation). While Karrfalt and Eggert (1977b) reported a propensity for gaining additional lobes in their study of I. tuckermanii (68% bilobate, 30% trilobate, 2% tetralobate), other observers, working on other taxa, have not found this degree of variability (Engelmann, 1882; F. D. Freund, personal observation of >200 specimens of I. howellii and >400 specimens of I. nuttallii). In fact, we have observed a total of <10 I. howellii and I. nuttallii specimens that were not either bilobate or trilobate, respectively (F. D. Freund, personal observation). In I. nuttallii, the trilobate morphology was readily observable even in very young plants, including two sporelings, one of which was still attached to the megaspore (F. D. Freund, personal observation). These results suggest that the degree of variability of I. tuckermanii may be species-specific, and they add further support to the value of sampling multiple individuals when examining and designating corm lobation numbers within a species. If at all possible, it is additionally worthwhile to look at very young plants, especially sporelings, to determine the base lobe number.

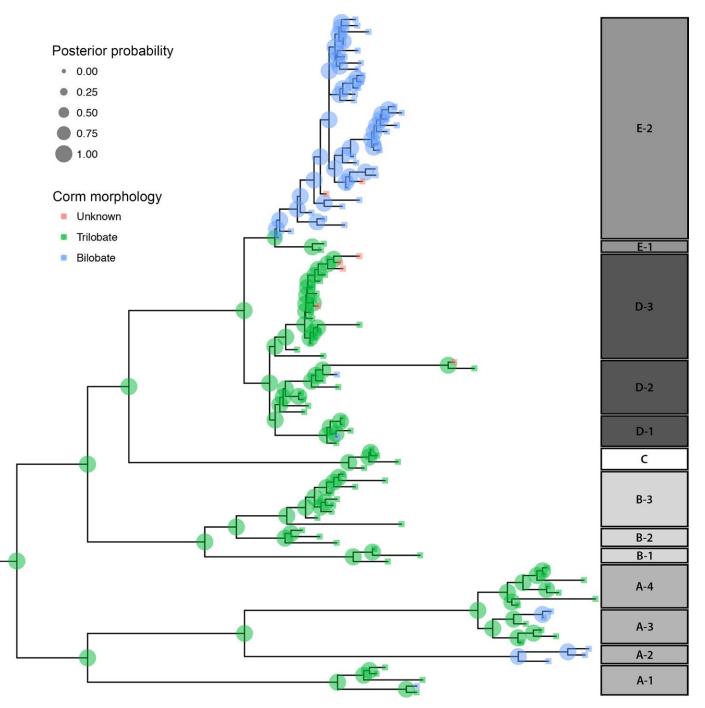


FIGURE 3. Bayesian model-averaged ancestral states of *Isoëtes* corm lobation inferred over the posterior tree sample from Larsén and Rydin (2016). Ancestral states are summarized on the maximum a posteriori phylogenetic tree. The size of the circles at each node represents the posterior probability of the most probable ancestral state, and the color represents the state: green = trilobate corms, blue = bilobate corms, and red = unknown. Boxes on the right reflect subclade designations to broad subclade: A = Gondwanan, B = Laurasian, C = Italian, D = Austro-Asian, and E = New World or American. For specific placement of taxa into subclades, see Appendix 2.

CONCLUSIONS

Our results support the hypothesis that crown *Isoëtes* has continued an evolutionary reduction in corm morphology from the larger arborescent lycophytes, with the best-supported model being one of irreversible evolutionary reduction. However, results from our simulation study showed that a dataset of this size only has weak statistical power to support irreversible models of character evolution, emphasizing the need for broader sampling of extant taxa. When we accounted for the uncertainty in character evolution models by making model-averaged estimates, we found strong support for the hypothesis of directional evolutionary reduction

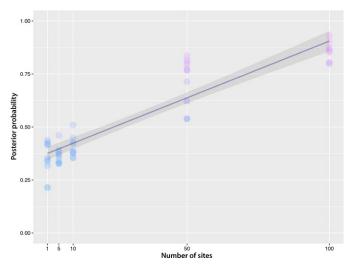


FIGURE 4. Statistical power to detect irreversible evolution as a function of the number of characters available. Each point plotted represents a different simulation replicate. The *y*-axis shows the posterior probability of the true irreversible model of character evolution. The *x*-axis shows the number of simulated characters (sites or columns in the data matrix). Ten replicates were simulated for 1, 5, 10, 50, and 100-character datasets, resulting in a total of 50 simulated datasets.

in corm number, with the rate of lobe loss estimated to be much higher than the rate of lobe gain. Furthermore, we found strong support that the ancestral state for all extant *Isoëtes* was trilobate, indicating that there have been multiple convergent reductions of the corms to a bilobate state.

