

Distribution of benthic diatoms in Korean rivers and streams in relation to environmental variables

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Abstract – The diatoms are an ecologically important group of algae that have been extensively studied by ecologists and taxonomists. However, the large-scale patterns of diatom distribution and the factors underlying this distribution are largely unknown. The aims of this study were to identify the large-scale spatial patterns of benthic diatom assemblages in Korean streams and rivers, and to assess the importance of numerous environmental factors on diatom distribution. We classified 720 study sites based on diatom flora. Benthic diatoms, water chemistry, altitude, and riparian land cover and use were characterized by multivariate analyses, Monte Carlo permutation tests, and indicator species analysis. In total, we identified 531 diatom taxa. Diatom assemblages were mostly dominated by species of the genera *Achnanthes*, *Navicula*, *Nitzschia*, *Cocconeis*, *Fragilaria* (*Synedra* included), *Cymbella*, *Gomphonema*, and *Melosira*. Cluster analysis partitioned all 720 sites into eight groups based on diatom species composition. Canonical correspondence analysis indicated that altitude, land cover and use, current velocity, electrical conductivity, and nutrient levels explained a significant amount of the variation in the composition of assemblages of benthic diatoms. At the national scale, a downstream ecological gradient was apparent, from fast-flowing, mostly oligotrophic highland streams to slow-flowing, mostly eutrophic lowland rivers. Our data suggest that spatial factors explain some of the variation in diatom distribution. The present investigation of the spatial patterns of benthic diatoms, the ecological determinants of diatom occurrence, and the identification of diatom indicator species contributes to development of a program for assessing the biological integrity of lotic ecosystems in Korea.

Key words: Benthic diatoms / spatial patterns / multivariate analyses / ecological gradient / bioassessment

Introduction

Diatoms are the most diverse group of algae in rivers and streams (Leland and Porter, 2000). They constitute a large proportion of the total algal biomass in many environments, and are a high-quality food source for higher trophic levels in aquatic food webs (Stevenson *et al.*, 1996).

Many studies have reported a wide distribution of benthic diatoms (Watanabe *et al.*, 1990; Choi *et al.*, 1995; Leland and Porter, 2000; Weckström and Korhola, 2001; Potapova and Charles, 2002; Soininen *et al.*, 2004; Leira and Sabater, 2005; Bona *et al.*, 2007; Wu *et al.*, 2009) and their tolerance to gradients of diverse environmental variables (Watanabe *et al.*, 1990; Potapova and Charles, 2002). However, other studies have reported that some species only occur in particular geographical locations, water bodies, or micro-habitats (Kociolek and Spaulding,

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Table 1. Characteristics of the five studied river watersheds in Korea, and the number of rivers and streams and sites in each watershed. Data from “Water Management Information System” (WAMIS, <http://www.wamis.go.kr>) and “A list of rivers in Korea” (The Ministry of Land, Transport and Maritime Affairs of Korea, 2008).

Watershed	Length of main stream (km)	Number of tributaries	Total stream length (km)	Watershed area (km ²)	Human population	Number of study streams	Number of study sites
Han River	560.0	912	8567.7	41 957.0	23 404 251	170	320
Nakdong River	470.0	1185	9637.6	31 785.0	14 431 507	75	130
Geum River	393.1	876	6134.9	17 537.0	5 721 207	76	130
Youngsan River	117.7	576	3540.4	12 833.4	3 800 240	47	76
Seomjin River	211.9	283	1928.8	4914.3	319 614	20	64
Total	1752.7	3832	29 809.4	109 026.7	47 676 819	388	720

2000). Rivers and streams are complex ecosystems in which many environmental factors vary on spatial and/or temporal scales. These factors include climate, geomorphology, and land use in the watershed as well as the physical, chemical, and biological properties of rivers and streams. Previous studies have shown that the distribution of diatoms depends on environmental factors (Pan *et al.*, 2006) such as climate and eco-hydrological regimes (Weckström and Korhola, 2001), geomorphic characteristics and land use (Leland and Porter, 2000), nutrient concentrations (Biggs and Smith, 2002), ionic composition (Potapova and Charles, 2003), and herbivory (Anderson *et al.*, 1999).

Diatoms are sensitive to physical, chemical, and biological changes in lotic ecosystems, and their very short generation times allow them to respond rapidly to these changes. The sensitivity of diatom physiology to habitat conditions manifests as a great ecological variability in biomass and species composition (Stevenson *et al.*, 1996). This variability, which can lead to uncertainties in ecological status assessment (Kelly *et al.*, 2009), is due to complex interactions among ecological variables that can affect diatom physiology and community composition (Stevenson, 1997). Despite these uncertainties, studies of diatom distribution provide an effective tool for assessing the ecological integrity of various lotic ecosystems (Kelly and Whitton, 1995; Whitton and Rott, 1996; Kelly, 2002).

Since Skvortzow first reported the presence of freshwater diatoms in Korea in 1929 (Skvortzow, 1929), there has been substantial development of diatom taxonomy. A total of 1457 species of diatoms have been identified in freshwater and marine ecosystems in Korea, among which 724 species have been reported in freshwaters (Choi *et al.*, 1995). However, there is limited information on diatom distribution and biogeography in Korean lotic ecosystems and on the use of benthic diatom assemblages for bioassessment (Lee and Chung, 1992; Hwang *et al.*, 2006; Kim, 2007). The present study was conducted as part of a Korean government-led nationwide biological survey of rivers and streams (MOE/NIER, 2008) that aims to develop national biological criteria under the National Aquatic Ecological Monitoring Program (NAEMP). The purpose of the NAEMP is to establish a national bio-monitoring network to assess the biological and ecological status of stream and river ecosystems, and to develop a strategy for the restoration and management of

disturbed systems. The NAEMP also aims to assess macro-invertebrates, fish, and riparian habitats.

We performed a synoptic study of the spatial distribution of benthic diatoms in Korea in relation to numerous environmental variables. Our specific objectives were to (i) characterize the geographic distribution of benthic diatom assemblages, (ii) identify the major environmental factors that affect diatom distribution, and (iii) identify diatom indicator species and the major factors that affect their presence.

Materials and methods

Study sites

Samples were collected at 720 study sites from 388 streams and rivers in the five major river systems of Korea during September and October 2009 as a part of NAEMP (Table 1, and see also Fig. 3). The Han River Watershed (HRW), located in the central region of the Korean peninsula, and the Nakdong River Watershed (NRW), in the southeast of the peninsula, include most of the study area and of the human population of Korea. The Han River runs through Seoul, the biggest city in Korea, and the Nakdong River runs through Busan and Daegu, the second and third biggest cities, respectively. The Geum River Watershed (GRW) and the Youngsan River Watershed (YRW) are in the western part of the country, and the Seomjin River Watershed (SRW) is between the NRW and the YRW (Fig. 3). The largest number of study sites were in the HRW (320) followed by the watersheds of the NR (130), the GR (130), the YR (76), and the SR (64) (Table 1). All sampling and field measurements were conducted according to the guidelines of the “National surveys for stream ecosystem health” (MOE/NIER, 2008).

Analysis of environmental data

Physico-chemical and hydrological factors were measured at all 720 sampling sites. These included measurements of water temperature, dissolved oxygen concentration, electrical conductivity (EC), and pH, which were measured *in situ* with a multi-probe meter (YSI 6920, YSI Inc., USA). Stream and river water was sampled for analysis of water quality variables. Three water samples

(2 L each) were collected in sterile plastic bottles at each site and transported to the laboratory on ice. Biological oxygen demand (BOD), total nitrogen (TN), NO₃-N, NH₃-N, PO₄-P, and total phosphorus (TP) were determined by standard methods (APHA, 2001). The current velocity was measured at each sampling site using a current meter (Model 2100, Swoffer Inc., USA). In addition, reach-scale riparian conditions were assessed in the area adjacent to each study site according to US EPA guidelines (Barbour *et al.*, 1999) for qualitative habitat analysis, and were recorded as a percentage of land use and land cover type (*e.g.*, forest, agriculture, urban).

Benthic diatom sampling and identification

Most field investigations were conducted in wadeable sites of the selected streams and rivers. To collect benthic diatoms, pebbles (~10 cm diameter) were collected at three to five regions in the riffle zone along a transect at each study site. For deeper sites, in which riffles were not present, substrates were collected at the edge of the target transect. A known surface area of the sampled substrate was scrubbed with a toothbrush and rinsed with distilled water on site to collect surface-attached diatoms. The collected material was placed in a plastic bottle and transported to the laboratory on ice and analyzed within one week. Prior to analysis, all diatom samples from each study transect were combined, and the composite samples were subsampled for subsequent determination of abundance and assemblage structure.

Diatom specimens for microscopy were mounted in Naphrax[®] following the methods of Barbour *et al.* (1999). Diatom counts and identification were performed at ×1000 magnification using a light microscope (Zeiss, Axioskop 2, Germany), and photographs were taken for subsequent use (Roper Scientific Photometrics, COOL SNAP[™]). For assessment of diatom density (cells.cm⁻²), at least 500 diatom cells were counted in each sample.

Diatoms were identified primarily according to Krammer and Lange-Bertalot (1986, 1988, 1991a, 1991b) and Watanabe (2005), although other relevant monographs, illustrations, and articles were also consulted. Identification and counting were performed using common taxonomic criteria based on morphotypes. We included some unidentified specimens in well-established taxa, based on comparisons of morphology and other relevant information, to enable combination of all counts into a single dataset.

Data analysis

Two different multivariate analyses, cluster analysis and canonical correspondence analysis (CCA), were used to characterize relationships between diatoms and environmental variables. Rare taxa, defined as those occurring at < 5% of all total sites, were excluded in the multivariate analysis. Prior to this analysis, benthic diatom density data

were log-transformed to reduce variation; a value of 1 was added to all data points to avoid the problem of log 0 being undefined. The data were rescaled in the range 0–1 based on the min–max transformation, which gave the same level of importance to all species in the analysis.

