Recognition of key regions for restoration of phytoplankton communities in the Huai River basin, China

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1. Introduction

The world’s ecosystems (both terrestrial and aquatic) are capital assets. They can yield numerous vital services, including the production of goods, life support processes, and life-filling conditions when properly managed. Ecosystem services are essential to human existence and operate on such a grand scale, and in such intricate and little-explored ways, that most could not be replaced by technology. Unfortunately, escalating impacts of human activities on ecosystems imperil their delivery. Ecosystems are undergoing rapid degradation and depletion (Daily, 1999; Daily et al., 2000) and are increasingly threatened by human-induced habitat loss (Kagalou et al., 2010). With economic development and population increase over the past decades, large quantities of pollutants have been discharged into rivers in China. This has resulted in degradation of water quality and greatly impaired aquatic ecosystems, which, in turn, has severely hindered the sustainable development of the economy and society. This is especially true for the Huai River, China.

The Huai River, located between the Yangtze River and the Yellow River, forms a geographical divide between northern and southern China. The Huai River basin (HRB) is one of the main grain-producing areas of China. Its population (0.165 billion by the year 2000) surpasses all other large basins in China (Zhao et al., 2010). Over the last half century, it has been severely affected by human activities, especially construction of dams and weirs and discharge of pollutants. Because of serious flood disasters and flood control requirements in the Huai River, around 11,000 dams and sluices had been built by the year 2000. The number of such structures on this river accounts for approximately half of those in China and a quarter of those in the world (Liu et al., 2011). These structures have brought tremendous economic benefit through flood control, increased irrigation, and power
generation in the basin. However, major counterarguments have been raised for many years regarding their detrimental impacts on the environment (Wang and Xia, 2010; Zhao et al., 2010). Dams and sluices usually lead to hydrologic “fragmentation”, which greatly harms aquatic ecosystems of the HRB. Water quality in more than 83% of rivers cannot reach the national criteria (GB3838–2002) and the quality in this basin is the worst in the nation's top seven basins (based on Chinese Environment Bulletin in 2005) (Zhang et al., 2010). The river has a history of disastrous pollution events. One severe water pollution event happened in 1994 because of major floods, made worse by inappropriate operation of sluices. Affected waterworks had to stop supplying water for 54 days and 1.5 million people suffered from a shortage of drinking water. This caused economic losses of at least US$200 million (Xia et al., 2011). Long-term mismanagement of dams and sluices for water use, plus excessive pollution discharge, has resulted in ecosystems in many middle and lower river reaches that are seriously degraded and extremely unstable (Zhao et al., 2008). Consequently there is a need to re-construct or restore stable, healthy aquatic ecosystems. The top priority should be given to the organisms that underpin the food webs of aquatic ecosystems – phytoplankton communities in the HRB.

In aquatic ecosystems, phytoplankton is the major microbial biomass. Light energy conversion and related synthesis of carbon compounds is carried out by three major primary producers – higher plants (macrophytes), phytoplankton, and photosynthetic bacteria. Phytoplankton are the main microorganisms involved in this process (Sigee, 2005). They play a major role in aquatic ecosystems as their biological activity affects the biogeochemical cycles of a number of macro and micronutrients (carbon, silicon, sulfur, nitrogen, iron, etc.) (Falkowski, 1994; Falkowski et al., 1998). Phytoplankton are also the principal primary producer of freshwater food webs and mainly depends on light energy and nutrients. However, too many nutrients often accelerate the growth of phytoplankton and an overpopulation of phytoplankton is extremely harmful to the local aquatic-related economy. For example, toxic algal blooms worsen water quality and greatly impair local fisheries. The abundance and biomass of phytoplankton should therefore be controlled within a moderate range. An index of “dominance” (Zhao et al., 2011), which denotes the importance or contribution of a species can be used to assess community structure and used to manage aquatic ecosystem health. To control the “dominance” of a species, understanding of the connection between its abundance and its habitat indices, especially indices of water quality are necessary.

The ecological niche can establish connections between phytoplankton species and indices of water pollution. It is one of the most important concepts in the exploration of biological communities’ structure and development, biodiversity, association of species with a particular environment, conservation planning and decision making (Pearce and Lindenmayer, 1998; Ferrier, 2002; Wiley et al., 2003). Hutchinson (1957) formalized the niche as an n-dimensional hyper volume whose axes are critical physical and environmental factors determining the existence of a species – this concept of niche as a function of measurable factors has provided a foundation for many theoretical and field studies (Smith, 1982).

