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Ecological determinants of Potamogeton taxa in glacial lakes: assemblage composition, species richness, and species-level approach --Manuscript Draft--

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Abstract:	Aquatic macrophytes are important primary producers in glacial lakes that provide critical habitat, improve water quality, and are reliable indicators of ecosystem condition. The diverse Potamogeton genus includes over 80 species of macrophytes that occur across a broad geographic range and have variable response to environmental conditions. This study evaluated how environmental and spatial variables structure assemblage composition and species richness of Potamogetons in the US states of Minnesota and Wisconsin. Variation partitioning based on partial redundancy analysis and partial linear regression was used to study the relative contribution of local, climate and spatial variables in explaining assemblage composition and species richness. Models were also developed for sixteen individual Potamogeton species using partial linear regression. Assemblage composition and total species richness was better explained by the pure effects of spatial and local variables as compared to the pure effects of climate variables. The spatial groupings among lakes characterized a strong latitudinal gradient of local variables related to eutrophication across the study region. Models for individual species were similar although some were disproportionately described by specific categories of explanatory variables. For example, invasive Potamogeton crispus was more tolerant of elevated trophic state than most species and was also described by a strong spatial grouping of lakes near a large urban area. This analysis is the first regional evaluation of factors related to the distribution of this ecologically important genus and the importance of landscape-level approaches to ecological conservation is emphasized.
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To:

Dr. Stuart E.G. Findlay Editor-in-Chief Aquatic Sciences

From:

Dr. Marcus W. Beck US Environmental Protection Agency beck.marcus@epa.gov, 850-934-2480

Enclosed please find our manuscript, entitled "Ecological determinants of *Potamogeton* taxa in glacial lakes: assemblage composition, species richness, and species-level approach", to be considered as a research article in Aquatic Sciences. The *Potamogeton* genus includes over 80 species of aquatic macrophytes with wide ranges that cover almost all freshwater systems across the globe. Our manuscript presents the first evaluation of this diverse genus to describe environmental and spatial factors that drive variation in community structure on a regional scale. A large dataset of more than 200 lakes in the Midwest United States was used to characterize variation attributed to spatial, local, and climate variables. Our results highlight the importance of landscape-level approaches to ecological conservation. The conclusions can also be generalized to freshwater communities of glacial lakes in other regions (e.g., northern Europe).

Due to misunderstandings with the reviewers in the past, we kindly request that our reviewers are ecologists and not taxonomically-oriented botanists. We consider that ecologists can give us a fair review, as our manuscript deals with general ecological questions which are not necessarily well-known to taxonomists. Please do not hesitate to contact us with questions regarding our manuscript. We greatly appreciate the opportunity to publish our work in Aquatic Sciences.

Respectfully,

Marcus W. Beck

Ecological determinants of *Potamogeton* taxa in glacial lakes: assemblage composition, species richness, and species-level approach

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Abstract

Aquatic macrophytes are important primary producers in glacial lakes that provide critical habitat, improve water quality, and are reliable indicators of ecosystem condition. The diverse *Potamogeton* genus includes over 80 species of macrophytes that occur across a broad geographic range and have variable response to environmental conditions. This study evaluated how environmental and spatial variables structure assemblage composition and species richness of *Potamogetons* in the US states of Minnesota and Wisconsin. Variation partitioning based on partial redundancy analysis and partial linear regression was used to study the relative contribution of local, climate and spatial variables in explaining assemblage composition and species richness. Models were also developed for sixteen individual *Potamogeton* species using partial linear regression. Assemblage composition and total species richness was better explained by the pure effects of spatial and local variables as compared to the pure effects of climate variables. The spatial groupings among lakes characterized a strong latitudinal gradient of local variables related to eutrophication across the study region. Models for individual species were similar although some were disproportionately described by specific categories of explanatory variables. For example, invasive *Potamogeton crispus* was more tolerant of elevated trophic state than most species and was also described by a strong spatial grouping of lakes near a large urban area. This analysis is the first regional evaluation of factors related to the distribution of this ecologically important genus and the importance of landscapelevel approaches to ecological conservation is emphasized.

Keywords: Aquatic plants; Dispersal; Glacial lakes; Metacommunity; Metapopulation; *Potamogeton* species

Introduction

Species distributions are explained by local environmental conditions, historical characteristics and biotic factors, of which species dispersal has intrigued scientists for centuries (Levin 1992). Dispersal is generally described as the movement of species from one location to another across a landscape and it strongly affects how individuals, populations and communities are organised among habitats (Logue et al. 2011; Baguette et al. 2013). A suitable habitat for the individual species occurs as a network of idealized habitat patches, varying in area, degree of isolation and quality, and is surrounded by equally unsuitable habitats (Hanski 1998). Dispersal among habitats with different quality leads to colonization of new suitable areas and extirpations of populations from unsuitable habitats. Populations which are connected in a landscape by dispersal are regarded as metapopulations (Hanski 1991; 1998). For dispersal among connected communities (i.e., metacommunities), the relative importance of environmental characteristics and dispersal in structuring metacommunities varies (Leibold et al. 2004; Heino 2011). Species sorting is primarily influenced by niche processes, although a minimum level of dispersal is needed for species to shift among suitable habitats. In the remaining three metacommunity perspectives (i.e., mass effects, neutral theory, and patch dynamics), dispersal or dispersal limitation are the most important forces explaining variation in metacommunities. Dispersal-related spatial processes can be particularly important in freshwater ecosystems which are distinct environments surrounded by an uninhabitable terrestrial matrix. Moreover, lake species are more dependent on dispersal compared to species in rivers, which are better connected through hydrologic networks (Heino 2011). However, recent studies have suggested that dispersal limitation between lakes and rivers is more likely related to biological groupings rather than differences in habitats (Padial et al. 2014; Alahuhta et al. 2015; Heino et al. 2015). Characterising the relative effects of forcing mechanisms of species dispersal could greatly inform understanding of ecological processes in aquatic systems.