Cluster analysis was conducted to classify the spatial distribution of benthic diatom communities using the flexible beta method ($\beta = -0.25$) with the Sorenson distance measure (McCune and Grace, 2002). Samples were classified into clusters based on similarities of community composition. A multi-response permutation procedure (MRPP; Mielke *et al.*, 1976), a non-parametric method for testing the hypothesis of “no difference between two or more groups of entities”, was used to evaluate the statistical significance of the clusters. Cluster analysis and the MRPP were conducted using PC-Ord (McCune and Mefford, 1999). Differences in environmental variables among clusters were evaluated using the Kruskal–Wallis test and Dunn’s non-parametric multiple comparisons test with STATISTICA software (StatSoft, 2004).

Indicator species analysis (IndVal; Dufrene and Legendre, 1997) was used to identify potential indicator species in each cluster defined in the cluster analysis. The indicator value for each species in a group is the product of its relative abundance and relative frequency (×100), and ranges from 0 (no indication) to 100 (perfect indication) (Petersen and Keister, 2003). Thus, a perfect indicator of a group should be consistently and uniquely associated with that group, and should never occur in other groups (McCune and Grace, 2002). In this study, a species in a cluster that had an IndVal more than five times greater than in any other cluster was defined as good indicator species for that cluster (Keister and Peterson, 2003). A Monte Carlo method was used to determine the significance of IndVals.

CCA was also used to evaluate the relationships between the distribution of diatom assemblage types and environmental variables by use of a previously described procedure (ter Braak, 1987). The significance of each CCA axis was tested using unrestricted Monte Carlo permutations. To assist interpretation of changes in community profile between dimension scores, Pearson’s correlation coefficients between scores and environmental variables were calculated. Cluster analysis, IndVal, and CCA were conducted using PC-Ord software (McCune and Mefford, 1999).

Results

Environmental factors

Table 2 shows a summary of selected variables pooled from studied rivers and streams of each major river watershed, including altitude, riparian land cover and type of use, current velocity, and water chemistry. The rivers we studied varied widely in water chemistry, physiography, and land cover and type of use. On average, rivers and streams of the HRW had the highest altitude

Table 2. Summary of selected environmental variables of rivers and streams in each major river watershed in Korea.

Watershed	Mean	Max	Min	Altitude (m)	Land cover/use type (%)			Current velocity (cm.s ⁻¹) ^a	Electric conductivity (μS.cm ⁻¹)	BOD (mg.L ⁻¹)	TN (mg.L ⁻¹)	TP (mg.L ⁻¹)	NH ₃ -N (mg.L ⁻¹)	NO ₃ -N (mg.L ⁻¹)	PO ₄ -P (mg.L ⁻¹)
					Forest	Agriculture	Urban								
Han River	Mean	147.5	35.8	24.7	31.7	51.6	248.6	2.1	2.6	0.11	0.13	2.0	0.07		
	Max	721.0	100.0	100.0	100.0	140.0	1600.0	7.8	17.8	1.56	7.71	9.6	0.88		
	Min	1.0	0.0	0.0	0.0	0.0	30.0	0.6	0.3	0.00	0.00	0.3	0.00		
Nakdong River	Mean	89.6	31.4	42.5	23.0	28.6	229.9	2.1	1.9	0.07	0.03	1.5	0.04		
	Max	629.0	100.0	100.0	100.0	115.7	1366.0	5.8	6.4	0.56	0.91	5.0	0.35		
	Min	1.0	0.0	0.0	0.0	0.0	43.3	0.6	0.3	0.00	0.00	0.3	0.00		
Geum River	Mean	57.7	17.9	46.7	30.9	54.6	262.4	1.9	2.4	0.12	0.20	1.3	0.08		
	Max	278.0	100.0	100.0	100.0	137.7	1138.0	18.7	29.0	2.90	5.98	8.8	2.69		
	Min	0.0	0.0	0.0	0.0	0.0	64.4	0.0	0.3	0.00	0.00	0.0	0.00		
Youngsan River	Mean	32.5	22.6	45.4	31.4	29.4	223.2	1.9	2.1	0.12	0.19	1.7	0.09		
	Max	211.0	100.0	90.0	90.0	94.7	1971.0	8.0	10.4	0.48	3.86	10.2	0.46		
	Min	0.0	0.0	0.0	0.0	2.5	25.3	0.0	0.4	0.01	0.00	0.1	0.00		
Seonjin River	Mean	118.1	48.8	35.9	14.6	12.9	131.5	0.8	1.2	0.08	0.02	1.0	0.03		
	Max	335.0	95.0	80.0	80.0	35.0	255.8	2.6	2.3	1.14	0.09	2.0	0.11		
	Min	1.0	1.0	0.0	0.0	0.0	29.1	0.0	0.3	0.01	0.00	0.2	0.00		
Overall	Mean	106.1	31.5	35.0	28.4	41.4	234.8	1.9	2.3	0.10	0.12	1.7	0.07		
	Max	721.0	100.0	100.0	100.0	140.0	1971.0	18.7	29.0	2.90	7.71	10.2	2.69		
	Min	0.0	0.0	0.0	0.0	0.0	25.3	0.0	0.3	0.00	0.00	0.0	0.00		

A value of 0 cm.s⁻¹ for current velocity indicates that the sampling site was at the edge of an unswimmable large river, not that the river is not flowing.

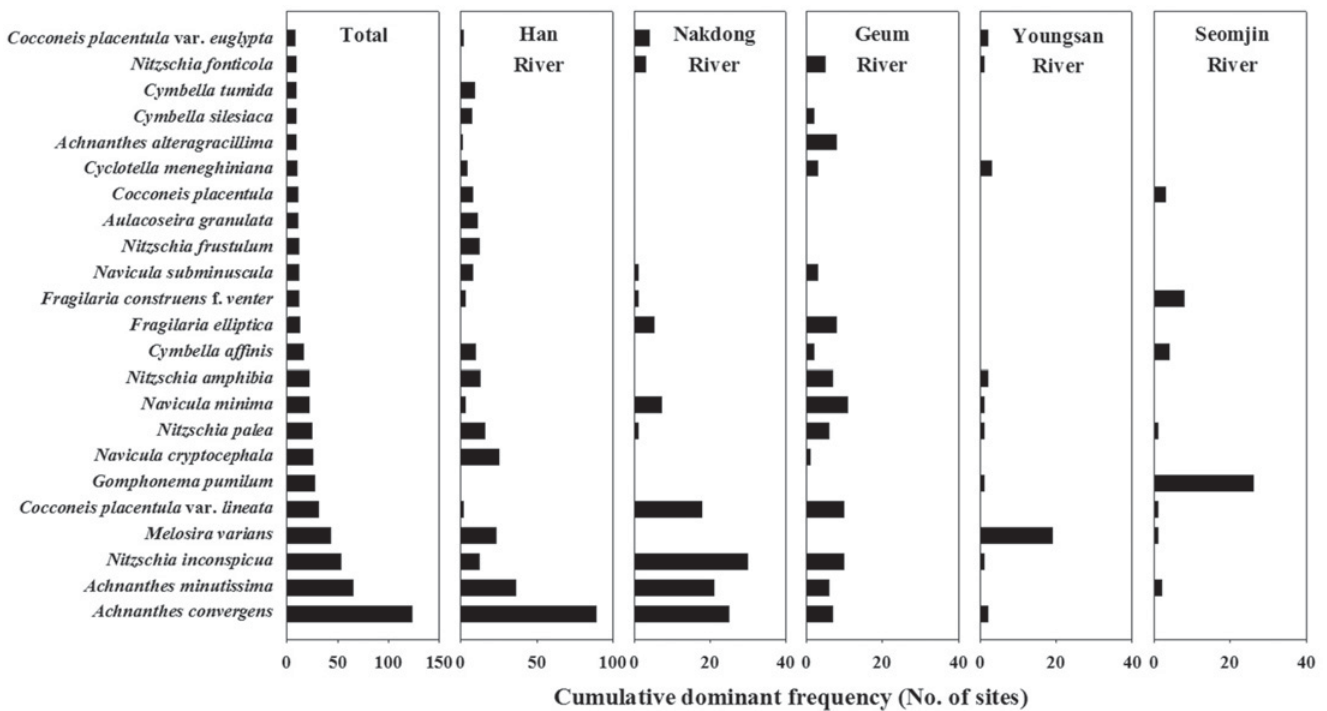


Fig. 1. Diatom distribution in the five major river systems in Korea. The abscissa indicates the percentage of sites in which each diatom species occurred, and a different scale was used for each river system.

(mean 147.5 m) and the highest proportion of forest land cover (35.8%); the altitude and proportion of forest cover of rivers and streams in the SRW were also significant. The rivers and streams in the NRW, the GRW, and the YRW were at low altitude and had relatively high proportions of agricultural land. Among all 720 study sites, the current velocity ranged from 0 to 140.0 $\text{cm}\cdot\text{s}^{-1}$; among the five major river watersheds, the average current velocity ranged from 12.9 to 54.6 $\text{cm}\cdot\text{s}^{-1}$. Relatively fast flow was recorded at the sites of the HRW and GRW, and slow flow at the SRW. The average levels of electric conductivity, BOD, and nutrients (N and P) were low at the SRW sites, indicating rather homogeneous water chemistry relative to the other river systems.

Community composition

A total of 531 diatom taxa were recorded from the 720 study sites. The YRW had the greatest number of species (340 taxa), followed by the HRW (287 taxa), GRW (259 taxa), NRW (179 taxa), and SRW (161 taxa) (see the taxonomic list in the Appendix). There were 81 taxa that occurred at more than 5% of all sites, and these taxa were used for subsequent site classification analysis.

The dominant taxa varied among the different river systems. *Achnanthes convergens* H. Kobayasi was the most common species and was present at 17% of all sites; this species was the most dominant species in the HRW, and the second most dominant species in the NRW (Fig. 1). Interestingly, *A. convergens* was rare in the Youngsan River System (YRS) and Seomjin River System (SRS);

instead, *Melosira varians* C. Agardh was predominant in the YRW and *Gomphonema pumilum* (Grunow) Reichardt & Lange-Bertalot was predominant in the SRW. *Achnanthes minutissima* Kützing was the second most common species nationwide (observed at 9.0% of all sites), and was predominant in the HRW and the NRW. *Nitzschia inconspicua* Grunow was present at 7.4% of all sites, and was particularly dominant in the NRW. *M. varians* occurred at 6.0% of all sites, and was the most dominant species in the YRW. *Cocconeis placentula* var. *lineata* (Ehrenberg) Van Heurck occurred at 4.3% of all sites, and was predominant in the NRW and the GRW. These six benthic diatom species accounted for 48% of the diatoms in all 720 study sites.