![Fig. 1. Phytoplankton sampling in the HRB (modified from Zhao et al. (2011)), the numbers on the map indicate the sampling sites.](image-url)
Studies on ecological niche have mainly focused on niche breadth and niche overlap (Thompson and Gaston, 1999; Jelke et al., 2000; Brändle et al., 2002). Ecological niche has been widely applied in research on habitat selection, species conservation, spatial distribution and temporal dynamics, temporal and spatial niche-partitioning, species delimitation, exotic species invasion and community succession, etc. (McNysst, 2005; Domínguez-Domínguez et al., 2006; Chen et al., 2007; Irfan-Ullah et al., 2007; Peterson et al., 2007; Raxworthy et al., 2007; Solano and Feria, 2007; Basille et al., 2008; Foulon et al., 2008; Friberg et al., 2008; Peterson and Nakazawa, 2008; Quero et al., 2008; Thorn et al., 2009; Waltari and Guralnick, 2009). The protection or restoration of whole ecosystems often represents the most effective way to sustain genetic, population, and species diversity (Vitousek et al., 1997). Understanding the consequences of biodiversity changes on ecosystem functioning is becoming increasingly critical in view of the profound influence of human activity on natural ecosystems and the goods and services humans receive from them (Vitousek et al., 1997; Daily et al., 2000; Giller et al., 2004). To achieve restoration of aquatic ecosystems across a very large area, selection of appropriate key regions or priority areas is essential.

Most studies in aquatic systems have used plants, invertebrates, fish and birds as indicators (Altaba, 1990; Crandall, 1998; Posadas and Crisci, 2001; Pérez-Losada et al., 2002; Turpie et al., 2002; Sánchez-Fernández et al., 2004; Abellán et al., 2005). Few studies focused on the phytoplankton, and furthermore, few considered many factors of dominance, biodiversity, water quality and ecological niche.

In view of the major role of phytoplankton in aquatic ecosystems, the objective of this paper is to identify the key regions for future restoration of degenerated phytoplankton communities resulting from severe water pollution. This study was based on assessments of dominance, biodiversity and ecological niche along a gradient of water quality indices.

### 2. Methodology

#### 2.1. The study area

The Huai River is the sixth largest river in China. It is located between the Yangtze River and the Yellow River of China (Wang and Xia, 2010). The area of the Huai River basin (HRB: 30°55′–36°36′N, 111°55′–121°25′E) covers 27,000 km². The HRB lies at China’s transition between the northern climate and southern climate (Gao et al., 2010). It is the most densely inhabited river basin and the main grain-producing area of China. In 2005, the total population and grain yield accounted for 13.1% and 16.1%, respectively, of the national total (Xia et al., 2011). The population density surpasses all other large basins in China (Zhao et al., 2010).

The HRB can be divided into eight regions: Main Stream (R1), Hongru River (R2), Shaying River (R3), Guo River (R4), Baohui River (R5), Yishu River (R6), Along East line of South-North Water Transfer Project (R7), Southern Mountain Area (R8), as shown in Fig. 1.

#### 2.2. Methods

To explore the impacts of water pollution on phytoplankton communities and therefore identify key regions for ecological restoration in the HRB, we sampled phytoplankton as well as water chemistry at 71 typical sites (Fig. 1). We sampled every day during a low-water period in the hot wet season (from July 10th to July 20th, 2008). This low-water period is the best for exploration of relationships between phytoplankton and water quality because during this period phytoplankton flourish and the impacts of dam on water quality and water ecosystems are greatly reduced since most water sluices are kept open.