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APPENDIX 1. Vouchers examined for morphological determinations in this study.

Isoëtes bolanderi: RSA (811637, 811638, 811639, 811643). *I. howellii*: RSA (796366, 796367, 796368, 796369, 796370, 796371), UC (F. Freund 255, 266.1). *I. nuttallii*: RSA (796374, 796375, 796376), UC (F. Freund 256, 257, 258, 259, 261, 263, 264, 265, 266). *I. occidentalis*: RSA (811640, 811641, 811642). *I. orcuttii*: UC (F. Freund 254, 262). *I. storkii*: GH00021453, CM0102, NY00144272 (virtual herbarium sheets). For herbarium acronyms, see Index Herbariorum (http://sweetgum.nybg.org/science/ih/).

APPENDIX 2. Corm lobation numbers for Isoëtes. Clade designations follow Larsén and Rydin (2016). For herbarium acronyms, see Index Herbariorum (http://
sweetgum.nybg.org/science/ih/).

Species	Observed by F. Freund	Clade	Lobes	Locality	Sources
<i>lsoëtes abyssinica</i> Choiv. – synonym of <i>I. welwitschii</i> (sensu Verdcourt and Beentje, 2002)		B-3	3	Africa, Ethiopia	Pfeiffer, 1922; Cook, 2004; Crouch et al., 2011
lsoëtes acadiensis Kott		?	2	USA (MA, ME, NH, NJ, NY, VT), CAN (NB, NF, NS)	Taylor et al., 1993
<i>lsoëtes adspersa</i> A. Braun		?	3	Algeria	Pfeiffer, 1922
<i>lsoëtes aequinoctialis</i> A. Braun		B-3	3	Angola, South Africa, Mali, Ghana, Tanzania, zambia, Zimbabwe, south to RSA: NAM, NC	Pfeiffer, 1922; Cook, 2004; Crouch et al., 2011
lsoëtes alpina Kirk		D-2	3	New Zealand	Pfeiffer, 1922
<i>lsoëtes alstonii</i> C.F. Reed & Verdc.		?	2	Egypt; Mozambique; Namibia; Sudan; Tanzania, United Republic of; Zambia; Zimbabwe	Crouch et al., 2011
<i>soëtes amazonica</i> A. Braun ex Kuhn		A-3	3	Brazil (north [Pará])	Pfeiffer, 1922
<i>soëtes anatolica</i> Prada & Rolleri		С	3	Turkey	Prada and Rolleri, 2005
lsoëtes andicola (Amstutz) L.D. Gómez (formerly Stylites andicola Amstutz)	Live plants	E-2	2	Peru (Lima, Pasco, Junin, Cuzco, Puno), Bolivia	Karrfalt, 1984b; Tryon et al., 1994
lsoëtes andina Spruce ex Hook.– synonym of I. triquetra		E-2	2	South America, Peru	Pfeiffer, 1922
<i>soëtes appalachiana</i> D.F. Brunt. & D.M. Britton	Live plants	E-2	2	USA (FL, GA, NC, PA, SC, VA)	Brunton and Britton, 1997
<i>soëtes araucaniana</i> Macluf & Hickey		?	2	Malleco (Chile)	Macluf and Hickey, 2007
soëtes asiatica Makino		E-2	2	Kamchatka, Sakhalin, the Kuriles, and Japan	Yi and Kato, 2001
<i>lsoëtes australis</i> R.O. Williams		A-2, D-2	2	Bruce Rock, West Australia	Williams, 1944
<i>soëtes azorica</i> Durieu ex Milde		?	2	Islands of Azores	Pfeiffer, 1922
<i>lsoëtes bolanderi</i> Engelm.	RSA (811637, 811638, 811639, 811643)	E-2	2	USA (AZ, CA, CO, ID, MT, NM, NV, OR, UT, WA, WY), CAN (AB)	Pfeiffer, 1922; Taylor et al., 1993
lsoëtes boliviensis U. Weber		?	2	Peru (Cajamarca, San Martin, Ancash, Lima, Pasco, Junin, Ayacucho, Cuzco, Puno), Bolivia (La Paz)	Tryon et al., 1994
<i>lsoëtes boryana</i> Durieu		?	3	France (Landes)	Pfeiffer, 1922
<i>lsoëtes bradei</i> Herter		A-3	3	Brazil, Sudeste (São Paulo)	Brazilian Flora 2020 (in construction). Rio de Janeiro Botanical Garden. Available at http://floradobrasil.jbrj.gov. br/, accessed Aug 15, 2016; Hickey, 1990
<i>Isoëtes braunii</i> Unger		?	2	USA (NH, VT, MA), CAN	Pfeiffer, 1922
<i>lsoëtes brevicula</i> E.R.L. Johnson		D-2	3	Western Australia	FloraBase, the Western Australian Flora, Western Australian Herbarium. https:// florabase.dpaw.wa.gov.au/,
					accessed Aug 15, 2016