Diatom-based site classification

Based on similarities of diatom community composition, cluster analysis (Sorensen's distance measure) classified all 720 study sites into four clusters or eight clusters (Figs. 2 and 3). Group 1 (318 sites) was divided into four subgroups (Group 1a, 89 sites; Group 1b, 45 sites; Group 1c, 67 sites; Group 1d, 117 sites); Group 2 (153 sites) was divided into two subgroups (Group 2a, 86 sites; Group 2b, 67 sites); Group 3 (57 sites) and Group 4 (192 sites) were not subdivided (Fig. 2). The MRPP indicated statistically significant differences among the four clusters and among the eight clusters ($A = 0.150$, $P < 0.001$).

Differences of clusters were associated with differences in environmental conditions and geographical location of sampling sites, especially with hydrological systems.

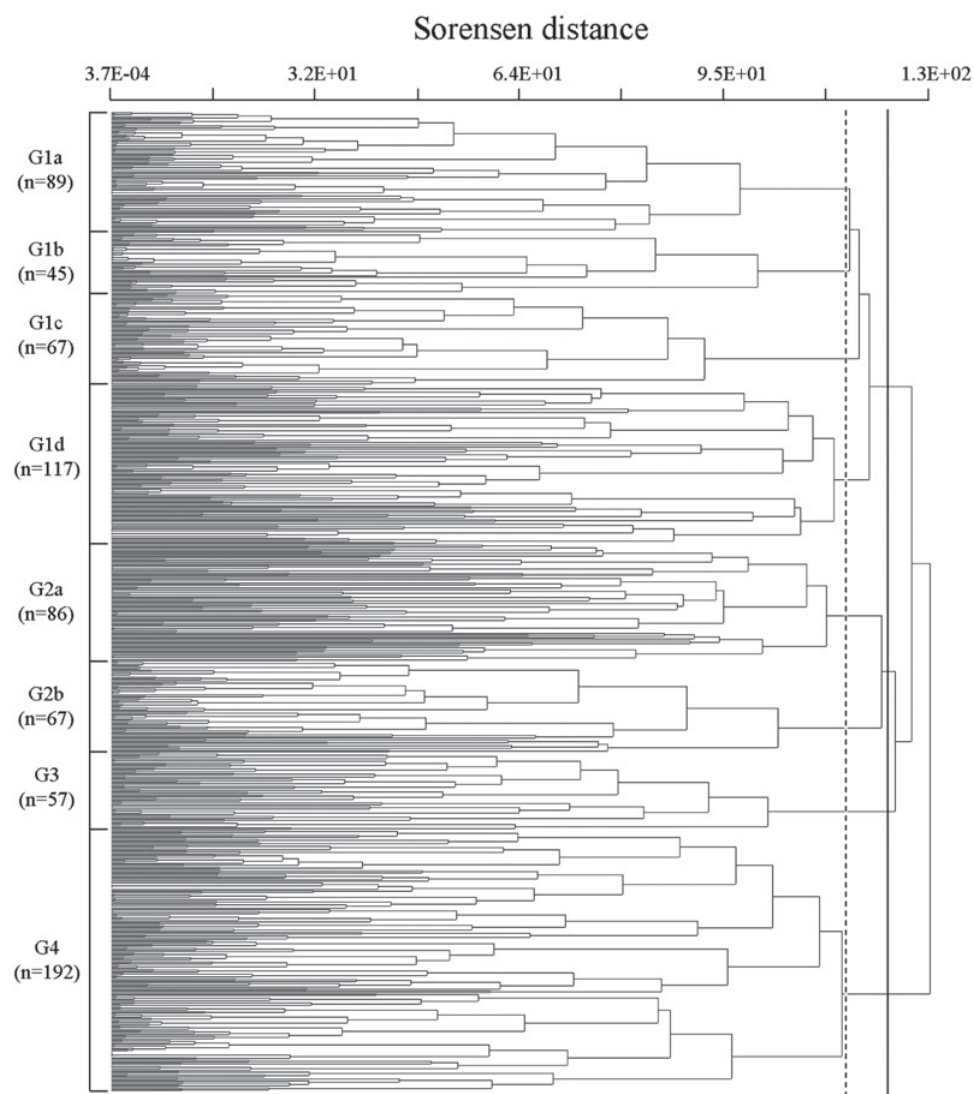


Fig. 2. Dendrogram of diatom-based stream site classification using Sorensen's clustering method. The solid line differentiates the four diatom-based groups and the dashed line differentiates the eight diatom-based groups. See the text for abbreviations of group names.

In particular, sampling sites in Group 1a were in the mountainous tributaries of the HRW and NRW (Fig. 3), and had the highest mean altitude (221 m) of all groups (Fig. 4). Sites of Group 1b had a high proportion of forest land cover (50%), and a low proportion of agricultural (28%) and urban land use (20%). The hydrogeography of Group 1a sites (*i.e.*, mountainous locations) seemed to affect their water chemistry, but the results did not indicate a clear association for all sites of this group, maybe due to their scattered distribution. Geographic location and water chemistry had a clearer relationship in the sites of Group 1b. Group 1b sites were mostly in the eastern mountainous region (Fig. 3), in which there is a high mean altitude (201 m), significant forest land (73%), significant current velocity (mean $63 \text{ m}\cdot\text{s}^{-1}$), low nutrient concentrations, and low EC (Fig. 4). Sites of Group 1c were almost all in regions downstream of the South Han River, where human activities, such as agriculture and urban development, were predominant (Fig. 3). In Group 1c, the

mean concentrations of TN ($2.9 \text{ mg}\cdot\text{L}^{-1}$) and $\text{NO}_3\text{-N}$ ($2.3 \text{ mg}\cdot\text{L}^{-1}$) were in the high range (Fig. 4). Sites of Group 1d were dispersed across the country (Fig. 3), had a variety of different land uses, and had nutrient concentrations in the low range. In particular, mean concentrations of TN ($0.03 \text{ mg}\cdot\text{L}^{-1}$) and $\text{PO}_4\text{-P}$ ($0.01 \text{ mg}\cdot\text{L}^{-1}$) were the lowest among all groups (Fig. 4).

The sites of Groups 2a and 2b were almost exclusively in lowland agricultural and urban areas, and included the downstream Han River and most of the YRW, respectively (Fig. 3). Both groups had relatively low flow and very eutrophic waters. Group 2b had the highest mean nutrient enrichment of all eight groups ($P < 0.01$) (Fig. 4).

The sites of Group 3 were mostly in the SRW (Fig. 3). These sites had very low proportions of urban land use (Fig. 4). The water quality was good (similar to Group 1b), as indicated by the lowest BOD ($0.8 \text{ mg}\cdot\text{L}^{-1}$) and relatively high nutrient concentrations. Group 3 had the slowest mean current velocity ($12 \text{ cm}\cdot\text{s}^{-1}$).

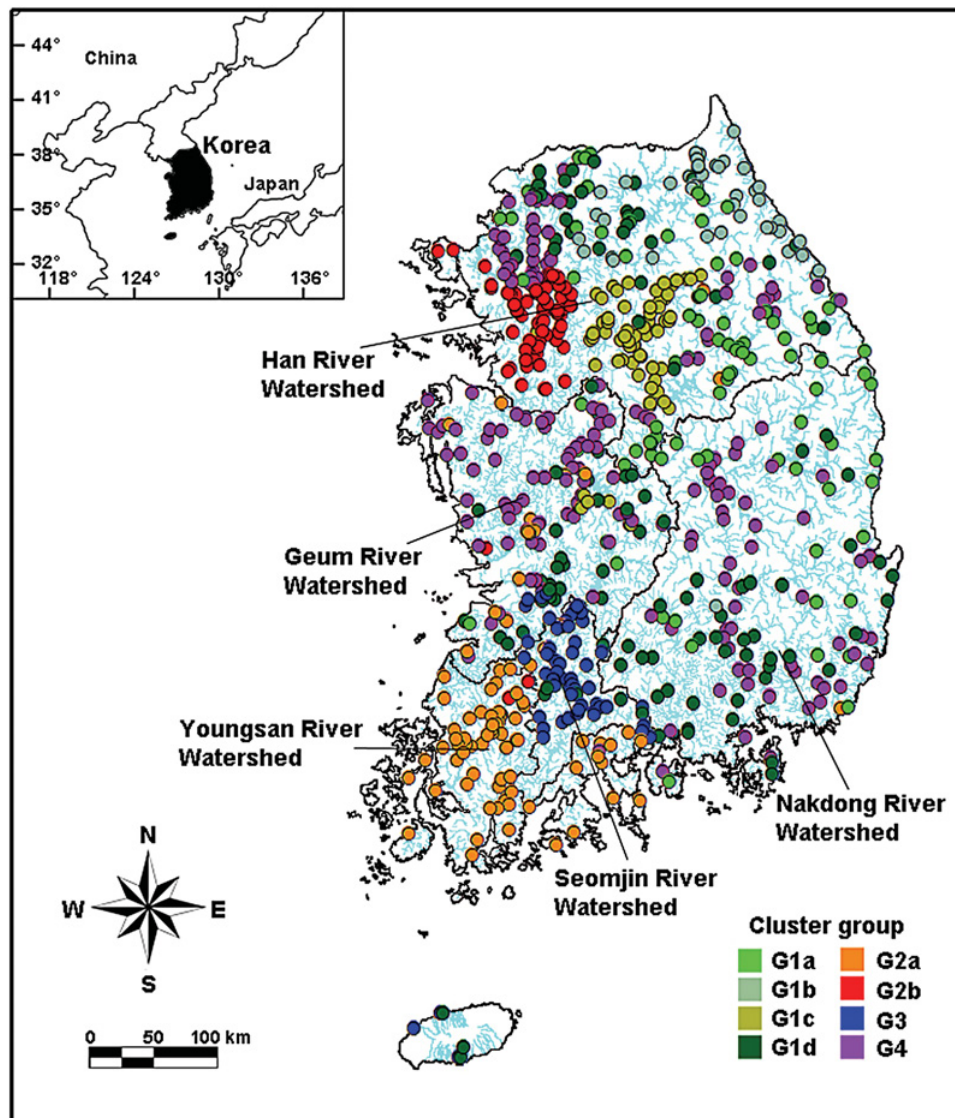


Fig. 3. Geographical position of the eight diatom-based site groups.