**2.2.1. Water chemistry**

We measured in situ indices of water chemistry indices: water temperature: WT; pH; and dissolved oxygen: DO by using a portable HACH PC101. Twenty-one additional indices were tested in the laboratory. Water samples were sent to laboratory within 24 h. Various instruments were used to analyze the additional indices. Among these instruments, Spectrophotometer (DR5000) was used to measure Ammonia nitrogen (NH₃–N), total phosphorus (TP), total nitrogen (TN) and hexavalent chromium; Atomic Absorption Spectrophotometer (Thermo M6) was used for test of copper (Cu), zinc (Zn), cadmium (Cd) and lead (Pb); Ion Chromatograph (DIONEX-600) was employed to measure sulfate, fluoride, chloride and nitrate; Automatic Flow Injection Analyzer (KALAR SAN**) was used to measure cyanide, volatile phenol, anionic detergent. Of the initial 21 water chemistry indices selected for analyses the concentrations of many of them were at or below the limits of detection at 69.0–100% of the sampling sites. Consequently we selected nine main factors (WT; pH; DO; NH₃–N; permanganate index; CODₘₜ; chemical oxygen demand; CODₚ; TP; TN; Fluoride) as the water chemistry indicators of the Huai River (Table 1).

**Table 1**

<table>
<thead>
<tr>
<th>No.</th>
<th>Environmental factor</th>
<th>Range</th>
<th>Mean ± SD</th>
<th>Mode</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Water temperature (WT)</td>
<td>27.3–33.4</td>
<td>29.9 ± 1.34</td>
<td>30.6 (8.82%)</td>
</tr>
<tr>
<td>2</td>
<td>pH</td>
<td>6.08–8.09</td>
<td>7.33 ± 0.39</td>
<td>7.5 (5.63%)</td>
</tr>
<tr>
<td>3</td>
<td>Dissolved oxygen (DO)</td>
<td>1.20–11.8</td>
<td>5.76 ± 2.37</td>
<td>7.8 (11.1%)</td>
</tr>
<tr>
<td>4</td>
<td>Ammonia nitrogen (NH₃–N)</td>
<td>0.14–12.4</td>
<td>1.13 ± 2.39</td>
<td>0.27 (7.04%)</td>
</tr>
<tr>
<td>5</td>
<td>Permanganate index (CODₘₜ)</td>
<td>1.96–18.7</td>
<td>5.58 ± 2.72</td>
<td>4.40 (7.04%)</td>
</tr>
<tr>
<td>6</td>
<td>Chemical oxygen demand (CODₚ)</td>
<td>11.1–54.7</td>
<td>22.6 ± 10.1</td>
<td>13.5 (6.45%)</td>
</tr>
<tr>
<td>7</td>
<td>Total phosphorus (TP)</td>
<td>0.017–1.742</td>
<td>0.16 ± 0.26</td>
<td>0.03 (8.45%)</td>
</tr>
<tr>
<td>8</td>
<td>Total nitrogen (TN)</td>
<td>0.32–15.4</td>
<td>2.66 ± 2.97</td>
<td>3.07 (4.23%)</td>
</tr>
<tr>
<td>9</td>
<td>Fluoride</td>
<td>0.5–0.9</td>
<td>0.65 ± 0.17</td>
<td>0.50 (52.1%)</td>
</tr>
</tbody>
</table>

Unit of water temperature (WT) is degrees Celsius, pH has no unit and the rest are all in mg L⁻¹. The mode value in the table is expressed as “Mode (percentage of number to total number)”.

**2.2.2. Phytoplankton sampling and taxa determination**

A 1000 mL-capacity organic glass bottle was used to sample water from 0 to 2 m below the water surface. As quickly as possible, 1.5% concentration Lugol’s solution was added to the bottle. In the laboratory, a 24-h sedimentation method was used to concentrate the phytoplankton sample to 30 mL. A 0.1 mL sub-sample was taken from the 30 mL concentrated sample and loaded into a 0.1 mL plankton counting chamber. Finally, the phytoplankton...
were counted using Utermöhl’s inverted plankton microscope. The biomass was converted from biovolume assuming a specific gravity of 1.0. To determine individual biovolume, individual size (length, height and breadth, or diameter) of a species was measured with the plankton microscope. Average size of at least 50 individuals was used to calculate average biovolume of a species (SL167-961).

2.2.3. Dominance assessment
Abundance and biomass are fundamental indices for biological monitoring. The two indices often rank differently, which makes it hard to objectively assess the dominance or importance of a species in a community (Zhao et al., 2011). To overcome this, Zhao et al. (2011) combined them into one index by using the following equation:

\[ I_{\text{importance}} = \omega_1 \frac{PCT_{\text{abundance}}}{C_{16}} + \omega_2 \frac{PCT_{\text{biomass}}}{C_{17}} \]

where \( I_{\text{importance}} \) stands for the dominance of a species; \( PCT_{\text{abundance}} \) and \( PCT_{\text{biomass}} \) refer to the ratio of the species’ abundance and biomass to the total for the communities, respectively; \( \omega_1 \) and \( \omega_2 \) are the weights of abundance and biomass, and let \( \omega_1 = \omega_2 = 0.5 \).