Aquatic macrophytes are important biological components of lake communities and are considered good indicators of long-term changes of freshwater ecosystems. Aquatic macrophytes respond to reduced light availability, increased sedimentation and nutrient concentrations, and hydromorphological changes, often originating from anthropogenic activities at different temporal scales (Lachavanne et al. 1992; Padial et al. 2009; Bornette and Puijalon 2011; Beck et al. 2014). Moreover, macrophytes have an essential functional role in freshwater ecosystems, as they provide habitat and shelter, breeding areas, and food resources for other aquatic and terrestrial species (Carpenter and Lodge 1986; Schmidt et al. 2005). One of the most important and diverse genera of macrophytes is *Potamogeton*, which includes over 80 species with wide ranges covering almost all global freshwater systems (Cook 1990). Potamogeton species have variable responses across environmental conditions and vary in morphology between species (Crow et al. 2000; Vestergaard and Sand-Jensen 2000; Heegaard et al. 2001; Chambers et al. 2008). Most *Potamogeton* species grow in alkaline waters, but some species, like P. amplifolius, P. natans, P. Epihydrus, and P. gramineus, can favour more acidic conditions (Toivonen and Huttunen 1995; Crow et al. 2000; Vestergaard and Sand-Jensen 2000). Potamogeton amplifolius, P. praelongus, and P. robbinsii are often found in deeper parts of lakes (Chambers and Kalff 1985; Crow et al. 2000). The morphology of the *Potamogeton* taxa varies from emergent broad-leaved plants (e.g., *P. Illinoensis*) to submerged thin-leaved species (e.g., P. diversifolius). In addition, vegetative and reproductive morphology varies considerably across the taxon (Crow et al. 2000). Although knowledge of Potamogeton species responses to local environmental conditions is relatively well-documented, little is known on how distribution of these species is affected by dispersal across the landscape.

The influence of water quality and climate in explaining aquatic plant distributions has rarely been studied within the same work (but see Alahuhta et al. 2011; Kosten et al. 2011), whereas the effects of local environmental conditions are relatively well known. Aquatic plants, and more importantly submerged species, are strongly affected by local water quality, but climate

also contributes to species distributions at broad spatial scales (Alahuhta et al. 2011; Netten et al. 2011). Moreover, aquatic plants are often investigated at small spatial scales (Jones et al. 2003; Saarneel et al. 2011) where climate has only a marginal effect on species distributions. Recent works examining the relationship between aquatic plants and ecological gradients at broad scales have focused on all species or life forms (e.g. Mikulyuk et al. 2011; O'Hare et al. 2012; Alahuhta and Heino 2013; Alahuhta et al. 2013), neglecting the keystone species of aquatic plants (cf. Paine 1995). To our knowledge, the contribution of water quality and climate, in addition to spatial determinants of community structure, have not been thoroughly investigated for keystone *Potamogeton* taxa.

This study examines the relative roles of local variables, climate, and geographic location in explaining the distribution of *Potamogeton* taxa in 214 lakes across the states of Minnesota and Wisconsin, USA. The aim was to study how assemblage composition and species richness respond to the three different groups of ecological variables (local, climate, and space), having implications for understanding dispersal mechanisms in landscape or regional contexts. The relationship between sixteen species and the three groups of explanatory variables was also examined to identify species-level trends separate from the whole community. Similar analyses have been used to examine macrophyte assemblage composition in glacial lakes (Mikulyuk et al. 2011; Beck et al. 2013; Alahuhta et al. 2013) but have not been used specifically to evaluate the diverse *Potamogeton* genus. As such, the results presented herein are relevant for understanding drivers of overall assemblage structure of *Potamogeton* taxa, including the most common species in this genus that occur in the upper Midwest United States and regions with comparable climatic and geological characteristics.

Material and methods

Data

Biological surveys of *Potamogeton* taxa from 214 lakes were used, covering the US states of Minnesota and Wisconsin (Fig. 1). The Minnesota and Wisconsin Departments of Natural Resources (MNDNR, WDNR) have collected macrophyte data using the point intercept method beginning in the early 2000s (Madsen 1999). All macrophyte species in each lake were surveyed in a grid design of evenly-spaced points throughout the littoral zone. Species were sampled during the growing season at each point by boat using a grapple that was sunk to the bottom and retrieved to identify species present. Data for each lake included total richness and frequency occurrence of individual species using the total number of survey points for which each *Potamogeton* species was found, scaled by total effort. Sampling effort was set at a point density (i.e., number of points per littoral hectare) that was sufficient to capture all but the most rare species (Mikulyuk et al. 2010; Beck et al. 2014).

Study lakes occurred across a spatial gradient of land use, climate, and morphometry such that the dataset included a variety of lake types. Lakes were situated in four ecoregions of the upper Midwest United States (level III, Omernik 1987), including the Northern Lakes and Forests ecoregion in the north, the North Central Hardwood Forests ecoregion in central areas, and the Northern Glaciated Plains and Western Cornbelt Plains of southern Minnesota (Fig. 1). In general, lake productivity decreases from the south to the north across the ecoregions, whereas overall richness is generally highest in moderately productive or mesotrophic lakes in central regions of each state.