The sites of Group 4 had a wide geographic distribution and were mostly present in lowland areas, particularly in the GRW and NRW. This group also included sites of a major tributary (Imjin River) that is located in the northwestern region of the HRW (Fig. 3). Land uses of these sites were variable and water quality was intermediate (Fig. 4).

We used IndVal to identify indicator species in each of the eight groups (Table 3). A total of 81 species were present in more than 5% of all study sites and different numbers of indicator species were present in the different groups, ranging from 17 species (Group 4) to 4 species (Group 1a). The indicator species with high indicator values (>50%) differed among the different groups. The highest indicator value (94%) was for *G. pumilum*, which was common in sites of the SRS (Fig. 1) and Group 3 (Table 3). *Cymbella tumida* (Brébisson) Van Heurck was predominant in the HRW (Fig. 1) and had a high indicator value (65%) in Group 1b (Table 3). *N. inconspicua* was

predominant in sites of the NRW (Fig. 1), and had a high indicator value (86%) in Group 4. *Navicula cryptocephala* Kützing was the predominant species in the HRW, particularly in the downstream region and in Group 2b. Other diatom taxa with indicator values more than 50% were *Aulacoseira granulata* (Ehrenberg) Simonsen, *Fragilaria construens* (Ehrenberg) Grunow, *Navicula minima* Grunow, *A. minutissima*, and *C. placentula* Ehrenberg.

Ordination of diatom assemblages

Next, sites clustered using Sorenson's method were plotted by CCA in the ordination space in relation to environmental variables (Fig. 5). The eigenvalues of the first CCA axis (0.201) and the second CCA axis (0.094) were both significant ($P < 0.01$; 99 Monte Carlo permutations test with 99 permutations), and the total

Table 3. Indicator values (%) for the most important species ($P < 0.05$) clustered in each group. Monte Carlo simulations (999 permutations) were used to assess the significance of each species as an indicator for each site group. There were 81 species that occurred at more than 5% of all sites, and these species were used for site classification analysis. Less important species are not shown.

	Cluster group								<i>P</i> value
	G1a	G1b	G1c	G1d	G2a	G2b	G3	G4	
<i>Achnanthes minutissima</i>	55	0	5	7	0	10	1	6	0.001
<i>Gomphonema clevei</i>	25	19	6	7	0	0	0	1	0.001
<i>Cymbella turgidula</i>	21	0	1	0	2	0	0	2	0.001
<i>Cymbella tumida</i>	2	65	0	1	0	1	0	0	0.001
<i>Achnanthes convergens</i>	22	32	29	2	0	0	0	1	0.001
<i>Gomphonema parvulum</i>	6	32	6	6	2	0	6	2	0.001
<i>Cocconeis placentula</i>	0	0	51	4	0	1	13	0	0.001
<i>Navicula viridula</i>	0	0	42	0	0	0	6	0	0.001
<i>N. viridula</i> var. <i>rostellata</i>	1	1	30	9	0	0	0	10	0.001
<i>Cocconeis placentula</i> var. <i>lineata</i>	14	10	0	26	0	0	3	12	0.001
<i>Nitzschia fonticola</i>	3	2	3	21	0	0	0	21	0.001
<i>Achnanthes alteragracillima</i>	4	0	0	18	0	0	0	1	0.001
<i>Cyclotella meneghiniana</i>	0	0	4	2	24	13	0	10	0.001
<i>Navicula bacillum</i>	0	0	1	0	21	0	0	1	0.001
<i>Cyclotella atomus</i>	0	0	0	0	21	0	0	3	0.001
<i>Navicula cryptocephala</i>	1	1	7	3	0	61	5	1	0.001
<i>Aulacoseira granulata</i>	0	0	0	1	0	60	0	1	0.001
<i>Fragilaria construens</i>	0	0	0	0	0	59	0	0	0.001
<i>Gomphonema pumilum</i>	0	0	0	1	0	0	94	0	0.001
<i>Rhoicosphenia abbreviata</i>	0	0	0	1	0	0	42	0	0.001
<i>Cymbella minuta</i>	1	0	7	7	1	0	34	0	0.001
<i>Nitzschia inconspicua</i>	1	0	1	3	0	0	0	62	0.001
<i>Navicula minima</i>	1	0	2	8	0	0	0	56	0.001
<i>Navicula subminuscula</i>	0	0	12	2	0	0	0	48	0.001
Total number of significant indicator species	4	8	10	10	7	14	7	17	

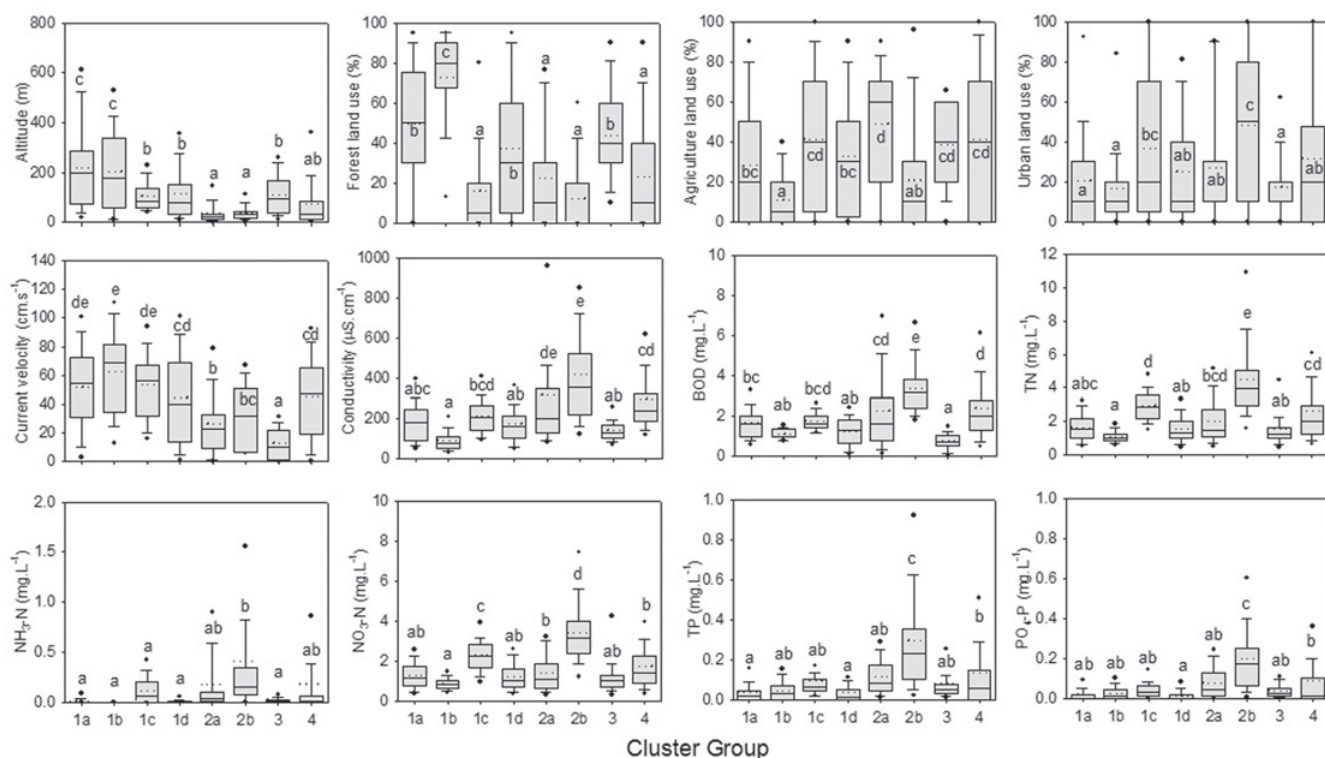


Fig. 4. Boxplots of selected environmental variables of cluster groups. The mean (horizontal dotted line), median (horizontal solid line), range from the 5th to the 95th percentile (dots at the bottom and top of each box), and standard deviation (error bar) are shown in each box. Letters in or on the bars indicate significant difference among cluster groups ($P < 0.05$).