The larger the \( I_{\text{importance}} \) is, the more the species contributes to its community, and the more important it is in the community.

2.2.4. Biodiversity
We employed the commonly used Shannon Index \((H)\) (Spellerberg and Fedor, 2003; Shannon and Weaver, 1949):

\[ H = -\sum_{i=1}^{s} \left( \frac{n_i}{N} \ln \frac{n_i}{N} \right) \]

where \( H \) stands for the biodiversity; \( n_i \) refers to the number of the \( i \)th species, in [individual L\(^{-1}\)]; \( N \) is the total number of all species in a sample, in [individual L\(^{-1}\)]; \( s \) refers to the species type number in a sample. When all species are equally abundant, \( H \) reaches its peak value.

2.2.5. Niche breadth and niche overlap
There are many models to calculate niche breadth and overlap (Levins, 1968; Pianka, 1974; Hurlbert, 1978; Smith, 1982). In this paper, we employ the widely-used Levins Breadth Model (Levins, 1968; Eq. (3)) and Pianka Overlap Model (Pianka, 1974; Eq. (4)) to get niche breadth and niche overlap, respectively.

Levins’ Breadth Model: \( B_i = 1/ \sum_{j=1}^{k} P_{ij}^2 \) \( \hspace{1cm} (3) \)

where \( B_i \) is the niche breadth of species \( i \); \( P_{ij} \) stands for the ratio of the number of individuals of species \( i \) in resource state \( j \) to the total number of individuals of species \( i \). \( R \) refers to the total number of resource states. Resource states are defined according to national water quality criteria. They stand for gradients along one available resource. Resources available include biochemical oxygen demand (BOD5), dissolved oxygen (DO), permanganate index (COD\(_{\text{Mn}}\)), ammonical nitrogen (NH\(_3\)-N), and so on.

Pianka Overlap Model: \( O_{ik} = \frac{1}{\sqrt{R \sum_{j=1}^{k} P_{ij}^2 \sum_{j=1}^{k} P_{kj}^2}} \)

where \( O_{ik} \) is the niche overlap of species \( i \) on species \( k \); \( P_{ij} \) and \( P_{kj} \) are respectively the ratios of numbers of individuals of species \( i \) and species \( k \) in resource state \( j \) to the total number of individuals of species \( i \) and \( k \); \( O_{ik} \neq O_{ki} \). Calculation of niche breadth and niche overlap was conducted using the software “Data Processing System (DPS)” (Tang and Feng, 2007).

2.2.6. Random forests (RFs) and statistical methods
There are many methods available for classification of sampling sites. Dudoit et al. (2002) compared and reviewed these (Mehrian et al., 2007). Clustering is important for pattern recognition, classification, model reduction and optimization (Hardin and Rocke, 2004; Shaﬁ et al., 2010). Most clustering algorithms require as input a dissimilarity measure between samples (Shi et al., 2005) whereas an unsupervised learning method – random forests (RFs) – does not require this. An RF predictor is an ensemble of individual classiﬁcation tree predictors (Breiman, 2001). For each observation, each individual tree falls into one class and the forest predicts the class that has the largest number of trees. The user has to specify the number of randomly selected variables \( \text{mtry} \) to be searched through for the best split at each node (Horvath et al., 2007). Injecting the right kind of randomness makes RFs accurate classiﬁers. Using out-of-bag estimation makes concrete the otherwise theoretical values of strength and correlation. Random inputs and random features produce good results in classiﬁcation (Breiman, 2001).

In addition, RFs have a number of theoretical advantages (Shi et al., 2005). First, the clustering results do not change when one or more covariates are monotonically transformed since the dissimilarity only depends on the feature ranks and one does not need to worry about symmetrizing skewed covariate distributions. Second, the random forest dissimilarity weights the contributions of each covariate on the dissimilarity in a natural way: the more related the covariate is to other covariates, the more it will affect the deﬁnition of the random forest dissimilarity. Third, the random forest dissimilarity does not require the user to specify threshold values for dichotomizing expressions. It automatically dichotomizes the expressions in a consistent, data-driven way based on individual tree predictors. Fourth, the random forest dissimilarity naturally accommodates missing values.