The explanatory variables used to characterize *Potamogeton* distributions were grouped into three categories: local, climate, and spatial. Local variables (water quality and morphometry) included alkalinity concentration (mg/L of CaCO₃), colour (Pt-Co units), lake area (km²), maximum depth (m), perimeter (km), Secchi depth (m) and total phosphorus (mg/L). The water quality variables (alkalinity, colour, Secchi depth, and total phosphorus) were obtained from three sources: MNDNR division of fisheries water quality data (http://www.dnr.state.mn.us/lakefind/), the

STORET database maintained by the United States Environmental Protection Agency (http://www.epa.gov/waterdata/storage-and-retrieval-and-water-quality-exchange), and the Wisconsin Lake Historical Limnological Parameters database (Papes and Vander Zanden 2010). The lake morphometry variables were obtained from the same sources or supplemented with Geographical Information System (GIS) databases (MN: https://gisdata.mn.gov/, WI: http://www.sco.wisc.edu/find-data.html). The climate variables for each lake based on geographic location included annual mean temperature (°C), maximum temperature of the warmest month (°C), minimum temperature of the coldest month (°C), annual precipitation (mm), and lake altitude (m.a.s.l.). The climate variables, with the exception of altitude, were derived from the WorldClim database (Hijmans et al. 2005).

Quantifying spatial variation

Spatial variables describing variation in location were derived from the Cartesian coordinates of geographic centers of each lake (North American Datum 1983). Following similar methods in Alahuhta et al. (2013), Principal Coordinates of Neighbor Matrices (PCNM) analysis was used to deconstruct the lake locations into orthogonally and linearly uncorrelated components (Borcard and Legendre 2002). PCNM is a special form of Moran's Eigenvector Map (MEM) functions that quantifies the spatial autocorrelation among geographic features based strictly on location and proximity between features. Significant spatial variation shown in PCNMs can indicate environmental autocorrelation, dispersal limitation or historical effects on species distributions (Dray et al. 2012). The eigenvectors for PCNM are derived from a Euclidean distance matrix for all locations which is then truncated by the longest distance in the minimum spanning tree linking all sites on the map. Principal Coordinates Analysis of the truncated neighbor matrix produces eigenfunctions for all eigenvectors, which are in turn described by Moran's I values that quantify spatial autocorrelation. Eigenfunctions with positive Moran's I values were retained to describe variation among locations related to positive spatial autocorrelation. In general, this analysis can be

used to describe a range of geographic patterns of spatial variation, with the first few eigenvectors describing large-scale spatial variation and the remaining describing finer-scale variation. The use of PCNM eigenvectors in statistical models provides a means to assess variation among biological communities across the landscape as explained strictly by physical location and relative to additional variables (e.g., climate or local). PCNM analysis was conducted using the PCNM package (Legendre et al. 2013) for the R statistical computing environment (RCT 2015).

Statistical analysis

The variation partitioning procedure provided by the varpart function in the vegan package for R (Oksanen et al. 2015) was used to evaluate effects of local, climate, and spatial variables on the *Potamogeton* data. Partial redundancy analysis (pRDA) was used to evaluate variation in the assemblage composition across lakes (i.e., counts of occurrence of *Potamogeton* species by lake) and partial linear regression (pLR) was used to evaluate variation in total species richness (logtransformed) of *Potamogetons* and frequency occurrence of sixteen individual species. Frequency occurrence data were arcsine-square root-transformed to use linear multivariate methods and Hellinger-transformed to minimize the effect of zeros (i.e., species absence, Legendre and Gallagher 2001). The most important explanatory variables among each category (local, climate, spatial) were selected prior to variation partitioning to reduce model complexity and to avoid type I error. Variable selection followed a standard forward selection technique with stopping criteria if the inclusion of an additional variable in a model produced a probability value less than the selected alpha-level. The significance of a variable was based on a Monte Carlo permutation test of the model residuals following the default procedure (999 permutations, $\alpha = 0.05$) in the forward.sel function in the packfor package for R (Dray 2013). The selected variables in each category were used to create combined models with all categories for variation partitioning. Total variation of the response (assemblage composition, total richness, occurrence of individual species) was

decomposed into separate fractions of: 1) pure local, 2) pure climate, 3) pure spatial, 4) shared local/climate, 5) shared local/spatial, 6) shared climate/spatial, 7) shared all categories, and 8) unexplained (1 – total explained) (Fig. 2, Anderson and Gribble 1998). Detailed information on the variation partitioning process is given in Legendre et al. (2005) and Borcard et al. (2011).

The variation partitioning was based on adjusted R² which provided unbiased estimates of the explained variation of the modelled response variables by individual and combined categories of explanatory variables (Fig. 2, Peres-Neto et al. 2006). The number of explanatory variables is also taken into account in the adjusted R² values, for which reason the different models can be compared to one another (Blanchet et al. 2008). The use of adjusted R² values often results in a decreased percentage of explained variation, that is, it generates a considerable amount of unexplained variation due to the high degree of stochasticity in species distributions (Capers et al. 2010; Alahuhta and Heino 2013).

