



March 2012

GAHCHO KUÉ PROJECT

2011 Lower Trophic Organisms Supplemental Monitoring Report

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REPORT



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1.0 INTRODUCTION

De Beers Canada Inc. (De Beers) is proposing to develop the Gahcho Kué Project (Project), a diamond mine in the Northwest Territories (NWT). The Project is located in the North Slave region of the NWT at Kennedy Lake, approximately 140 kilometres (km) northeast of Łutsek'e and 280 km northeast of Yellowknife.

Baseline studies have been conducted to support the Environmental Impact Assessment (EIS) for the Project and the Environmental Impact Review (EIR) Process. These data were reported in the December 2010 EIS (De Beers 2010a). Baseline data reported in the 2010 EIS are sufficient to support the environmental assessment within the EIS. However, De Beers is committed to ongoing data collection in advance of regulatory approval of and the permitting process for the Project. As such, supplemental baseline data have been collected in 2011, and will continue to be collected and reported annually, until such time that these activities are no longer required prior to Project construction or evolve into future monitoring programs associated with an approved Project.

The purpose of collecting and reporting the supplemental baseline data for the Project is to support a consistent and transparent baseline program. In general, the goals of the supplemental data collection are to:

- reduce uncertainty and increase the level of confidence in impact predictions;
- broaden the baseline areas of investigation; and
- contribute to long-term future monitoring and adaptive management of the Project.

The focus of the 2011 supplemental data collection reported herein is lower trophic communities (e.g., phytoplankton, zooplankton, and benthic invertebrates). The purpose of this report is to provide supplemental baseline information on the lower trophic resources in the area of the Project. This report supplements the data presented in the EIS (De Beers 2010a, Annex J [Fisheries and Aquatic Resources Baseline] and Addendum JJ [Additional Fish and Aquatic Resources Baseline Information]). The supplemental information presented in this report was collected in late summer 2011 to enhance the existing baseline data (e.g., 2004, 2005, 2007, and 2010).

“Plankton” is a general term referring to small, usually microscopic organisms that live suspended in the water. For the purpose of this study, the term “phytoplankton” refers to the algal component of plankton and includes the following seven major taxonomic groups:

- cyanobacteria;
- Chlorophyceae (chlorophytes);
- Chrysophyceae (chrysophytes);
- Cryptophyceae (cryptophytes);
- Bacillariophyceae (diatoms);
- Dinophyceae (dinoflagellates); and
- Euglenophyceae (euglenoids).



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The term “zooplankton” refers to microscopic animals and includes Rotifera (rotifers) and crustaceans, specifically Cladocera (cladocerans or water fleas), Cyclopoida (cyclopoid copepods), and Calanoida (calanoid copepods). Cyclopoid and calanoid copepods are considered separately because of taxonomic and ecological differences. Calanoids are typically herbivorous, feeding on phytoplankton; whereas cyclopoids are typically omnivorous, feeding on phytoplankton and small zooplankton (Brönmark and Hansson 1998). Additionally, calanoids are almost exclusively pelagic (i.e., open-water), while cyclopoids are dominated by littoral (i.e., near-shore) species, although a few pelagic species of cyclopoids can account for a major component of the planktonic community.

Benthic invertebrates are small aquatic animals that lack backbones; they live on the bottoms of waterbodies such as lakes and streams. Freshwater benthic invertebrates include mostly insect larvae, crustaceans, worms, leeches, snails, and clams. They form diverse communities often consisting of thousands of individuals per square metre. Benthic invertebrates live on the surface of the sediments or burrow into sediments, although some species are closely associated with aquatic plants. They are frequently sampled to monitor the environmental quality of lakes for the following reasons (Rosenberg and Resh 1993):