Continued

APPENDIX 2 Continued

Species	Observed by F. Freund	Clade	Lobes	Locality	Sources
<i>lsoëtes butleri</i> Engelm.		?	2	USA (AL, AR, GA, IL, KS, KY, MO, OK, TN, TX)	Pfeiffer, 1922
<i>lsoëtes capensis</i> A.V. Duthie <i>Isoëtes caroli</i> E.R.L. Johnson		A-1 D-2	3 3	South Africa (Western Cape) Western Australia	Cook, 2004; Crouch et al., 2011 FloraBase, the Western Australian Flora, Western Australian Herbarium. https:// florabase.dpaw.wa.gov.au/, accessed Aug 15, 2016
<i>lsoëtes caroliniana</i> (A.A. Eaton) Luebke		?	2	USA (NC, TN, VA, WV)	Taylor et al., 1993
<i>lsoëtes coreana</i> Y.H. Chung & H.K. Choi		D-3	3	Korea	Chung and Choi, 1986
<i>lsoëtes coromandeliana</i> L.f.		A-4	3	India	Pfeiffer, 1922
<i>lsoëtes cubana</i> Engelm. ex Baker		A-3	3	Cuba (Pinao del Rio), Belize, Mexico (Yucatan)	Pfeiffer, 1922
<i>lsoëtes dispora</i> Hickey		?	2	Laguna Tembladera, Lambayeque, Peru	Tryon et al., 1994
<i>Isoëtes dixitei</i> Shende		B-3	3	Panchgani, India	Pant and Srivastava, 1962
<i>lsoëtes drummondii</i> A. Braun		D-2	3	Australia (South Australia, Western Australia, Victoria, New South Wales)	Osborn, 1922; Pfeiffer, 1922
<i>Isoëtes duriei</i> Bory		B-1	3	Algeria, Corsica, France, Italy, Turkey	Pfeiffer, 1922
<i>Isoëtes eatonii</i> R. Dodge	Live plants.	E-2	2	USA (CT, MA, NH, NJ, NY, PA, VT), CAN (ON)	Pfeiffer, 1922
<i>lsoëtes echinospora</i> Durieu		E-2	2	CAN (AB, BC, MB, NB, NT, NS, ON, PE, QC, SK, YT), USA (AK, CA, CO, ID, ME, MA, MI, MN, MT, NH, NJ, OH, OR, PA, VT, WA, WI), British Isles, Circum boreal	Pfeiffer, 1922
<i>Isoëtes elatior</i> A. Braun		?	3	Tasmania	Pfeiffer, 1922
<i>lsoëtes eludens</i> J.P.Roux, Hopper & Rhian J.Sm.		?	3	South Africa (Kamiesberg Mountains, Namaqualand)	Roux et al., 2009; Crouch et al., 2011