Results

Potamogeton distribution and lake characteristics

Abundance and distribution of *Potamogetons* varied considerably across the study region (Fig. 1, Table 1). Species richness among lakes was generally highest in the North Central Hardwood Forests and Northern Lakes and Forests Ecoregions. The most abundant species among ecoregions was *P. pectinatus*, except in the Northern Lakes and Forest ecoregion which was dominated by *P. zosteriformis* (Table 1). Lake characteristics also varied with distinct differences among the local, climate, and spatial categories (Table 2). In general, local and climate variables were more strongly related to longitudinal and latitudinal gradients as compared to non-monotonic differences between ecoregions and states.

Spatial groupings of lakes through analysis of geographic coordinates with PCNM produced 38 axes with positive Moran's I scores significantly different than zero ($\alpha = 0.05$). As described below, selection of spatial axes during initial model development selected some axes more often

than others (Fig. 3). Broad-scale variation in spatial groupings described by the first and second axes characterized variation in lake locations in central Minnesota, with the first axis describing a grouping in the Northern Lakes and Forests ecoregion and the second describing a grouping in the North Central Hardwood Forests ecoregion (Fig. 3). The fifth axis included central groupings similar to the first and second axes but also included a Wisconsin grouping in the north-central region of the state. Spatial variation described by the ninth axis was less clear and appeared to describe spatial groupings not captured by the other three axes.

Assemblage composition and species richness

Variation partitioning with pRDA and pLR for assemblage composition and total Potamogeton richness, respectively, indicated that local, climate, and spatial variables explained a total of 33% and 47% of the variation in each (Fig. 4, Table 3). For assemblage composition, initial variable selection by category in order of decreasing importance (Table 4, based on adjusted R² for each variable, p < 0.05 for all) was alkalinity, maximum depth, total phosphorus, Secchi depth, lake color, and lake area for local variables, maximum temperature, precipitation, lake altitude, mean temperature, and minimum temperature for climate variables, and twenty-two spatial axes from PCNM (axes one and two were most important). Variation of assemblage composition explained by category (Fig. 4, Table 3) indicated that the pure effect of spatial variables (6.6%) exceeded the pure effects of climate (0.6%) and local variables (3.8%). The joint effects of variable categories suggested that climate and space had the largest shared variation (6.2%), whereas variation in assemblage composition explained by all three categories was 11.8%. For total richness, significant local variables (p < 0.0001 for all) were maximum lake depth and total phosphorus (Table 4). Linear models for the parameter estimates for richness indicated a positive association with lake depth and a negative association with total phosphorus. For the climate variables, a negative association with minimum temperature (p < 0.0001) and a positive association with precipitation (p< 0.05) were observed with richness. Twelve of the fifty-eight spatial variables were selected for

richness. Variation of total richness explained by each category indicated that the pure effects of spatial variables (14.9%) were largest, although pure local effects explained a comparable amount of the variation (11.7%) (Fig. 4, Table 3). Likewise, the joint effects (two-way and all three categories) were smaller compared to the pure effect of local or spatial variables, although shared variation between all three categories was 11.3%.

Species level

The combined dataset for Minnesota and Wisconsin included 25 *Potamogeton* species, of which only sixteen were sufficiently abundant to create regression models for variation partitioning. Species present but not modelled included *P. alpinus*, *P. bicupulatus*, *P. filiformis*, *P. nodosus*, *P. oakesianus*, *P. obtusifolius*, and *P. vaseyi* (n = 1-7 lakes). For the remaining species, total explained variance ranged from 67.9% (*P. pectinatus*) to 12.2% (*P. strictifolius*) with an average of 31.7% for all species (Fig. 4, Table 3). For the pure effects of each variable category, local effects ranged from 14.0% (*P. gramineus*) to zero explained variance (numerous spp.), climate effects ranged from 2.8% (*P. pusillus*) to zero (numerous spp.), and spatial effects ranged from 20.0% (*P. crispus*) to 0.4% (*P. strictifolius*). Within the joint effects, explained variance that was shared between categories was generally lowest for local plus climate effects (average 0.5% for all species), whereas climate plus space and local plus space had similar averages (both ~4%). Average shared explained variance among all three categories was relatively large (9.9%), suggesting species response to shared effects was better explained by covariation among all explanatory variables rather than shared effects between pairwise categories.

An evaluation of variables within each category provided additional information on drivers of species occurrence (Table 4, Figs. 5-7). For local variables, alkalinity, lake color, maximum depth, and total phosphorus were most commonly selected from forward selection (Table 4). Species that included these variables typically showed negative associations with alkalinity and total phosphorus and positive associations with maximum lake depth. These trends were reversed

for some species, most notably for *P. crispus* and *P. pectinatus*. For climate variables, lake altitude and precipitation were most commonly selected. Species associations with climate variables did not show any regular patterns with the exception of temperature variables, such that species were most often negatively associated with increases in temperature. However, *P. crispus* and *P. pectinatus* had positive associations with maximum temperature. Selection of spatial variables typically did not include more than a few axes, suggesting variation by location was explained primarily by the first few spatial axes. However, some species had models with a relatively high number of axes, including *P. crispus*, *P. friesii*, *P. pectinatus*, *P. praelongus*, and *P. robbinsii*.