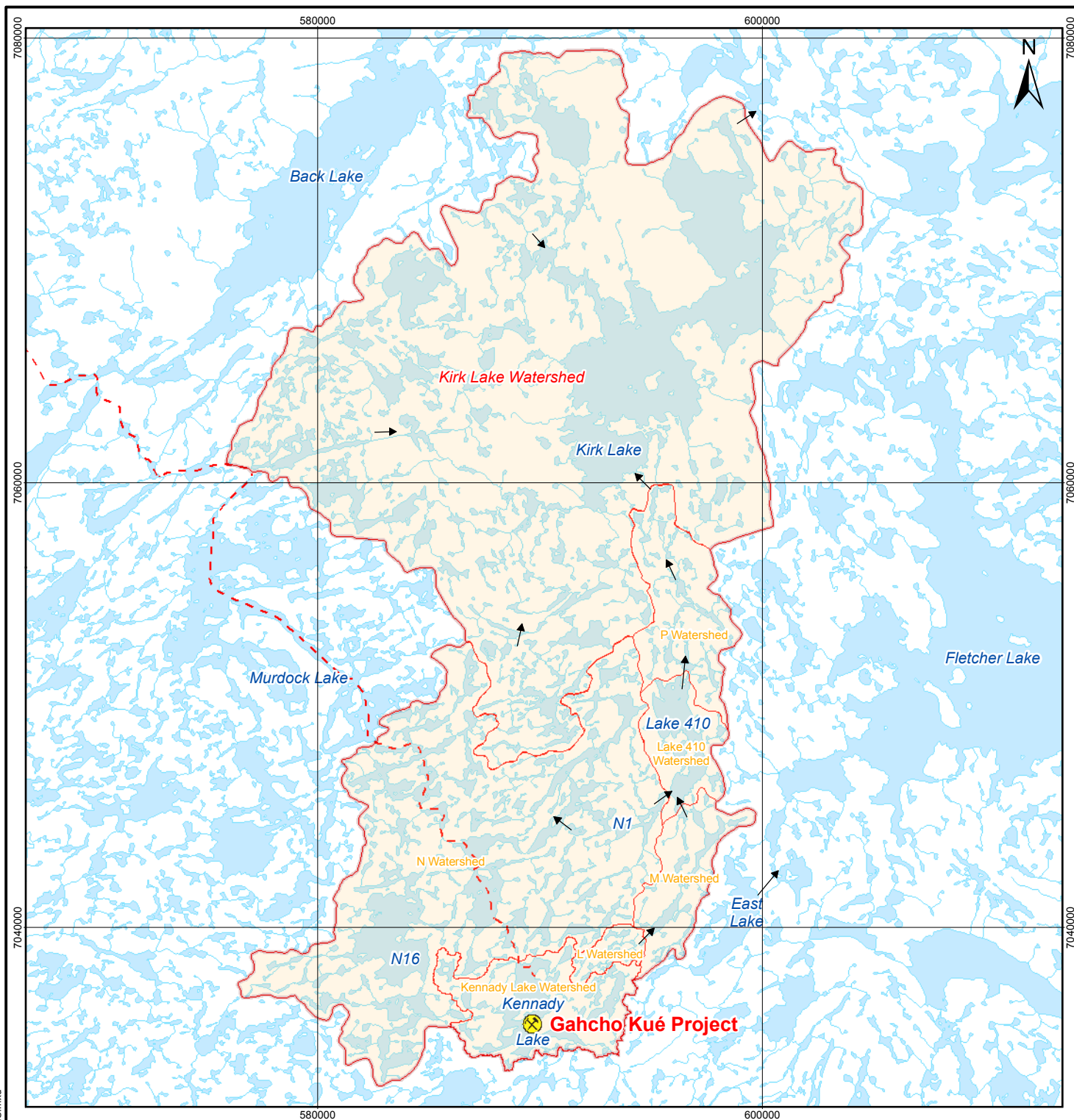
- they are present in nearly all waterbodies and are usually abundant;
- they remain in a small area throughout the aquatic phase of their life cycle;
- they obtain food by various means, including the filtering of fine particulates and feeding on algae, decaying organic material, aquatic plants, or other invertebrates;
- they have relatively long life cycles ranging from months to years, thereby integrating the effects of disturbances over a relatively long period;
- they are an important food source for organisms at higher trophic levels such as fish;
- they are sensitive to a large variety of disturbances, including the addition of sediment, toxins, nutrients, and organic material; low dissolved oxygen (DO) levels; and alteration of flow, substratum, and temperature;
- they respond to disturbances in a predictable manner;
- they can be relatively easily collected and identified; and,
- the wide range of species inhabiting any given location assures that animals of varying sensitivity are present.

This report describes results of plankton and benthic invertebrate sampling in August 2011, with a focus on deep open-water stations in East Lake (Reference Lake) and Lake N11, and open water locations in the L and M lakes (Lakes L2, M1, M2, M3, and M4).