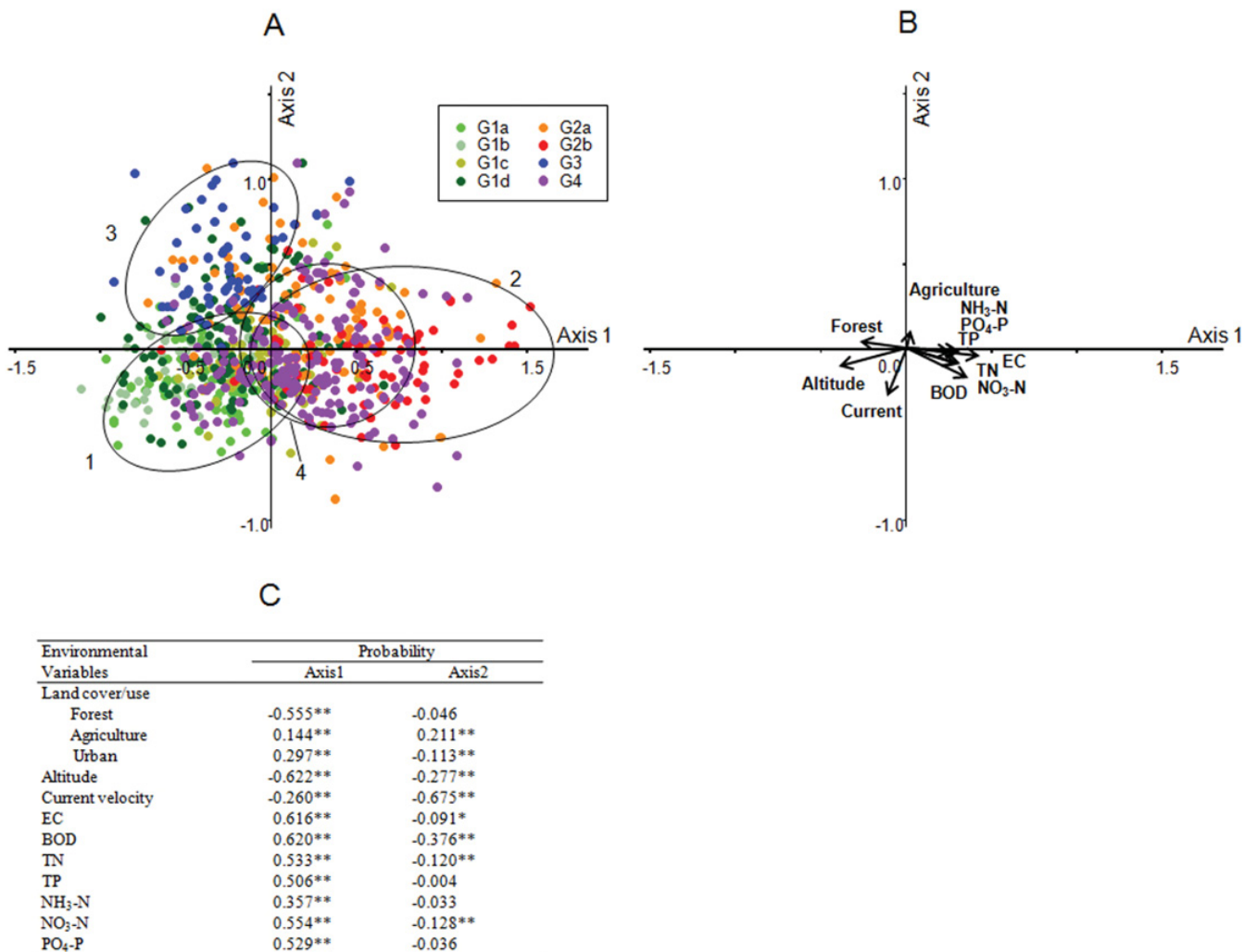


Fig. 5. (A) Ordination diagram showing the distributions of the four and eight cluster groups (denoted by numbers and corresponding circles). (B) Relative contributions of environmental variables in the CCA space. (C) Correlation coefficients and probabilities of environmental variables and the first two CCA axes (* $P < 0.05$, ** $P < 0.01$).

explained variance in the species data was 6.5%. The diatom–environment correlations for CCA axis-1 (0.758) and CCA axis-2 (0.573) were high, indicating a relatively strong relationship between diatom species and measured environmental variables. All the variables included in this analysis were significantly correlated with axis-1, and among these, altitude ($r = -0.622$, $P < 0.01$) was the most significant contributor (Fig. 5C); EC, BOD, and forest land cover were also important variables. Analysis of nutrients indicated that NO₃-N, TN, and PO₄-P had strong correlations with axis-1. Current velocity ($r = -0.675$, $P < 0.01$) was most significantly correlated with axis-2, followed by BOD and altitude (Fig. 5C).

Overall, clusters of diatom assemblages were not significantly separated in the ordination space (Fig. 5A), indicating that site specificities of many diatom species were not strong at the nationwide scale. This result is also supported by the geographical dispersion of sampling sites across Korea (Fig. 3). In particular, the sites of Group 4

were mostly located in the middle of the ordination space, and there was a large overlap with Groups 2a and 2b. Otherwise, the overall site distribution and separation of some site clusters (Groups 1 all together, 2b, and 3) is clear. Altitude and nutrient concentrations were most important in separating the clusters of Groups 1 and 3 from Group 2; current velocity was the most important factor for separating the clusters of Group 3 (Figs. 5B and 5C).

The indicator species in each group were analyzed in relation to land cover and use, hydrographical variables, and water quality variables (Table 4). In spite of some variations in the correlation results, many indicator species reflected the overall conditions of their habitats. The most significant indicator species in Group 1a (*A. minutissima*) had positive correlations with forest land cover and altitude, and negative correlations with nutrients and EC, indicating its preference for oligotrophic waters. *Cymbella turgidula* Grunow was similar to *A. minutissima*, but

Table 4. Indicator species in each cluster, and their correlation coefficients with environmental variables (* $P < 0.05$, ** $P < 0.01$).

Group	Indicator species	N	Land use			Altitude	Current	Conductivity	BOD	TN	TP	NH ₃ -N	NO ₃ -N	PO ₄ -P
			Forest	Agriculture	Urban									
G1a	<i>Achnanthes minutissima</i>	444	0.20**	-0.09	-0.05	0.22**	0.08	-0.14**	-0.11*	-0.12**	-0.09	-0.07	-0.13**	-0.11*
	<i>Gomphonema clevei</i>	243	-0.06	-0.04	0.11	-0.05	-0.04	-0.12	-0.11	-0.12	-0.08	-0.06	-0.11	-0.08
	<i>Cymbella turgidula</i>	101	0.23*	-0.22*	-0.05	0.16	0.11	0.03	-0.04	-0.14	-0.09	-0.11	-0.13	-0.07
G1b	<i>Cymbella tumida</i>	192	0.26**	-0.20**	-0.04	0.09	0.14	-0.26**	-0.11	-0.20**	-0.10	-0.10	-0.19**	-0.08
	<i>Achnanthes convergens</i>	353	0.20**	-0.18**	0.01	0.05	0.11*	-0.17**	-0.07	-0.09	-0.05	-0.08	-0.08	-0.01
	<i>Gomphonema parvulum</i>	340	0.12*	-0.08	-0.02	0.11	0.09	-0.06	-0.06	-0.04	0.00	-0.00	-0.10	0.00
G1c	<i>Cocconeis placentula</i>	168	0.12	-0.00	-0.12	0.13	0.20**	-0.15	-0.06	0.04	-0.07	0.04	0.05	-0.07
	<i>Navicula viridula</i>	71	-0.18	-0.06	0.06	-0.23	-0.23	0.09	-0.02	-0.02	-0.05	-0.05	0.00	-0.06
	<i>N. viridula</i> var. <i>rostellata</i>	208	-0.12	0.07	0.01	-0.14*	-0.03	0.06	0.15*	0.12	0.16*	0.10	0.11	0.14
G1d	<i>Cocconeis placentula</i> var. <i>lineata</i>	359	-0.01	0.06	-0.05	0.06	0.01	-0.09	-0.05	-0.04	-0.06	-0.05	-0.01	-0.06
	<i>Nitzschia fonticola</i>	259	0.07	-0.09	0.03	-0.08	0.01	0.04	-0.10	-0.05	-0.03	-0.01	-0.05	-0.02
G2a	<i>Achnanthes altergracillima</i>	87	0.04	0.00	-0.02	0.08	0.23*	-0.24*	-0.37**	-0.13	-0.19	-0.13	-0.14	-0.16
	<i>Cyclotella meneghiniana</i>	310	-0.07	0.07	-0.02	-0.20**	-0.16**	0.14*	0.27**	0.07	0.08	0.06	0.04	0.05
	<i>Navicula bacillum</i>	61	-0.00	-0.11	0.14	-0.38*	-0.30*	0.37**	0.19	-0.01	0.35**	0.28*	-0.10	0.41**
G2b	<i>Cyclotella atomus</i>	63	-0.16	0.14	0.02	-0.19	-0.32*	0.13	0.39**	0.06	0.17	0.08	-0.02	0.14
	<i>Navicula cryptocephala</i>	315	-0.09	-0.16**	0.19**	-0.08	-0.18**	0.06	0.14*	0.13*	0.09	0.11*	0.17**	0.08
	<i>Aulacoseira granulata</i>	112	-0.09	-0.27**	0.20*	0.16	-0.08	-0.01	0.11	0.02	0.14	-0.05	0.11	0.13
G3	<i>Fragilaria construens</i>	52	0.07	-0.04	-0.17	0.36**	0.11	0.01	-0.23	-0.16	-0.14	-0.13	-0.17	-0.11
	<i>Gomphonema pumilum</i>	90	0.21*	0.03	-0.19	0.04	-0.17	-0.22*	-0.18	-0.05	0.10	-0.07	-0.02	-0.07
	<i>Rhoicosphenia abbreviata</i>	41	-0.23	-0.07	0.37*	-0.23	-0.10	0.00	-0.19	0.46**	0.21	0.30	0.55**	0.33*
G4	<i>Cymbella minuta</i>	162	-0.03	0.01	0.04	0.13	-0.04	-0.14	-0.22**	-0.08	-0.06	-0.05	-0.09	-0.05
	<i>Nitzschia inconspicua</i>	221	0.01	0.10	-0.11	-0.10	-0.00	0.17**	0.00	-0.01	0.01	-0.05	0.04	0.02
	<i>Navicula minima</i>	279	-0.20**	0.12*	0.05	-0.11	-0.02	-0.01	0.08	0.01	0.11	0.08	-0.07	-0.02
	<i>Navicula subminuscula</i>	233	-0.10	-0.12	0.23**	-0.07	0.01	0.16*	0.10	0.20**	0.39**	0.15*	0.26**	0.42**

its correlation with nutrients was not significant. All major indicator species (*C. tumida*, *A. convergens*, and *Gomphonema parvulum* Kützing) were in Group 1b and were significantly correlated with a mountainous character and high proportion of forest land cover. None of the indicator species in Groups 1c and 1d had significant correlations to proportion of forest land cover. Overall, almost all of the indicator species in site clusters of Group 1 (a + b + c + d) had negative correlations with variables of water chemistry, indicating that they preferred oligotrophic waters. This result was supported by the habitat conditions related to the land cover and use patterns near the sampling sites. For example, the correlations between indicator species and nutrients were stronger for Groups 1a and 1b than for Groups 1c and 1d. However, *Navicula viridula* var. *rostellata* (Kützing) Cleve (Group 1c), which seemed to prefer eutrophic lowland waters, was a notable exception. The preference for slowly flowing eutrophic waters is striking for indicator species in the site clusters of Group 2a, particularly *Navicula bacillum* Ehrenberg and *N. cryptocephala* of Group 2b. *Rhoicosphenia abbreviata* (C. Agardh) Lange-Bertalot (Group 3) and *Navicula subminuscula* Manguin (Group 4) also had strong correlations with nutrient concentrations.