Discussion

Assemblage composition and species richness

Of the pure fractions of variation, assemblage composition and species richness of *Potamogetons* were most strongly explained by spatial variables. This suggests either a minimal effect of geographically structured environmental variables or historical limitations on species distributions. The models included many important environmental variables that structure macrophyte distributions (Vestergaard and Sand-Jensen 2000; Jones et al. 2003; Mikulyuk et al. 2011; Alahuhta et al. 2013; Beck et al. 2013) and the notable difference in the effects of local and climate variables relative to spatial variables provided evidence that geographic variation in environmental variables was unimportant. Additionally, the current range of *Potamogetons* in North America was likely established immediately after deglaciation (Sawada et al. 2003) and phylogenetic analyses have indicated that North America may be the ancestral geographic area (Lindqvist et al. 2006). This suggests that historical effects are not driving the strong spatial patterns of assemblage composition and species richness. Further, aquatic macrophyte communities may be dispersal-limited in unstable freshwater ecosystems (e.g., floodplain lakes, Padial et al. 2009; Padial et al. 2014), although environmental instability may not be relevant for glacial lakes that are geologically stable (O'Hare et al. 2012; Alahuhta et al. 2015). Mikulyuk et al. (2011) found

that spatial variables contributed a relatively high amount of explained variation for macrophyte communities in Wisconsin. They suggested that dispersal limitation related to strong latitudinal gradients may explain the strong effects of spatial variables. Although, many *Potamogetons* favour more nutrient- and alkaline-rich waters, the geographic centers were located in the middle of Minnesota and northern Wisconsin where lakes are more mesotrophic (Fig. 1). Thus, *Potamogeton* assemblage composition and species richness may well be dispersal limited by eutrophication in the southern region of the study area.

Pure effects of local environmental variables also explained a significant but smaller amount of variation for assemblage composition and species richness. All local variables were significantly related to *Potamogeton* assemblage composition, whereas only maximum depth and total phosphorus were correlated with species richness. The positive association between species richness and maximum depth is likely related to habitat availability, as habitat heterogeneity increases with an increasing depth gradient (Tolonen et al. 2001; Alahuhta et al. 2013). Increasing lake depth has also been associated with increases in biotic integrity of macrophytes in glacial lakes (Beck et al. 2010). Increasing lake depth and species richness could also be linked to effects of wave exposure. Macrophytes in deeper lakes are less susceptible to uprooting from wave action during high wind events (Riis and Hawes 2003). Turbidity in shallow lakes is also influenced by wind, which could indirectly limit macrophyte growth by light scattering. The negative influence of total phosphorus on species richness further emphasizes that the nutrient status of many lakes exceeds levels for sustaining *Potamogeton* species. Similar results related to latitudinal gradients in water quality have been found for species richness of all submerged macrophytes in Midwest glacial lakes of the United States (Beck et al. 2014; Alahuhta 2015). Interestingly, neither alkalinity nor Secchi depth were related to *Potamogeton* species richness, although both of these variables can affect growth patterns of submerged vascular plants through carbon and light availability, respectively (Chambers and Kalff 1985; Madsen et al. 1996).

Climate variables were weakly correlated to the distribution of *Potamogeton* communities. Previous studies have similarly reported that local water quality and morphometric conditions are often more important than climate for aquatic macrophytes at spatial scales similar to our analysis (Santamaría 2002; Chappuis et al. 2014; Alahuhta 2015). For assemblage composition, all individual climate variables were significant for *Potamogetons*, whereas species richness was only positively correlated with annual precipitation and negatively with minimum temperature of the coldest month. Higher annual precipitation has been correlated to higher nutrients and suspended solids that are leached from terrestrial sources during intense rainfalls (Cobbaert et al. 2014). However, this contradicts our results that showed a negative association between species richness and total phosphorous. Annual precipitation can function as a proxy for water-induced dispersal as plant propagules are more easily transported through stream networks during high flows (Riis 2008), which may explain the positive association observed from the model. The increase in Potamogeton species richness along decreasing winter temperature contradicts the commonly known influence of temperature on plant species richness (e.g., Pip 1989). Harsh winter conditions could restrict macrophyte growth through thick ice cover limiting the availability of carbon, oxygen, and light, freezing bottom sediments, or increasing ice erosion (Lind et al. 2014). However, Alahuhta (2015) and Johnson et al. (2010) have similarly described a reversed latitudinal gradient in species richness of all macrophyte taxa in the Midwest United States. Cold climate may be an important environmental filter for *Potamogetons* such that only tolerant species can occur at northern latitudes with lower minimum temperatures. A historical latitudinal gradient has likely existed for wetland plant species in the Great Lakes region, although anthropogenic activities have steepened the gradient (Johnson et al. 2010). Our results provide additional support that the latitudinal gradient is partially based on climatic differences, whereas land-use changes along this gradient have further affected water quality in the southern parts of the states. Water quality may be

a primary filter limiting species from the regional pool such that the majority of species cannot survive in the eutrophic lakes at southern latitudes.

Finally, the shared fractions of variation, either as joint effects or shared among all three categories of variables, explained a substantial amount of variation in assemblage composition and total species richness. For assemblage composition, the joint effects of climate and space were equally high compared to the highest pure effects of individual categories. Similarly, the joint effects of climate and space for total richness were comparable to those for assemblage composition, although the pure effects of space and local variables were higher compared to assemblage composition. This finding indicates that climate variables are geographically structured, related to the decreasing gradient with latitude in climate (Beck et al. 2013; Alahuhta 2015). The shared effects of all three variable groups were also high for assemblage composition and species richness, indicating that their individual influences cannot be statistically distinguished. Similar results were found by Beck et al. (2013) where a majority of variation of a macrophyte-based index of biotic integrity for Minnesota was explained by shared effects of environmental and anthropogenic variables, rather than the pure effects of each. These results emphasize challenges in variance partitioning when variables are correlated across spatial gradients.