2.0 STUDY AREAS

The LSA is a 739 square kilometres (km²) area that includes the watersheds of the lakes and streams that may be directly affected by the Project (Figure 1). The regional study area (RSA) was defined as the Lockhart River watershed. The 2011 lower trophic field program was conducted within the local study area (LSA), with the exception of East Lake (Reference Lake), which is located outside of the LSA, but within the RSA. The study areas are described in Annex J, Section J2 of the 2010 EIS (De Beers 2010a).



LEGEND

- Gahcho Kué Project
- Winter Access Road
- Watercourse
- Waterbody
- K1 Lake Identifier
- Watershed Boundary
- Local Study Area
- Flow Direction

NOTES

Base data source: National Topographic Base Data (NTDB) 1:250,000

GAHCHO KUÉ PROJECT

Local Study Area

PROJECTION: UTM Zone 12
DATUM: NAD83

Scale: 1:250,000
5 2.5 0 5
Kilometres

FILE No:
B2011-Fish-020-GIS

DATE:
January 27, 2012

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Figure 1



3.0 METHODS

This section summarizes the methods used during the 2011 lower trophic supplemental baseline field programs.

3.1 Lower Trophic Communities

3.1.1 Phytoplankton

Phytoplankton samples were collected in August 2011. A single location was sampled in Lakes M1, M2, M3, M4 and L2. Samples were also collected from five stations in Lake N11 and East Lake (Reference Lake; Table 1 and Figure 2). Discrete water samples were collected at 2 metre (m) intervals within the euphotic zone at each site, using a Kemmerer® water sampler. The water samples were mixed thoroughly and used to fill a 250 millilitre (mL) amber Nalgene® bottle at each site. The samples were preserved with 10 mL formalin acetic acid solution and 5 mL acid Lugol's solution. Samples were stored in the dark and shipped immediately for taxonomic identification and analysis.

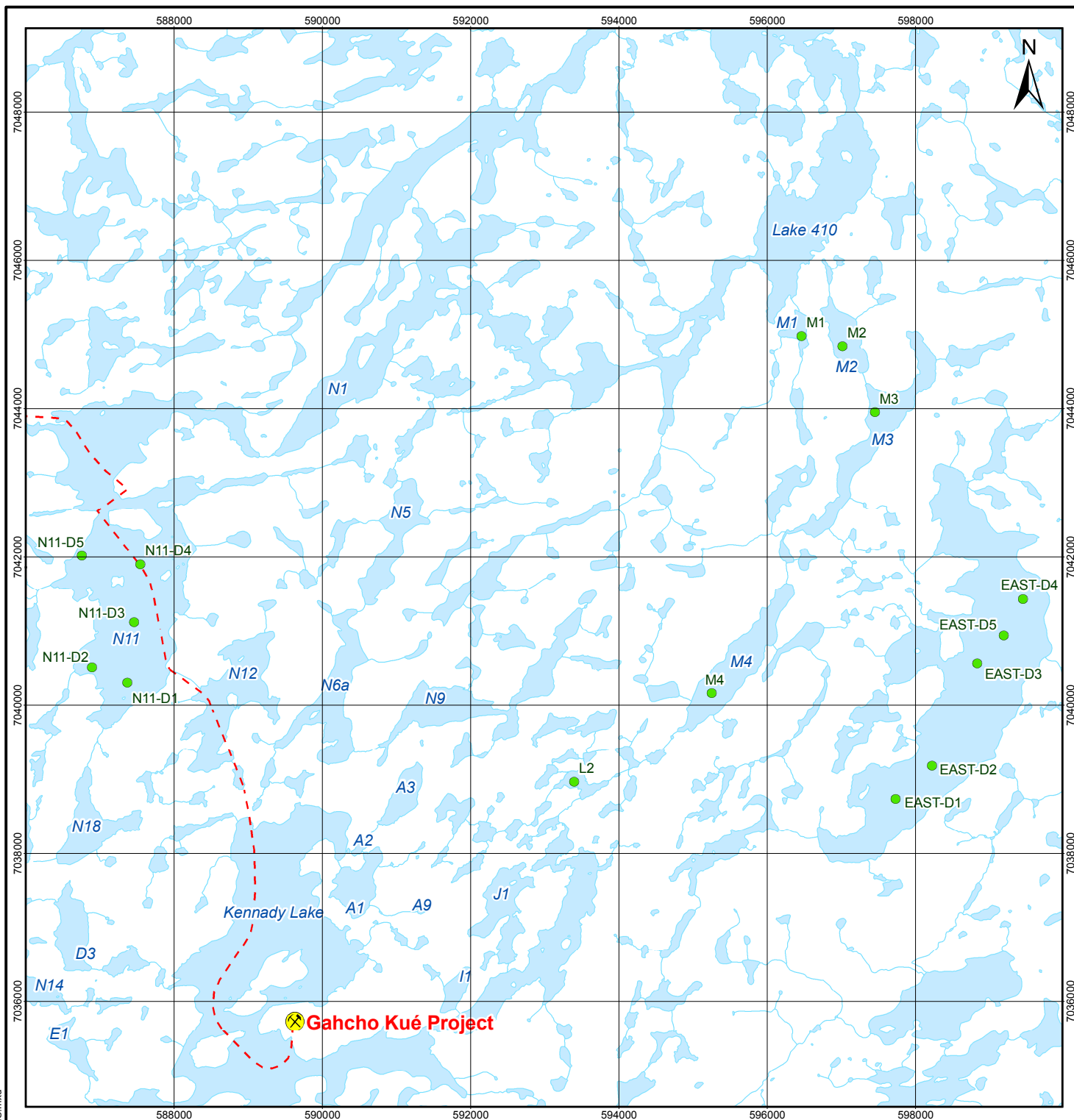
Secchi depths and limnological profiles (i.e., specific conductivity, dissolved oxygen, water temperature and pH) were also measured at each location using a YSI 600QS water meter (Appendix I, Table I-1).