Nationwide distribution of benthic diatoms

The major benthic diatoms identified as indicator species in each group exhibited large spatial variations at the national scale (Fig. 6). In particular, *A. minutissima* and *A. convergens* had high abundances almost nationwide (Figs. 6A and 6D), indicating adaptability to a broad range of lotic ecosystems in Korea. Similarly, *Gomphonema clevei* Fricke and *C. tumida* had widespread distributions, but their dominance was regional (Figs. 6B and 6C). *G. clevei* dominated in the four major river systems, except for the SRW, and *C. tumida* was dominant in upstream and headwater sites in mountainous regions of the HRW. *C. placentula* and *N. viridula* (Kützing) Ehrenberg had regionally clumped but distinct spatial separation; their dominance was concentrated in the middle region of the HRW and SRW (Figs. 6E and 6F). *G. pumilum* and *R. abbreviata* had rather restricted distribution in the SRW (Figs. 6M and 6N). The high density of *N. bacillum* and *Cyclotella meneghiana* Kützing was evident in the YRW (Figs. 6I and 6J). However, the occurrence of *C. meneghiana* is peculiar, with dominance mostly in the western region of the country. *N. cryptocephala* and *A. granulata* were found particularly in the downstream regions of the HRW (Figs. 6K and 6L), but the distribution of *N. cryptocephala* was much broader, and it covered more regions of the SRW than *A. granulata*. Several diatoms had a broad occurrence (Figs. 6G, 6H, 6O and 6P). In particular, *C. placentula* var. *lineata* and *Nitzschia fonticola* Grunow largely dominated in the GRW and NRW (Figs. 6G and 6H). *N. inconspicua* and *N. minima* had similar spatial patterns (Figs. 6O and 6P); both were

scattered in many regions, but were absent in the SRW and northeastern mountainous region of the HRW, where *C. tumida* was dominant.

Discussion

Taxonomic composition

We recorded a total of 531 diatom taxa in this nationwide survey of rivers and streams in Korea. This represents about 73% of the 724 freshwater diatom taxa (planktonic and benthic species) that have been identified in Korea (Lee, 1988). The number of diatoms in Korea is lower than that reported in the USA, in which 1548 diatom species (including 87 planktonic forms) were recorded in 2735 samples from rivers (Potapova and Charles, 2002). Diatoms in Korean rivers and streams were dominated by species of *Achnanthes*, *Navicula*, *Nitzschia*, *Cocconeis*, *Fragilaria* (*Synedra* included), *Cymbella*, *Gomphonema*, and *Melosira* (Fig. 1, and also see the Appendix). The major diatom species present in Korean rivers and streams have been recorded in rivers throughout the world, including China (Wu *et al.*, 2009), Hong Kong (Dickman *et al.*, 2005), Japan (Watanabe *et al.*, 1986), the USA (Potapova and Charles, 2003), Spain (Leira and Sabater, 2005), Italy (Bona *et al.*, 2007), Germany (Werner and Köhler, 2005), and Finland (Soininen *et al.*, 2004).

In the present study, we found that many diatom taxa occurred in diverse rivers and streams, but that other taxa had more restricted distributions. For example, *G. pumilum* occurred almost exclusively at sites of the SRW (Fig. 1) and *M. varians* was very common and had the highest frequency in sites of the YRW, but was not exclusive to this system. In addition, the major species appeared to have different abundances in the five major river watersheds (Fig. 1). Based on our cluster analysis, the most common benthic diatoms differed among the different sites of the five river watersheds, but other taxa occurred in two or more systems.

Relationships of benthic diatom distribution and environmental variables

A variety of environmental parameters, singly or in combination, directly and/or indirectly affect the species composition and distribution of stream biota (*e.g.*, Giller and Malmqvist, 1998). In a previous study of diatom ecology, Stevenson (1997) organized various multi-scale factors into a hierarchical framework, in which high-level factors (*e.g.*, climate, geology, land use, and physiography) can restrict low-level factors. Low-level, proximate factors including resources (nutrients), stressors (temperature, toxic substances, ionic strength, flow regime), and habitats directly affect diatom distribution. At spatial and temporal scales, the effects of proximate factors can be constrained by high- or intermediate-level factors.

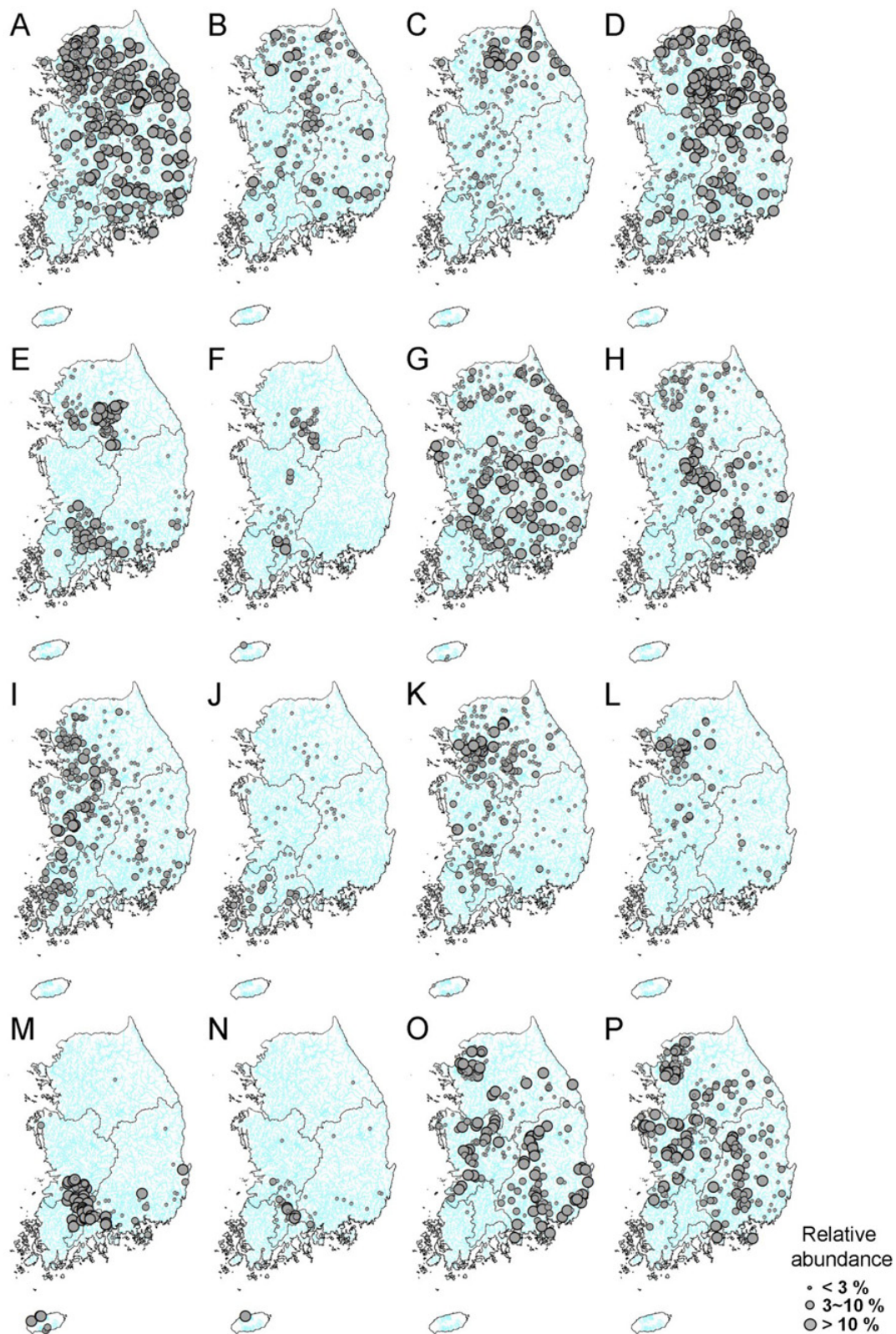


Fig. 6. Spatial distribution of the major diatom indicator species of benthic diatoms in each site cluster. (A) *Achnanthes minutissima*, (B) *Gomphonema clevei*, (C) *Cymbella tumida*, (D) *Achnanthes convergens*, (E) *Cocconeis placentula*, (F) *Navicula viridula*, (G) *Cocconeis placentula* var. *lineata*, (H) *Nitzschia fonticola*, (I) *Cyclotella meneghiniana*, (J) *Navicula bacillum*, (K) *Navicula cryptocephala*, (L) *Aulacoseira granulata*, (M) *Gomphonema pumilum*, (N) *Rhoicosphenia abbreviata*, (O) *Nitzschia inconspicua*, and (P) *Navicula minima*. See [Table 3](#) for corresponding cluster groups.

Thus, the species-specific sensitivity of diatoms to many environmental conditions manifests as great variations of species composition and diatom assemblages in different rivers and streams (Stevenson, 1997; Potapova and Charles, 2002). Previous research has established the effects of various multi-scale factors on diatom community structure and distribution in many rivers and streams (Pan *et al.*, 1999; Leland and Porter, 2000; Potapova and Charles, 2002; Leira and Sabater, 2005; Bona *et al.*, 2007; Wu *et al.*, 2009). The results of the present nationwide study of diatoms in Korea agree with these previous results.

In the present study, cluster analysis and CCA both showed that various multi-scale factors were important in explaining variations in the structure of benthic diatom assemblages of Korean rivers and streams. Our results indicated that a physiographic variable (altitude), land cover and use patterns, chemical variables (EC, BOD, and nutrients), and a physical variable (current velocity) were important factors in site classification. Our CCA analysis showed that altitude was the most important variable separating diatom site groups along axis-1. There were other important factors related to altitude, including forest land cover and current velocity. Considered together, these factors suggest the involvement of a “downstream gradient” (Potapova and Charles, 2002) in the structuring of diatom assemblages at the national scale in Korea. This gradient may not be due to a single factor (*e.g.*, altitude), because a downstream gradient is likely to be complex and associated with other factors, such as nutrient levels, land use, and water flow. In this study, nutrient enrichment appeared to be strongly associated with this complex downstream gradient, as shown previously (*e.g.*, Leira and Sabater, 2005). The site clusters located on the left side of CCA axis-1 represent upstream characteristics, including high altitude, high proportion of forest in the riparian structure, low EC, low BOD, and low nutrients (N and P). Upstream sites of the HRW (Groups 1a and b) and almost all sites in the SRW (Group 3) were clearly separated in the ordination space (Fig. 5). However, the sites of the SRW did not have fast current velocities despite their location at relatively high altitude. This may be due to the structure of channels and river beds in the SRW. Most sites in the SRW are long runs, rather than combinations of riffles and pools, and the former tend to have lower flow velocity. Most of our measurements and sampling were in the runs of river reaches in the SRW, and so our results do not indicate that the flows of sampling sites of the SRW were not as fast as those of other river systems.