Species level

Similar conclusions about the effects of local, climate, and spatial variables for assemblage composition and total richness can be generalized for individual species. Overall, the pure effects of spatial and local variables were much larger than climate effects and the combined effects of variable categories were generally larger than the pure effects for any given species. The latter conclusion was particularly true for effects shared between all categories, local plus space, and climate plus space. However, some differences between species were observed that potentially explains geographic variation in the relative distributions of each. For example, the total explained variation of separate models for *P. pectinatus*, *P. pusillus*, and *P. crispus* exceeded the total

explained variation for assemblage composition and total richness. The geographic centers of each species in relation to the spatial gradients among the explanatory variables provided a potential explanation for the relatively high amount of explained variation of the models. Both P. crispus and P. pectinatus were centered more closely than the other species to the southern area of Minnesota where lakes are more nutrient-rich, alkaline, and warmer. For *P. pectinatus*, local variables were also geographically patterned, likely in the southern lakes. Both species are commonly regarded as tolerant of eutrophic conditions (Beck et al. 2010) and relatively intolerant of harsh climate conditions. Valley and Heiskary (2012) provide evidence that P. crispus in Minnesota may be lightlimited during harsh winters by thicker snow depth on frozen lakes. Geographic climate variation and the positive association with maximum temperature suggests that P. crispus is likely restricted in the northern lakes by climate, although future expansion north may be mediated by warmer winter conditions associated with climate change. However, the largest pure effect for P. crispus was attributed to spatial variables, suggesting that lake location or proximity was an important factor explaining distribution. Interestingly, the second spatial axis from PCNM had the strongest correlation to the distribution of *P. crispus*. This axis explained spatial correlation among lakes in the North Central Hardwood Forests ecoregion of central Minnesota, which is also near a large metropolitan area (Fig. 3). Species transport between lakes with heavy recreational use in urban areas is a well-known vector of exotic species invasion (e.g., Miro and Ventura 2013). As such, the importance of spatial groupings among lakes for *P. crispus* provides further support of physical transport as a means of dispersal in the Midwest United States.

Other species also had disproportionately large amounts of variation explained by specific categories of the explanatory variables. As noted above, *P. pusillus* had a large amount of total explained variation compared to the remaining species. Although most of this variation was from shared effects, a large percentage of pure variation was explained by spatial variables. The geographic center of *P. pusillus* was located in northern Wisconsin and was well-described by the

ninth spatial axis that characterized a lake grouping in that region (Figs. 3 and 7). Additionally, the influence of climate (pure fraction and geographically-structured climate effect) was large for P. pusillus relative to the remaining species. The climate variables of lake altitude and precipitation were positively correlated with the distribution of *P. pusillus*, which suggests that the species is more commonly found in lakes that are higher in the hydrologic network and that receive more precipitation. These lakes likely have variable water levels as they fill from precipitation and drain quicker than lakes lower in the watershed. Some studies have suggested that P. pusillus is an early colonizer of habitats affected by hydrological alteration (Boedeltje et al. 2001; Van Geest et al. 2005), which may explain the strong association of the species with the climate variables. Local variables also explained a disproportionate amount of pure effects for some species, particularly P. illinoensis and P. gramineus. Both species were correlated with lake depth and lower total phosphorus, suggesting relative intolerance of eutrophic conditions. Additionally, P. illinoensis is more common in hardwater lakes, which was supported by geographically structured local variables (local/space) and especially a strong correlation with alkalinity. Both species are also morphologically similar suggesting an adaptation for similar habitats based on phenotypic characteristics.

Conclusions

This study provided a unique decomposition of factors that influence the distribution of *Potamogeton* species in glacial lakes of the Midwest United States. To our knowledge, no studies have evaluated the contributing factors of local, spatial, and climate variables on the distribution of this keystone genus. Variation partitioning analyses revealed that assemblage composition and total species richness were best explained by spatial groupings of the study lakes, particularly lake groups along a strong latitudinal gradient. Further evaluation suggested that spatial variables were descriptive of dispersal limitations of *Potamogeton* species related to eutrophication such that most species in the analysis were unable to colonize high-nutrient lakes. An additional, but minor,

confounding effect of climate described a temperature limitation along the latitudinal gradient such that most species were unable to colonize northern regions of the study area that had lower minimum temperatures. Models for individual species generally supported the results from the community models, with the exception of some species that were disproportionately described by specific categories of variables. For example, the invasive species *P. crispus* was strongly related to eutrophication variables but also showed a strong relationship with spatial variables, suggesting a vector for invasion related to recreational use of lakes in urban areas.

Overall, this analysis provides an argument that the management and conservation of this important keystone genus could focus on drivers of assemblage composition that are spatially aggregated across the landscape, in addition to traditional management efforts that focus on local characteristics related to eutrophication. Management efforts for individual species could be similar but dependent on lake characteristics and their variation across the landscape relative to the species of interest. Finally, considering the integration of *Potamogeton* species within the larger community of macrophytes and other biota may also be important for developing a landscape-level understanding of factors that drive variation in the entire community, having implications for regional efforts of lake management and conservation.