Table 1 Phytoplankton Sampling Locations in Lakes, 2011

Watershed	Lake/Station	UTM Coordinates 12V NAD 83	
		Easting	Northing
M	M1	596466	7045014
	M2	597001	7044838
	M3	597425	7043953
	M4	595226	7040172
L	L2	593390	7038966
N	N11-D1	587363	7040300
	N11-D2	586942	7040514
	N11-D3	587460	7041112
	N11-D4	587546	7041905
	N11-D5	586853	7041956
East Lake (Reference Lake)	Ref Lake-D1	597734	7038738
	Ref Lake-D2	598219	7039180
	Ref Lake-D3	598832	7040567
	Ref Lake-D4	599424	7041434
	Ref Lake-D5	599198	7040935

UTM = Universal Transverse Mercator; NAD = North American Datum.

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LEGEND

- Gahcho Kué Project
- Winter Access Road
- Watercourse
- Waterbody
- Phytoplankton and Zooplankton Sampling Location

NOTES

Base data source: National Topographic Base Data (NTDB) 1:50,000

GAHCHO KUÉ PROJECT

Phytoplankton and Zooplankton Sampling Locations, 2011

PROJECTION:

UTM Zone 12

DATUM:

NAD83

Scale: 1:75,000

1,000 500 0 1,000

Metres



FILE NO:

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Figure 2



Phytoplankton samples were analyzed for taxonomic composition, abundance and biomass by Bio-Limno Research and Consulting Inc. (Bio-Limno), Halifax, Nova Scotia. Aliquots of 7 mL of the preserved phytoplankton samples were allowed to settle overnight in sedimentation chambers following the procedure of Lund et al. (1958). Algal units were counted from randomly selected transects on a Zeiss Axiovert 40 CFL inverted microscope. Counting units were individual cells, filaments, or colonies depending on the organization of the algae. A minimum of 400 units were counted for each sample. The majority of the samples were analyzed at 500 times magnification (500x), with initial scanning for large and rare organisms (e.g., *Ceratium* sp.) completed at 250x. Taxonomic identifications were based primarily on Geitler (1932); Skuja (1949); Findlay and Kling (1976); Anton and Duthie (1981); Huber-Pestalozzi (1961, 1972, 1982, 1983); Tikkanen (1986); Prescott (1982); Whitford and Schumacher (1984); Starmach (1985); Krammer and Lange-Bertalot (1986, 1988, 1991a,b); Komárek and Anagnostidis (1998a,b, 2005); and Wehr and Sheath (2003).

Fresh weight biomass was calculated from recorded abundance and specific biovolume estimates based on geometric solids (Rott 1981), assuming a specific gravity of 1 gram per cubic centimetre (g/cm^3). The biovolume (cubic millimetres per cubic metre [mm^3/m^3] wet weight) of each species was estimated from the average dimensions of 10 to 15 individuals. The biovolumes of colonial taxa were based on the number of individuals within each colony. All calculations for cell concentration and biomass were performed with Hamilton's (1990) computer program.