Sites located on the right side of axis-1 reflect downstream characteristics, including low altitude, high proportion of urban land use, and very high concentrations of ions, organic matter, and nutrients. Sites clustered in Group 2a were almost exclusively confined to the YSW region, a region that is nutrient rich and has other downstream properties.

Although our results indicate a downstream gradient of diatom composition, further studies are needed to describe the distribution patterns in relation to physiographic

characteristics (*e.g.*, ecoregions). Such studies require a greater number of study sites in particular regions. For example, some of our groups (*e.g.*, Groups 1d and 4) had wide geographic distributions across river systems, and the site variation was large in the NRW (Fig. 3). The reason for this pattern is not clear, but the use of more study sites in the NRW would provide a better understanding of such scattered distributions. Although nutrient enrichment is generally higher in downstream regions, our results clearly show that a gradient from unpolluted to polluted rivers is very important in structuring benthic diatom assemblages. In particular, our CCA showed a response of diatom communities to conductivity, BOD, and nutrients. Previous research has identified conductivity as an important factor in determining the geographic variation of diatom community structure in various rivers (Potapova and Charles, 2003; Soininen *et al.*, 2004; Bona *et al.*, 2007). Future studies could provide clearer evidence of the role of conductivity in diatom distribution by considering individual ions in multivariate analyses. An increase in ionic content very often accompanies organic and nutrient enrichment (Leland and Porter, 2000; Potapova and Charles, 2002; Leira and Sabater, 2005).

The present study also indicated the potential importance of flow regimes in shaping the structure of benthic diatom assemblages. Our CCA indicated that current velocity was the most important factor separating groups along axis-2. Flood frequency and water velocity have been reported as significantly affecting species composition and biomass development of periphytic algae in streams (Petersen, 1996; Clausen and Biggs, 1997). However, velocity effects seem to vary with the trophic status of streams, probably due to a negative association between biomass accumulation and detachment (which increases with current velocity) (Petersen and Stevenson, 1990; Biggs and Hickey, 1994). The effect of velocity on benthic diatom community structure is not as clear as it is for filamentous benthic algae. However, as benthic diatoms (*e.g.*, stalked forms) often occur in filamentous algal mats, and as diatom structure varies with substratum type (Leland and Porter, 2000), velocity may also play an important role in benthic diatom community structure.

Our results indicated that both *A. convergens* and *A. minutissima* were the most common diatom species in Korean rivers and streams (Fig. 1). These tightly adherent prostrate species are generally considered to be pioneer species (Barbour *et al.*, 1999; Kwon and Lee, 2007) that typically dominate in streams with *Cocconeis* spp. and small *Navicula* spp. after severe scouring events (Leland and Porter, 2000). We believe that the predominance of these taxa in most Korean rivers and streams is related to the monsoon climate pattern of Korea, in which typhoons and consequent severe hydrological scouring events occur regularly during the summer (June–August). The influence of stream geomorphology on diatom distribution is noteworthy, as it affects the physical habitats of diatoms. Pan *et al.* (2006) showed that dominant diatom assemblages in central valley streams of California (USA) are mainly affected by channel morphology, instream habitat,

and riparian conditions; however, they showed a weak association of diatom assemblages with water chemistry, indicating that diatoms can also be used as indicators of alterations of the physical habitat. We also expect that grazing pressure may affect stream diatom community composition (Koetsier, 2005; Chessman *et al.*, 2009), because benthic diatoms are the principal food source of many aquatic herbivores, such as insects and snails.

Spatial patterns of diatom distributions: implications for biological water quality assessment

This study showed that there is a considerable spatial variation in benthic diatom assemblages in Korea. Although part of this variation may result from spatially structured parameters that were not measured or from the relatively small number of rivers investigated in some regions (*e.g.*, the NRW), some diatom taxa indicative of Groups 1c and 3 (*e.g.*, *G. pumilum* and *C. placentula*) had regionally restricted patterns (Fig. 6). This indicates that the distribution of some taxa is better explained by spatial parameters than environmental parameters. Even if there were significant correlations of environmental parameters with indicator diatom taxa of Groups 1c and 3, the overall environmental effect on these groups was relatively small relative to other groups (Table 4). Thus, other parameters, such as spatial factors, may also be involved. An exception is the diatoms in Group 4, which had scattered nationwide distributions and were not clearly distinguishable from other groups (Figs. 3 and 5). However, none of the diatom taxa that played important roles in river benthic diatom communities were limited to one region; variation in their dominance among regions may be related to overriding local parameters (Pan *et al.*, 1999; Soininen *et al.*, 2004).

The spatial variation of river benthic diatom communities is due to complicated multi-scale effects (*e.g.*, Stevenson, 1997), with many interrelated factors (Leland and Porter, 2000; Potapova and Charles, 2002). Higher-level parameters (*e.g.*, climate, geomorphology, vegetation, and land use), which are basic attributes of ecoregion classification, are likely to operate at the watershed and regional levels, imposing constraints on the local biotic interactions in streams. The ecoregion concept was developed to describe landscape characteristics that influence regional patterns of aquatic and terrestrial resources (Omernik, 1987). Subsequently, many studies have used the ecoregion concept to interpret regional biological phenomena and ecosystem function (Hughes and Larsen, 1988; Roth *et al.*, 1996; Butcher *et al.*, 2003; Simbora *et al.*, 2005; Borja *et al.*, 2007). However, there is often less correspondence between spatial patterns of biological communities and ecoregional classification (Whittier *et al.*, 1988; Pan *et al.*, 1999; Potapova and Charles, 2002), probably because of the greater impact of local geomorphic factors (*e.g.*, habitat alteration and destruction, channel straightening, and river bed modification) and disturbance factors (*e.g.*, pollution), which interact in a complicated

manner (Biggs *et al.*, 1990). The spatial distribution of benthic river diatoms that we found in the present study is worthy of further investigations in relation to ecoregion classification. The South Korean landscape is divided into 16 ecoregions (Shin and Lee, 2004). At present, the correspondence between diatom spatial patterns and this ecoregional classification is unclear. However, our initial comparison of these relationships suggests a correlation in some regions, including the HRW, YRW, and SRW, where sites of Groups 1b, 1c, 2a, and 3 are located (data not shown).

Potapova and Charles (2002) have described other potential causes of spatial variation of benthic river diatoms. Although many previous studies have investigated the association of various environmental variables with the spatial variation of diatoms, it is difficult to include all possibly related parameters in one study. Future carefully designed studies of benthic river diatom communities are necessary to better understand the relationships between the spatial distribution of diatoms and the biological assessment of water quality. Given the evidence for spatial patterns in diatom community composition, a biological assessment program that uses benthic diatoms would benefit from a better understanding of diatom geographical distribution. For example, some indicator species that are predominant in particular geographical regions may help to identify space-specific reference sites. However, local instream factors that affect environmental and disturbance conditions are often more important than spatial factors in explaining diatom distribution (*e.g.*, Leland and Porter, 2000; Potapova and Charles, 2002). Thus, a combination of regional discrimination and empirical modeling based on local environmental features might provide the most robust framework for diatom-based assessment of the biological integrity of streams and rivers. This strategy has been applied to other stream biota, including macroinvertebrates and fish. However, the degree of variation among different trophic levels may depend on the specific environmental and habitat conditions to which they respond (Roth *et al.*, 1996; Passy *et al.*, 2004). Thus, bioassessment of running waters should benefit from analysis of multiple taxonomic groups.

Conclusions

This study was the first large-scale investigation of benthic diatoms in Korea. We sampled 720 sites from the five major river systems and their tributaries and found a large number of benthic diatom taxa. Most species were rare and only 15% of taxa accounted for more than 5% of the diatoms at individual sites. Multivariate analysis allowed assessment of the spatial patterns and ecological determinants of benthic diatom assemblages. CCA indicated that site classification using groups of benthic diatom assemblages was not strongly discriminatory, as indicated by low eigenvalues and low total variance in species data. This was probably because of parameters

that we did not measure, and unresolved problems in taxonomic identification associated with such a large study. However, diatom–environmental factor correlations were high, indicating a strong relationship between diatoms and measured environmental variables. Our results show that multi-scale factors are important in explaining most of the variation in benthic diatom assemblages at the national scale, but that a “downstream gradient” was evident, with significant changes occurring from fast-flowing and mostly oligotrophic rivers in mountainous areas to slow-flowing eutrophic lowland rivers. Our results also suggest that a gradient of water mineral content (indicated by conductivity) can affect diatom community structure. However, because a downstream gradient is largely associated with eutrophication (nutrient enrichment), it was difficult to identify individual effects in each gradient. We suggest that future studies on this topic investigate the contribution of individual ions to the conductivity gradient. The information obtained in the present study on the spatial distribution of benthic diatom indicator species and their significance in relation to environmental variables contributes to development of a larger program for assessment of the biological integrity of lotic ecosystems.

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Appendix 1. Diatom taxa ($n = 126$) that occurred in the greatest density at more than one site among the five major river watersheds in Korea. Asterisks indicate abundance. *Presence of taxa regardless of their abundance. **Dominant taxa ($> 10\%$ in overall abundance).