<i>lsoëtes engelmannii</i> A. Braun (includes <i>I. valida</i>)	Live plants.	E-2	2	CAN (ON), USA (AI, AR, CT, DE, FL, GA, IL, IN, KY, MD, MA, MI, MO, NH, NJ, NY, NC, OH, PA, RI, SC, TN, VT, VA, WV)	Pfeiffer, 1922; Taylor et al., 1993
<i>lsoëtes flaccida</i> A. Braun		E-2	2	USA (FL, GA)	Pfeiffer, 1922; Taylor et al., 1993
Isoëtes flettii (A.A. Eaton) N. Pfeiff.		?	2	USA (WA - Spanaway Lake)	Pfeiffer, 1922
<i>Isoëtes foveolata</i> A.A. Eaton ex R. Dodge		?	2	USA (CT, MA, NH), CAN (ONT)	Pfeiffer, 1922
<i>lsoëtes gardneriana</i> Kunze ex Mett.		?	3	Brazil (Midwest [Goias]), Paraguay	Pfeiffer, 1922
<i>lsoëtes georgiana</i> Luebke		E-2	2	USA (GA)	Taylor et al., 1993
<i>lsoëtes giessii</i> Launert		?	3	Namibia (Erongo Mountains and in seasonally wed depressions in acacia scrub)	Crouch et al., 2011
<i>Isoëtes gunnii</i> A. Braun		D-2	3	Tasmania (Lake Fenton on Mt. Field)	Pfeiffer, 1922
Isoëtes habbemensis Alston		D-3	(2)-3-(4)	New Guinea	Croft, 1980
<i>Isoëtes hallasanensis</i> H.K. Choi, Ch. Kim & J. Jung		D-3	3	Jeju Island, Korea	Choi et al., 2008
<i>lsoëtes hawaiiensis</i> W.C. Taylor & W.H. Wagner		E-2	2	Hawaii	Taylor et al., 1993
Isoëtes heldreichii Wettst.		?	3	Greece (plains of Thessaly, base of Pindus Mts.)	Pfeiffer, 1922
<i>Isoëtes herzogii</i> U. Weber		E-2	2	Bolivia	Weber, 1922
Isoëtes hewitsonii Hickey		?	2	Celendin, Cajamarca, Peru	Tryon et al., 1994
<i>lsoëtes histrix</i> Bory & Durieu		E-1, E-2	3	Algeria, Italy, France, islands of the Mediterranean	Pfeiffer, 1922
<i>lsoëtes howellii</i> Engelm.	RSA (796366, 796367, 796368, 796369. 796370, 796371)	E-2	2	USA (MT, ID, WA, OR, CA)	Pfeiffer, 1922; Taylor et al., 1993
<i>Isoëtes humilior</i> A. Braun	,	D-1	2	Tasmania (So. River Esk)	Pfeiffer, 1922; Taylor et al., 1993