The research of Horvath et al. (2007) indicates that the results of a RF analysis are highly robust with respect to the RF parameter \( \text{mtry} \) (the number of variables considered at each split). The default value for the number of random features is the square root of the number of variables. A low value of \( \text{mtry} \) is appropriate when most variables are highly correlated to the outcome. If there is no \( a \) priori knowledge, to choose a high value of \( \text{mtry} \) is a better choice. Also, Horvath et al. (2007) found that large values of \( \text{mtry} \) (5000) leads to higher prediction accuracy and a lot of trees (30,000) make the estimate of the importance measure stable.

In this paper, RFs were employed to cluster the 71 sampling sites, taking as inputs a total of 29 variables (dominance, biodiversity, nine water quality indices, nine niche breadth and nine niche overlap variables along the water chemistry indices’ gradients). The sites were regarded as points in two-dimensional multidimensional scaling plots. Then the characteristics of every cluster were identified, based on which key regions were recognized.

To test whether the differences between clusters were signiﬁcant, a nonparametric test, the Kruskal–Wallis Test, was employed in our study. This test neither makes assumptions for a distribution, nor assumes that any particular distribution is being used. It is often used in tests for multiple independent samples with different sample sizes. Every RF clustering was accompanied by a Kruskal–Wallis Test. The clustering result with minimum asymptotic signiﬁcance (\( P \)-value) was selected as the optimized outcome.

The RFs clustering and Kruskal–Wallis Test analyses were conducted with the freely available software R (R Development Core Team, 2011).
3. Results and discussion

3.1. Phytoplankton community structure and representative species selection

There were, in total, 5 classes and 39 species in the sampled Huai River phytoplankton communities. Among them were 17 species of Chlorophyta; 12 Cyanophyta; 6 Bacillariophyta; 2 Euglenophyta; and 2 Pyrrophyta. Cyanophyta was the overwhelmingly dominant class in the HRB phytoplankton communities (94.38% of total abundance); Chlorophyta ranked second but only accounted for 5.38% of the total abundance; Euglenophyta for 3%, Pyrrophyta, for 2% and Bacillariophyta, for 0.19%

Among the Cyanophyta, Anabaenomenon flos-aquae had the greatest abundance (accounting for 40.00% of the total in the HRB phytoplankton communities) while Oscillatoria had the greatest biomass (61.32% of total communities). On the whole, A. flos-aquae and Oscillatoria made the largest contribution in terms of their dominance values in the HRB phytoplankton communities (24.27% and 39.01%, respectively). The 10 species with dominance values larger than 1% are listed in Table 2. Their dominance values summed to 92.71%, so they were selected to represent the whole HRB phytoplankton communities in the study period.

3.2. Ecological niche of the representative species

We computed the ecological niche breadth of the 10 representative species along the nine main water chemistry indices (Table 3) based on Eq. (3).

Generally, a species with a wider niche breadth has a greater adaptability while one with a narrower niche breadth is sensitive to environmental change. The former often has a much greater chance of survival than the latter under conditions of limited resources; however, the latter is usually more competitive in its local habitat when resources are abundant because of its higher efficiency in use of resources (Chen et al., 2009).

Among the 10 species, Oscillatoria had the broadest mean breadth value (3.536), suggesting greater adaptability in a changing environment. It had the broadest niche breadth along the TP gradient (5.252) and the narrowest one along TN (2.341). In contrast, Anabaena azotica had the narrowest mean niche breadth (1.889), signifying a poor adaptability to changing environmental conditions.

The 10 species had broader niche breadths along the WT, TP and CODCr gradients on average (means: 4.143, 3.465 and 3.368, respectively). The narrowest breadth on average was along the NH3–N gradient, as discussed above. Compared with niche breadth (Table 3), niche overlap in Table 4 approximates a reverse trend, which implies that species in the HRB are coexisting with each other with little competition for WT, TP, CODCr, and DO.

The overlap values for each species are listed in Table 5. D. acicularis had the largest total overlap with other species. Spirulina major had the second largest overlap with others, while A. azotica, with the smallest mean breadth, had the least total overlap with other species. According to the research of Jiang et al. (2009), we concluded that D. acicularis has the greatest similarity in resource use with the others while A. azotica has the least similarity. Overall, the total overlap and mean breadth of the phytoplankton species in the Huai River have the same trend (Total_overlap = 1.13 * Mean_breadth + 3.90, R² = 0.702. F = 20.64 > F0.01 = 5.35) – a wider mean niche breadth of a species usually leads to a greater overlap with other species.