Diatom taxa	Han River	Nakdong River	Geum River	Youngsan River	Seomjin River
Order Centrales					
Suborder Coscinodiscineae					
Family Thalassiosiraceae					
<i>Aulacoseira alpigena</i> (Grunow) Krammer	*		**	*	
<i>Aulacoseira granulata</i> (Ehrenberg) Simonsen	**	*	*	*	
<i>Cyclotella atomus</i> Hustedt	*	**	**	**	
<i>Cyclotella comta</i> (Ehrenberg) Kützing	**		*	*	
<i>Cyclotella meneghiniana</i> Kützing	**	*	**	**	*
<i>Cyclotella stelligera</i> (Cleve & Grunow) Van Heurck	*	*	**	**	
<i>Cyclotella</i> spp.				**	
<i>Stephanodiscus alpinus</i> Hustedt	**		*		
Family Melosiraceae					
<i>Melosira varians</i> C. Agardh	**	*	*	**	**
<i>Stephanopyxis</i> spp.					**
Family Hemidiscaceae					
<i>Actinocyclus normanii</i> (W. Gregory) Hustedt	*	**	*		
Order Pennales					
Suborder Araphidineae					
Family Fragilariaceae					
<i>Diatoma hiemale</i> var. <i>quadratum</i> (Kützing) R. Ross	**			**	
<i>Diatoma vulgare</i> Bory	**	*	*	*	
<i>Diatoma</i> spp.	**			*	
<i>Fragilaria capitata</i> (Ehrenberg) Lange-Bertalot				**	
<i>Fragilaria capucina</i> Desmazières	**	*	*	*	**
<i>Fragilaria capucina</i> var. <i>capitellata</i> (Grunow) Lange-Bertalot	*		*	**	
<i>Fragilaria capucina</i> var. <i>mesolepta</i> (Rabenhorst) Rabenhorst	**	*	*	*	**
<i>Fragilaria construens</i> (Ehrenberg) Grunow	**		*	*	
<i>F. construens</i> f. <i>binodis</i> (Ehrenberg) Hustedt	*	*	**	*	*
<i>F. construens</i> f. <i>venter</i> (Ehrenberg) Hustedt	**	**	*	*	**
<i>Fragilaria crotonensis</i> Kitton	**	*	*	*	
<i>Fragilaria elliptica</i> Schumann	*	**	**		
<i>Fragilaria vaucheriae</i> (Kützing) J. B. Petersen	**		*	*	
<i>Fragilaria vaucheriae</i> var. <i>capitellata</i> (Grunow) R. M. Patrick			*	**	
<i>Fragilaria</i> spp.			*		**
<i>Hannaea arcus</i> (Ehrenberg) R. M. Patrick	**			*	
<i>Synedra acus</i> Kützing	**	*	**	*	*
<i>Synedra fasciculata</i> (Kützing) Grunow	*	**	*	*	
<i>Synedra inaequalis</i> H. Kobayasi	**	*	*	*	
<i>Synedra ulna</i> (Nitzsch) Ehrenberg	*	**	**	**	*
<i>Synedra ulna</i> var. <i>contracta</i> Østrup			*	**	
Suborder Raphidineae					
Family Achnantheae					
<i>Achnanthes alteragracillima</i> Lange-Bertalot	**		**	*	
<i>Achnanthes amoena</i> Hustedt	**	*			
<i>Achnanthes biasoletiana</i> (Kützing) Grunow	**	**	*	*	
<i>Achnanthes bioretii</i> Germain	**				
<i>Achnanthes brevipes</i> C. Agardh	*		**		
<i>Achnanthes catenata</i> Bily & Marvan					**
<i>Achnanthes conspicua</i> A. Mayer				**	
<i>Achnanthes convergens</i> H. Kobayasi	**	**	**	**	*

Appendix 1. Continued.

Diatom taxa	Han River	Nakdong River	Geum River	Youngsan River	Seomjin River
<i>Achnanthes delicatula</i> ssp. <i>engelbrechtii</i> (Choln.) Lange-Bertalot				**	
<i>Achnanthes exigua</i> Grunow	*	*	**	**	*
<i>Achnanthes impexa</i> Lange-Bertalot		*	**		
<i>Achnanthes laevis</i> Østrup	**			*	
<i>Achnanthes lanceolata</i> (Brébisson) Grunow	**	**	*	*	*
<i>Achnanthes lanceolata</i> ssp. <i>dubia</i> (Grunow) Lange-Bertalot	**	*		*	
<i>Achnanthes microcephala</i> (Kützing) Grunow				**	
<i>Achnanthes minutissima</i> Kützing	**	**	**	*	**
<i>Achnanthes minutissima</i> var. <i>saprophila</i> Kobayasi & Mayama		*	*	**	
<i>Achnanthes minutissima</i> var. <i>scotica</i> Kützing				**	
<i>Achnanthes subhudsonis</i> Hustedt	*	*	*	**	
<i>Achnanthes</i> spp.	*	*	*	*	**
<i>Cocconeis placentula</i> Ehrenberg	**	*		*	**
<i>Cocconeis placentula</i> var. <i>euglypta</i> (Ehrenberg) Grunow	**	**		**	*
<i>Cocconeis placentula</i> var. <i>lineata</i> (Ehrenberg) Van Heurck	**	**	**	*	**
<i>Cocconeis</i> spp.	*			*	**
Family Naviculaceae					
<i>Amphora ovalis</i> (Kützing) Kützing	**		*	*	
<i>Amphora</i> spp.	**		*	*	*
<i>Caloneis bacillum</i> (Grunow) Cleve	**	*	*	*	
<i>Cymbella affinis</i> Kützing	**	*	**	*	**
<i>Cymbella amphicephala</i> Nägeli	*			**	
<i>Cymbella cinuata</i> W. Gregory	**	**	*		
<i>Cymbella cistula</i> (Hemprich & Ehrenberg) O. Kirchner	**			*	*
<i>Cymbella lacustris</i> (C. Agardh) Cleve	**	*	*	*	
<i>Cymbella minuta</i> Hilse ex Rabenhorst	*	*	**	*	*
<i>Cymbella silesiaca</i> Bleisch	**	*	**	*	*
<i>Cymbella tumida</i> (Brébisson) Van Heurck	**	*	*	*	*
<i>Cymbella turgidula</i> Grunow	**	*	*	*	*
<i>Cymbella</i> spp.			*	**	
<i>Gomphonema lagenula</i> Kützing	**				
<i>Gomphonema angustum</i> C. Agardh	**	*	*	*	*
<i>Gomphonema clavatum</i> Ehrenberg	**		*	*	*
<i>Gomphonema clevei</i> Fricke	**	**	**	*	
<i>Gomphonema dichotomum</i> Kützing				*	**
<i>Gomphonema herculeana</i> Ehrenberg			*	**	
<i>Gomphonema insigne</i> Gregory	*			**	
<i>Gomphonema minutum</i> (C. Agardh) C. Agardh	**	**	*		
<i>Gomphonema olivaceum</i> (Hornemann) Brébisson	**			**	
<i>Gomphonema parvulum</i> Kützing	**	*	**	**	**
<i>Gomphonema pumilum</i> (Grunow) Reichardt & Lange-Bertalot	*	*	*	**	**
<i>Gomphonema truncatum</i> Ehrenberg	**	*	*	*	*
<i>Navicula cincta</i> (Ehrenberg) Kützing	*	*	*	*	**
<i>Navicula contenta</i> Grunow			**	*	
<i>Navicula cryptocephala</i> Kützing	**	*	**	*	*
<i>Navicula cryptotenella</i> Lange-Bertalot	**	**	*	*	*
<i>Navicula elginensis</i> var. <i>cuneata</i> (M. Moller ex. Foged) Lange-Bertalot				**	
<i>Navicula goeppertiana</i> (Bleisch) H. L. Smith	**	**	**	*	**
<i>Navicula minima</i> Grunow	**	**	**	**	
<i>Navicula mutica</i> var. <i>ventricosa</i> (Kützing) Cleve & Grunow		*	*	**	
<i>Navicula neoventricosa</i> Hustedt		**		*	
<i>Navicula nipponica</i> (Skvortzow) Lange-Bertalot	*			**	*
<i>Navicula nivalis</i> Ehrenberg	*		**	**	
<i>Navicula notha</i> Wallace	*	*	*	**	*
<i>Navicula novasiberica</i> Lange-Bertalot				**	
<i>Navicula perminuta</i> Grunow	*	*	*	**	
<i>Navicula pupula</i> Kützing	*	*	**	**	*
<i>Navicula recens</i> (Lange-Bertalot) Lange-Bertalot		**			
<i>Navicula saprophila</i> Lange-Bertalot & Bonik	**		*	**	
<i>Navicula schroeteri</i> var. <i>symmetrica</i> (Patrick) Lange-Bertalot			**		
<i>Navicula seminulum</i> Grunow	**	**	*	*	*
<i>Navicula subminuscula</i> Manguin	**	**	**	*	*

Appendix 1. Continued.

Diatom taxa	Han River	Nakdong River	Geum River	Youngsan River	Seomjin River
<i>Navicula subtilissima</i> Cleve		*	*	**	
<i>Navicula tenelloides</i> Hustedt				**	
<i>Navicula viridula</i> (Kützing) Ehrenberg	*		*	*	**
<i>Navicula</i> spp.	*	*	**	**	**
<i>Neidium dubium</i> (Ehrenberg) Cleve	**				
<i>Pinnularia divergens</i> W. Smith	**				
<i>Pinnularia gibba</i> Ehrenberg	**	*	*	*	*
<i>Rhicosphenia abbreviate</i> (C. Agardh) Lange-Bertalot	*	*	*	**	**
<i>Stauroneis anceps</i> Ehrenberg	**			*	*
Family Epithemiaceae					
<i>Epithemia adnata</i> (Kützing) Brébisson	*				**
Family Bacillariaceae					
<i>Denticula tenuis</i> Kützing	**				
<i>Nitzschia acicularis</i> (Kützing) W. Smith	*	*	**		
<i>Nitzschia amphibia</i> Grunow	**	*	**	**	*
<i>Nitzschia capitellata</i> Hustedt	**	*		*	
<i>Nitzschia diversa</i> Hustedt			**	*	
<i>Nitzschia filiformis</i> (W. Smith) Van Heurck	*	*	**	*	
<i>Nitzschia fonticola</i> (Grunow) Grunow	*	**	**	**	
<i>Nitzschia fossilis</i> Grunow	**				
<i>Nitzschia frustulum</i> (Kützing) Grunow	**	*	*	*	
<i>Nitzschia gracilis</i> Hantzsch	*	**	*	**	**
<i>Nitzschia inconspicua</i> Grunow	**	**	**	**	
<i>Nitzschia intermedia</i> Hantzsch ex Cleve & Grunow	*	*	*	**	
<i>Nitzschia palea</i> (Kützing) W. Smith	**	**	**	**	**
<i>Nitzschia</i> spp.	*	*	**	*	*
Family Surirellaceae					
<i>Surirella minuta</i> Kützing	*	*	**	*	