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Figure captions

- **Fig. 1** *Potamogeton* species richness (top) for each study lake (n = 214) and geographic centers (bottom) for each species in Minnesota and Wisconsin. County boundaries (lines) and shading for each level III ecoregion (Omernik 1987) are also shown. Ecoregions with study lakes included North Central Hardwood Forests (NCHF), Northern Glaciated Plains (NGP), Northern Lakes and Forests (NLF), and Western Cornbelt Plains (WCBP). Geographic centers were estimated as the average latitude/longitude weighted by species occurrence. The richness legend applies only to the top plot. PA: *P. amplifolius*, PC: *P. crispus*, PE: *P. epihydrus*, PF: *P. foliosus*, PG: *P. gramineus*, PI: *P. illinoensis*, PN: *P. natans*, POP: *P. praelongus*, POPU: *P. pusillus*, POR: *P. richardsonii*, POS: *P. spirillus*, PP: *P. pectinatus*, PR: *P. robbinsii*, PS: *P. strictifolius*, and PZ: *P. zosteriformis*
- **Fig. 2** Conceptual representation of variation partitioning analyses used to estimate explained variance of *Potamogeton* assemblage composition, total richness, and frequency occurrence of individual species. Categories of explanatory variables for each lake included local, climate, and spatial information. Partial redundancy analysis and partial linear regression models were used to explain total variation (area covered by all circles), pure variation (non-overlapped local, climate, or spatial areas), shared variation (overlapping areas between categories or all three), and unexplained (total area circle area). Additional details are in Legendre et al. (2005) and Borcard et al. (2011)
- **Fig. 3** Scores for selected axes from principal components of neighbouring matrices. The axes shown were the four most frequently retained during variable selection of models to explain *Potamogeton* distribution. Points for each lake are scaled relative to axis scores to show spatial autocorrelation from geographic location. The Moran's eigenvectors (top) are a combined measure of spatial autocorrelation described by each axis. See Fig. 1 for the ecoregion labels
- **Fig. 4** Fractions of variation explained by local, climate, and spatial variables for *Potamegeton* assemblage composition, richness, and individual species. The top plot shows total explained variance (sum of pure and shared effects), the middle shows pure variation explained by each variable category, and the bottom shows shared fractions of variation for different combinations of the variable categories. See Fig. 2 for a conceptual representation of the fractions of variation
- **Fig. 5** Redundancy analysis biplots for *Potamogeton* spp. relative to local lake characteristics (see Table 2). The plot was created using all variables to explain relationships affecting species distribution. See Fig. 1 for species abbreviations
- **Fig. 6** Redundancy analysis biplots for *Potamogeton* spp. relative to climate characteristics (see Table 2). The plot was created using all variables to explain relationships affecting species distribution. See Fig. 1 for species abbreviations
- **Fig. 7:** Redundancy analysis biplots for *Potamogeton* spp. relative to spatial variables from principal coordinates of neighbouring matrices (PCNM). The PCNM axes used in the figure were the top ten most frequent axes that were retained after variable selection for individual species models. See Fig. 1 for species abbreviations

Table 1 Summary of *Potamogeton* distribution by ecoregion (Fig. 1, Omernik 1987). Species shown are those with variation partitioning models in Fig. 4. The mean, minimum, and maximum are based on frequency occurrence values by lake. Lake counts in each ecoregion are in parentheses. NCHF: Northern Lakes and Forests Ecoregion, NGP: Northern Glaciated Plains, NLF: Northern Lakes and Forests, WCBP: Western Cornbelt Plains

	NCHF (78)		NGP (3)		NLF	(122)	WCBP (11)		
	n	mean (min -	n	mean (min -	n	mean (min -	n	mean (min -	
		max)		max)		max)		max)	
P. amplifolius	29	0.12 (0-0.57)	0	0 (0-0)	100	0.29 (0-0.78)	1	0.01 (0-0.15)	
P. crispus	50	0.31 (0-0.93)	0	0 (0-0)	26	0.08 (0-0.67)	8	0.41 (0-0.86)	
P. epihydrus	2	0.01 (0-0.34)	0	0 (0-0)	27	0.07 (0-0.64)	0	0 (0-0)	
P. foliosus	3	0.01 (0-0.41)	1	0.14 (0-0.42)	17	0.04 (0-0.48)	0	0 (0-0)	
P. friesii	22	0.09 (0-0.64)	1	0.2 (0-0.59)	41	0.1 (0-0.6)	3	0.1 (0-0.45)	
P. gramineus	19	0.08 (0-0.64)	0	0 (0-0)	72	0.18 (0-0.55)	0	0 (0-0)	
P. illinoensis	36	0.16 (0-0.71)	0	0 (0-0)	54	0.14 (0-0.69)	1	0.03 (0-0.32)	
P. natans	24	0.08 (0-0.4)	0	0 (0-0)	61	0.14 (0-0.68)	1	0.02 (0-0.25)	
P. pectinatus	66	0.35 (0-0.84)	2	0.42 (0-0.64)	50	0.11 (0-0.4)	11	0.6 (0.34-1)	
P. praelongus	37	0.16 (0-0.71)	0	0 (0-0)	80	0.21 (0-0.7)	1	0.03 (0-0.32)	
P. pusillus	16	0.08 (0-0.66)	0	0 (0-0)	48	0.17 (0-0.92)	0	0 (0-0)	
P. richardsoni	40	0.18 (0-0.51)	2	0.37 (0-0.61)	82	0.21 (0-0.56)	4	0.18 (0-0.64)	
P. robbinsii	15	0.08 (0-0.69)	0	0 (0-0)	59	0.2 (0-1)	0	0 (0-0)	
P. spirillus	1	0 (0-0.27)	0	0 (0-0)	16	0.04 (0-0.62)	0	0 (0-0)	
P. strictifolius	5	0.02 (0-0.51)	0	0 (0-0)	23	0.06 (0-0.85)	0	0 (0-0)	
P. zosteriformis	51	0.29 (0-0.75)	1	0.07 (0-0.21)	96	0.34 (0-0.94)	5	0.21 (0-0.61)	