APPENDIX 2 Continued

Species	Observed by F. Freund	Clade	Lobes	Locality	Sources
<i>Isoëtes inflata</i> E.R.L. Johnson		?	2	Australia (West Australia)	FloraBase, the Western Australian Flora, Western Australian Herbarium. https:// florabase.dpaw.wa.gov.au/, accessed Aug 15, 2016 Pfeiffer. 1922
<i>Isoëtes japonica</i> A. Braun		D-3	3	Japan (Yokohama)	/ -
<i>lsoëtes jejuensis</i> H.K. Choi, Ch. Kim & J. Jung		D-3	3	Jeju Island, Korea	Choi et al., 2008
<i>Isoëtes kirkii</i> A. Braun		D-1	3	New Zealand	Pfeiffer, 1922
<i>lsoëtes labri-draconis</i> N.R. Crouch		?	3	South Africa (Drakensberg Range in KwaZulu-Natal)	Crouch et al., 2011
Isoëtes lacustris L.		E-2	2	CAN (MB, NB, NL, NT, NS, ON, OQ, SK), USA (ME, MA, MI, MN, NH, NY, VT, WI)	Pfeiffer, 1922; Taylor et al., 1993
<i>lsoëtes laosiensis</i> C. Kim & H.K. Choi		A-4	3	Laos	Kim et al., 2010
Isoëtes lechleri Mett.		?	2	Argentina, Equador, Colombia, Peru	Pfeiffer, 1922; Tryon et al., 1994
lsoëtes lithophila N. Pfeiff.		E-2	2	USA (TX)	Pfeiffer, 1922; Tryon et al., 1994
lsoëtes longissima Bory & Dur.		B-3	3		Pfeiffer, 1922
– As I. velata forma longissima (sensu Pfeiffer, 1922)		D-2	2	Algeria	Pieliiel, 1922
Isoëtes macrospora Durieu		?	2	Newfoundland to USA (MN)	Pfeiffer, 1922
Isoëtes malinverniana Ces. &		C	3		Pfeiffer, 1922
De Not.				Italy	
<i>lsoëtes maritima</i> Underw.		E-2	2	Alaska, British Columbia, Washington	Underwood, 1888
<i>lsoëtes martii</i> A. Braun ex Kuhn		E-2	2	Brazil (Southeast [Minas Gerais, Rio de Janeiro], South [Rio Grande do Sul])	Brazilian Flora 2020 (in construction). Rio de Janeiro Botanical Garden. Available at http://floradobrasil.jbrj.gov.br/ accessed Aug 15, 2016
<i>lsoëtes maxima</i> Hickey, Macluf & Link-Pérez		?	2	Brazil (South [Rio Grande do Sul])	Hickey et al., 2009
<i>lsoëtes melanopoda</i> Gay & Durieu		E-2	2	USA (AL, AR, GA, IA, ID, IL, IN, KS, KY, LA, MN, MO, MS, MT, NC, NE, NJ, OK, SC, SD, TN, TX, UT, VA)	Pfeiffer, 1922; Taylor et al., 1993
<i>lsoëtes melanospora</i> Engelm.		E-2	2	USA (GA - Stone Mtn.)	Pfeiffer, 1922; Taylor et al., 1993
<i>Isoëtes mexicana</i> Underw.		E-2	2	Mexico (Chihuahua, Hidalgo, Mexico, Morelos, Michoacan)	Pfeiffer, 1922
<i>Isoëtes mongerensis</i> E.R.L. Johnson		?	3	Western Australia	FloraBase, the Western Australian Flora, Western Australian Herbarium. https:// florabase.dpaw.wa.gov.au/, accessed Aug 15, 2016
<i>lsoëtes muelleri</i> A. Braun		D-3	3	Eastern Australia (Rockhampton)	Pfeiffer, 1922
<i>lsoëtes muricata</i> Durieu		E-2	2	North America	Pfeiffer, 1922, as <i>I. braunii</i>
<i>lsoëtes nigritiana</i> A. Braun		?	3	Nigeria (along the Niger River, Nupe)	Pfeiffer, 1922
<i>lsoëtes novo-granadensis</i> H.P. Fuchs		E-2	2	Columbia, Peru	Tryon et al., 1994
<i>lsoëtes nuttallii</i> A. Braun ex Engelm.	RSA (796374, 796375, 796376)	B-2	3	USA (CA, OR, WA), CAN (Vancouver)	Pfeiffer, 1922; Taylor et al., 1993
<i>lsoëtes occidentalis</i> L.F. Hend.	RSA (811640, 811641, 811642)	E-2	2	USA (CA, WY, CO, ID)	Pfeiffer, 1922; Taylor et al., 1993
<i>lsoëtes olympica</i> A. Braun		B-3	3		Pfeiffer, 1922
<i>Isoëtes orcuttii</i> A.A. Eaton	Live plants	B-2	3	USA (CA)	Pfeiffer, 1922; Taylor et al., 1993
lsoëtes ovata N. Pfeiff.	Enc plurio	?	3	French Guiana, Guyana	Pfeiffer, 1922
Isoètes ovata N. Pieni. Isoètes panamensis Maxon & C.V. Morton	Museo Nacional de Costa Rica (sheet no. not recorded)	? A-3	3 2	Brazil (northeast [Maranhão, Bahia], midwest [Mato Grosso])	Brazilian Flora 2020 (in construction). Rio de Janeiro Botanical Garden. Available at http://floradobrasil.jbrj.gov.br/