3.3. Spatially clustering sites using random forests

To make the estimate of importance measure more stable and keep a higher prediction accuracy, we chose a large number of trees (ntree: 30 000) for each random forest fit and used a large number of random features (mtry: 5000) as recommended by Horvath et al. (2007). The optimized outcome (P = 0.02 < 0.05 with the Kruskal–Wallis Test) showed that the 71 sites were grouped into six clusters with different sizes of 20, 10, 14, 13, 5 and 9 sites, respectively. The P-value indicated that there were significant dissimilarities among the six clusters. The optimized clustering results are shown in Fig. 2.

To facilitate study of the characteristics of every cluster, we divided every factor of dominance, biodiversity, water chemistry, mean ecological niche breadth and mean ecological niche overlap into six different grades: highest, higher, middle higher (MH); middle lower (ML), lower and lowest. Then we analyzed the characteristics of the six clusters. We found the following:

- Cluster 1 was associated with lower-grade pollution, consisting of a middle-lower NH3–N concentration, lower concentration of CODMn, CODCr, TP, TN and fluoride; a lower dominance and a higher biodiversity, a middle lower mean breadth value and a middle-higher mean overlap value.

- Cluster 2 had the lowest pollution including the lowest concentration of NH3–N, CODMn, TP, TN and fluoride plus a middle lower CODCr concentration; the lowest dominance and a lower biodiversity; a lower mean breadth and a middle lower mean overlap.
Cluster 3 had a middle-lower pollution level, with middle-lower concentrations of TP, TN and fluoride, a lower NH₃–N concentration, and the middle concentration of CODₘₙ and CODₖₜ. It also had a higher dominance but the lowest biodiversity in addition to the lowest mean breadth and overlap.

Cluster 4 had a higher pollution level, with higher concentrations of NH₃–N, CODₘₙ, CODₖₜ, TP and TN, plus the highest fluoride concentration. Besides, it had a midly higher dominance and a middle lower biodiversity. Its mean breadth ranked the highest in the six clusters and its mean overlap was higher.
Cluster 5 had the highest pollution level because of the highest concentration of NH₃–N, CODMn, CODCr, TP and TN, and a higher fluoride concentration. It had the highest dominance and biodiversity. Its mean breadth was middle higher and its mean overlap was lower.

Cluster 6 had a middle higher pollution level, composed of middle higher concentrations of NH₃–N, TP, TN and fluoride, plus a middle lower CODMn and the lowest CODCr concentration. Additionally, it had a middle lower dominance and a middle higher biodiversity. Its mean breadth was higher and the mean overlap ranked the highest in the HRB phytoplankton communities during our study period.

The spatial distribution of the six clusters is shown in Fig. 3. The effect of dams made all clusters disperse throughout the HRB except for clusters 4 and 5. Cluster 1 was scattered in the southern and eastern regions of the HRB; cluster 2 was mainly distributed along the main stream of the Huai River; most of clusters 3 and 6 were mainly found in the eastern region of the HRB, with a few sites scattered in the western region; clusters 4 and 5, were uniformly concentrated in the central-northern region of the HRB.

The regions of cluster 5 (thick cross in Fig. 3) have been severely polluted and the concentrations of NH₃–N, CODMn, CODCr, TP, TN were extremely high. Meanwhile, phytoplankton in these regions made the greatest contributions to the HRB phytoplankton communities due to their highest dominance and biodiversity.

The regions of cluster 4 (thin cross in Fig. 3) were less polluted compared with those of cluster 5. The biodiversity of phytoplankton in these regions was low, which resulted in a lower self-purifying rate in these waters. That in turn makes water quality worse. The emphasis should be laid on improvement of biodiversity and pollutant control in these regions.

In summary, regions of clusters 4 and 5 (within the ellipse in Fig. 3) were severely polluted. Phytoplankton there are highly important to the HRB phytoplankton communities due to the high
dominance and biodiversity in these regions. While ecological restoration measures might cause damage to the habitat and the biota, the relatively high mean breadth values in these regions indicate that the phytoplankton there have strong adaptability to environmental change. Ecological restoration focusing on aquatic environment improvement in the regions of clusters 4 and 5 is, therefore, feasible, and the importance of the phytoplankton there means that these are undoubtedly key regions for future phytoplankton-related ecological restoration.