Table 2 Summary of local and climate variables used to evaluate distributions of *Potamogetons*

	Mean	Minimum	Maximum	SD
Local variables				
Alkalinity (mg L ⁻¹ CaCO ₃)	100.16	5.08	282.50	60.57
Area (km ²)	2.84	0.08	22.50	3.78
Color (Pt-Co units)	24.53	0.00	160.00	22.24
Maximum depth (m)	12.33	1.52	63.40	9.57
Perimeter (km)	8.46	0.00	57.49	10.05
Secchi depth (m)	2.90	0.25	40.88	3.14
Total phosphorus (mg L ⁻¹)	0.07	0.01	0.83	0.09
Climate variables				
Altitude (m.a.s.l.)	376.11	180.00	593.00	76.24
Annual mean temperature (°C)	5.01	2.62	7.42	1.20
Annual precipitation (mm)	18.89	12.00	31.00	3.80
Maximum temperature of the warmest month (°C)	27.14	22.50	29.30	1.30
Minimum temperature of the coldest month (°C)	-18.92	-23.40	-12.60	1.89

Table 3 Explained variance of assemblage composition, total species richness, and individual *Potamogeton* species among local, climate, and space variables. Values are adjusted R² (PeresNeto et al. 2006) from partial redundancy analysis and partial least squares

				Local +	Climate	Local +		
	Local	Climate	Space	Climate	+ Space	Space	All	Total
Assemb. comp.	3.83	0.63	6.63	0.33	6.17	3.38	11.84	32.81
Richness	11.66	-0.47	14.94	0.31	5.38	3.84	11.32	46.99
P. amplifolius	3.83	1.43	3.15	2.81	1.20	2.75	21.06	36.23
P. crispus	2.34	-0.09	20.02	0.69	9.30	-0.84	15.93	47.35
P. epihydrus	1.41	2.48	3.04	0.79	1.49	5.06	13.91	28.18
P. foliosus	0.08	2.09	7.69	-0.29	1.75	-0.50	2.26	13.09
P. friesii	0.98	-0.16	9.22	0.03	5.87	-1.16	1.48	16.27
P. gramineus	13.98	0.35	7.59	1.87	3.95	2.25	5.92	35.91
P. illinoensis	11.45	0.56	10.34	-0.84	4.30	10.87	6.20	42.88
P. natans	4.03	-0.33	7.28	-0.04	4.63	1.38	5.41	22.37
P. pectinatus	5.86	-0.16	13.00	0.17	0.67	19.11	29.20	67.86
P. praelongus	1.90	-0.34	16.11	-0.01	4.68	5.82	2.89	31.05
P. pusillus	-0.17	2.78	8.61	0.18	15.80	3.54	19.92	50.66
P. richardsoni	4.61	1.04	6.19	1.77	0.53	0.67	0.51	15.31
P. robbinsii	7.55	0.38	13.54	-0.56	0.71	4.64	16.76	43.02
P. spirillus	2.28	0.24	6.39	0.22	6.22	2.95	8.68	26.99
P. strictifolius	-0.29	0.73	0.42	0.12	5.18	0.37	5.64	12.17
P. zosteriformis	0.28	0.39	4.28	0.45	10.01	0.34	2.76	18.51

Table 4 Significant local, climate, and spatial variables used for variation partitioning. Values are by category from individual models created with forward variable selection. Direction of the effect is indicated by + and - symbols from the selected model, excluding results for assemblage composition that modelled multiple species with redundancy analysis. For spatial variables, only the number of selected significant eigenvectors is shown. Explained variance (%) indicates how much the model explains the dependent variable based on adjusted R^2 and does not consider joint effects with the remaining categories (see Table 3). ***: p < 0.0001, *: p < 0.005, ns: p > 0.05

	Local							Climate						Space	
	alk	area	color	depth	secchi	tp	%	alt	prec	tmax	tmean	tmin	%	n	%
Assemb. comp.	**	*	*	**	*	**	20.08	**	**	**	**	*	19.38	22	28.71
Richness				***+		***_	27.14		*+			***_	16.54	12	36.35
P. amplifolius	***_					**_	30.44		**+	ns+	*_		26.50	8	29.34
P. crispus	***+			*_		*+	18.12			***+			25.83	12	44.41
P. epihydrus	***_		*+				21.17	*+	**+	*_			18.67	6	22.32
P. foliosus	*_						1.56	**+					5.82	3	11.21
P. friesii	ns+						1.33		***_				7.22	11	20.32
P. gramineus				**+	***+	*_	24.02	*_			***_		12.09	5	19.71
P. illinoensis	***+		*_	**+		***_	27.69	***_			*_		10.22	6	30.72
P. natans				**+		*_	10.78					***_	9.67	7	18.70
P. pectinatus	***+	*_				*+	54.34			***+			29.88	21	61.98
P. praelongus			*_			***_	10.60					***_	7.22	10	30.55
P. pusillus	***_						23.47	***+	***+				36.40	8	47.87
P. richardsoni		***+					7.56		*_				3.85	4	10.30
P. robbinsii	***_		*_	*+			28.39			***_			17.30	11	35.64
P. spirillus	***_		*+				14.14	***+	**+				15.37	5	23.00
P. strictifolius	**_						5.84	*+	***+				11.67	3	13.16
P. zosteriformis				*+			3.83	*_				***_	13.62	5	18.62



