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accessed Aug 15, 2016

APPENDIX 2 Continued

species	Observed by F. Freund	Clade	Lobes	Locality	Sources
oëtes parvula Hickey		?	2	Laguna Yaurihuiri, Ayacucho, Perú	Tryon et al., 1994
<i>pëtes pedersenii</i> H.P. Fuchs ex .I. Meza & Macluf		?	2	Argentina (Corrientes - Mburucuyá National Park)	Macluf et al., 2010
oëtes philippinensis Merr. & R.H. Perry		D-3	3	Philippine islands	Merrill and Perry, 1940
pëtes piperi A.A. Eaton		?	2	USA (WA)	Pfeiffer, 1922
<i>pëtes pringlei</i> Underw.		?	2	Mexico (Guadalajara, state of Jalisco)	Pfeiffer, 1922
pëtes prototypus D.M. Britton		?	2	CAN (NB, NS), USA (MA)	Taylor et al., 1993
pëtes pseudojaponica M. Fakamiya, Mits. Watan. & K. Dno		D-3	3	Japan	Takamiya et al., 1997
<i>pëtes quiririensis</i> J.B.S. Pereira & Labiak		?	2	Brazil (Serra do Quoriri)	Pereira and Labiak, 2013
<i>pëtes riparia</i> Engelm. ex A. Braun		?	2	CAN (southern region), USA (New England south to DE and PA)	Pfeiffer, 1922
<i>etes saccharata</i> Engelm.		?	2	USA (DE, DC, MD, VA)	Pfeiffer, 1922
ëtes sampathkumarinii L.N. ao		D-2	2	India	Rao, 1944
ëtes saracochensis Hickey		?	2	Laguna Saracocha, Puno, Peru	Tryon et al., 1994
pëtes savatieri Franch.		E-2	2	Argentina (Puerto Bueno)	Hickey et al., 2003
oëtes schweinfurthii Baker		A-4	3	South Africa (Namibia, Botswana, Limpopo), Sudan, Madagascar, Zambia	Pfeiffer, 1922; Cook, 2004; Crouch et al., 2011
pëtes setacea Lam.		E-1	3	France (Montpellier, Hérault, Pyréneés-Orientales); Morocco; Portugal; Spain (Baleares)	Pfeiffer, 1922
<i>iëtes stellenbossiensis</i> A.V. Duthie		A-1	3	South Africa (western part of Western Cape province)	Cook, 2004; Crouch et al., 201
<i>etes stephansenii</i> A.V. Duthie		A-1	2	South Africa (flats around Stellenbosch in the Western Cape)	Crouch et al., 2011
pëtes stevensii J.R. Croft		D-3	(2)-3-(4)	New Guinea	Croft, 1980
<i>pëtes storkii</i> T.C. Palmer	GH00021453, CM0102, NY00144272 (virtual herbarium sheets)	E-2	2	Costa Rica	JStor Global Plants virtual herbarium. http://plants.jsto org/search?filter=name& so=ps_group_by_genus_ species+asc&Query= isoetes+storkii, accessed Dec 28, 2016
<i>iëtes subinermis</i> Cesca – ynonym of <i>I. histrix</i>		B-1	3		Cesca and Peruzzi, 2001
etes taiwanensis De Vol	Live plants	D-3	3	Taiwan	Chiang, 1976
pëtes tegetiformans Rury		E-2	2	USA (GA)	Rury, 1978; Taylor et al., 1993
<i>pëtes toximontana</i> Musselman & J.P. Roux		A-1	3	South Africa (Giftberg endemic)	Cook, 2004; Crouch et al., 201
<i>pëtes transvaalensis</i> Jermy & Schelpe		?	3	Lesotho, South Africa (Limpopo, Mpumalanga, Free State, KwaZulu-Natal)	Cook, 2004; Crouch et al., 201
pëtes triquetra A. Braun		E-2	2	Peru	Pfeiffer, 1922
<i>bëtes truncata</i> (A.A. Eaton) Clute		?	2	CAN (Vancouver Island), USA (Alaska)	Pfeiffer, 1922
pëtes tuerckheimii Brause		E-2	2	Dominican Republic (Santo Domingo)	Pfeiffer, 1922
<i>pëtes valida</i> (Engelm.) Clute		E-2	2	USA (NC, TN, VA, WV)	Taylor et al., 1993
pëtes velata A. Braun		B-3	3	Italy (Algeria, Corsica, Sicily)	Pfeiffer, 1922
<i>pëtes welwitschii</i> A. Braun ex Kuhn		?	3	Angola, South Africa (Mpumalanga, KwaZulu-Natal), Madagascar	Pfeiffer, 1922; Cook, 2004; Crouch et al., 2011
pëtes wormaldii R. Sim		?	3	South Africa (Eastern Cape - small area between Grahamstown and East London)	Pfeiffer, 1922; Cook, 2004; Crouch et al., 2011
		D-3	3	China	Wang et al., 2002