Several previous studies on the HRB have also found that the northern plain regions of the HRB (approximate the ellipse area in Fig. 3), and especially the Hongru (R2), Shaying (R3) and Guo (R4) Rivers, have the most severe water pollution (Wang and Ongley, 2004; Cheng et al., 2005; Tang et al., 2008; Zhang and Shan, 2008; Xia et al., 2011). These regions have a high population and high levels of industrial activities (e.g. mining of sulfate and chloride minerals), and runoff from agricultural activities (Zhang et al., 2011; Zhao et al., 2008). Zhang et al. (2011) studied the chemistry of these rivers of the HRB, and found that the northern plain regions in the HRB have high concentrations of ions, with the spatial patterns and ionic composition reflecting the intensive human activities in the region. The middle and lower reaches of most rivers have fragile or even unstable aquatic ecosystems, with sub-healthy or unhealthy aquatic habitats (Zhao et al., 2008). Our methods also identified these areas as high-priority sites key sites for restoration, demonstrating that our methods are effective and practical.

We aimed to explore the impact of water pollution on phytoplankton communities after we had identified water pollution as the most important factor influencing the quality of phytoplankton communities. For this purpose, some variables that directly control phytoplankton species composition such as light, silica and residence time were not included within the study. This may result in some uncertainties in results.

The quantity and nature of the required data can limit the effectiveness of methods for identification of key regions in a basin with severe water pollution. Turpie et al. (2002) devised a method for prioritizing South African estuaries on the basis of conservation importance. Estuaries were scored in terms of their size, type and biogeographical zone, habitats and biota (plants, invertebrates, fish and birds). This method considered as many factors as possible and was developed to aid in decision-making regarding the freshwater requirements of estuaries, and in the development of a sound management strategy for estuaries. However, too many data requirements and uncertainties in estimates of the indices limited its application in data-scarce areas. Sánchez-Fernández et al. (2004) identified high-priority areas for conservation using only data on water beetle presence and distribution. However, they did not consider the impact of habitat factors such as water chemistry and other factors controlling phytoplankton species compositions. Filipe et al. (2004) presented a practical way of ranking watercourses for conservation based on the probability of occurrence of species and criteria for rarity, abundance, and endemic value. However, failure to incorporate critical habitat variables with the proper scale may result in incorrect classifications. Similarly, when Posadas and Crisci (2001) set priorities in conservation by using phylogenetic diversity measures, they failed to consider the influences of habitat factors on aquatic species. Therefore these methods are not satisfactory for recognizing key regions in ecologically degenerated area with severe pollution.

In most developing countries and regions, water pollution is the principle factor hindering the restoration of aquatic ecosystems. At the same time, collection of detailed data on factors influencing the growth of aquatic biota is costly. Consequently, such datasets are usually scarce. In those countries/regions, our methods can be easily employed to identify key regions for aquatic ecosystem restoration by using simple critical impact factors.

In this study, some uncertainties in the conclusions may occur because of limited data and because the results are based on only one sampling time. In the process of ecological restoration, regular, long-term monitoring on aquatic ecosystems will be required. Long-term monitoring is likely to increase the precision of prediction of the key regions for rehabilitation.

4. Conclusions

To identify the key regions for future phytoplankton related ecological restoration in the Huai River basin (HRB), China, we sampled phytoplankton and water chemistry in the HRB. Species dominance, biodiversity and ecological niches (niche breadth and niche overlap) at all sampling sites were calculated. Then, the random forests (RFs) clustering approach was used to classify all sampling sites into six clusters.

Analyses of the characteristics of the six clusters showed that two clusters in a severely-polluted region of the Northern Plain area of the HRB were key candidates for restoration. The phytoplankton in these regions were an important component of the whole HRB biota due to their high dominance and biodiversity. Their wide niche breadths indicated that they could tolerate the environmental disruption associated with restoration activities. During phytoplankton-related restoration in the HRB, water temperature, concentration of total phosphorus and chemical oxygen demand can be altered with few adverse effects on phytoplankton communities, while measures increasing Ammonia Nitrogen concentration would be highly detrimental.

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References