3.1.1.1 Data Analysis

Phytoplankton data were summarized as total taxonomic richness, abundance and biomass, and taxonomic richness, abundance and biomass of the major taxonomic groups:

- Chlorophyceae;
- Chrysophyceae;
- cyanobacteria;
- Cryptophyceae;
- Bacillariophyceae;
- Euglenophyceae; and
- Dinophyceae.

Community composition was summarized as relative abundance and biomass of each of the major taxonomic groups.

3.1.2 Zooplankton

Zooplankton samples were collected from the same waterbodies and stations as the phytoplankton samples (Table 1 and Figure 2). Five samples were collected at each site using a 25 centimetre (cm) diameter, 73 micron (μm) mesh plankton Wisconsin® net. The net was lowered to 1 m above the lake bottom allowing collection of the full water column. If the sites were too shallow for a full water column vertical haul, horizontal tows were completed. Haul or tow depths and/or lengths were recorded for each sample (Table 2) and were used to calculate the volume of water filtered through the net. Filtering efficiency was assumed to be 100percent



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(%), based on the low productivity in the lakes sampled, which was expected to result in low suspended sediment concentrations.

The 250-mL sample bottles were filled with 125 mL of sample and preserved with one half of an Alka-Seltzer tablet to avoid shock or contortion of the zooplankters and then with 125 mL sugar formalin. Samples were stored in the dark and shipped immediately for taxonomic identification.

Table 2 Zooplankton Haul Depths in each Lake, August 2011

Watershed	Lake/Station	Depth/Length of Haul [m]
M	M1	3.5
	M2	4
	M3	6
	M4	7
L	L2	3
N	N11-D1	5.5
	N11-D2	5
	N11-D3	5
	N11-D4	5
	N11-D5	5
East Lake (Reference Lake)	Ref Lake-D1	5.9
	Ref Lake-D2	13
	Ref Lake-D3	14
	Ref Lake-D4	8
	Ref Lake-D5	5

Zooplankton samples were analyzed for abundance and biomass of crustaceans and rotifers by Salki Consultants Inc., Winnipeg, Manitoba. Each sample underwent three levels of analysis, as follows:

- 1/40 or 1/80 of each sample was examined under a compound microscope at 63x to 160x, and all specimens of crustaceans and rotifers were identified to the lowest taxonomic level (typically species) and assigned to size categories;
- a second sub-sample, representing 11 % of the sample volume, was examined under a stereoscope at 12x for the large species (i.e., *Heterocope septentrionales*, *Holopedium gibberum*, and *Daphnia middendorffiana*) and rare species, which were enumerated and assigned to size categories; and
- the entire sample was examined under the stereoscope to improve abundance estimates for the largest species (i.e., adult male and female *Heterocope septentrionales*, *Holopedium gibberum*, and *Daphnia middendorffiana*).

All Cyclopoida and Calanoida specimens (mature and immature) were identified to the species level, with the exception of nauplii, which were classified as either Calanoida or Cyclopoida. All Cladocera were identified to the species level. Rotifers were identified to genus. Zooplankton abundance was reported as individuals per litre (ind./L). Taxonomic identifications were based primarily on Brooks (1957), Wilson (1959), and Yeatman (1959).



Biomass estimates for each taxon were obtained using mean adult sizes determined during the analysis of the zooplankton samples and length-weight regression equations developed by Malley et al. (1989). Additional measurements were made on newly encountered species and to validate consistency of adult sizes. Zooplankton biomass was reported as milligrams (wet weight) per cubic metre (mg/m^3). Wet weights were converted to dry weight by assuming that dry weight equals 7% of wet weight, based on the results of Malley et al. (1989) (Appendix I Table I-7).

3.1.2.1 Data Analysis

Zooplankton data were summarized as total taxonomic richness, abundance and biomass, and taxonomic richness, abundance, and biomass of the major taxonomic groups:

- Calanoida
- Cyclopoida
- Cladocera
- Rotifera

Community composition was summarized as relative abundance and biomass of the major taxonomic groups.

3.1.3 Quality Control

Seven samples, accounting for approximately 10% of the total number of zooplankton samples, were re-counted by the same taxonomist to verify counting efficiency (Appendix I Table I-6). The same procedure was not performed for phytoplankton samples collected in 2011, but is planned for 2012. Both the zooplankton and phytoplankton data were entered into electronic format by the taxonomist that did the counts and were double-checked upon entry; errors were corrected as necessary before transferring the electronic files.

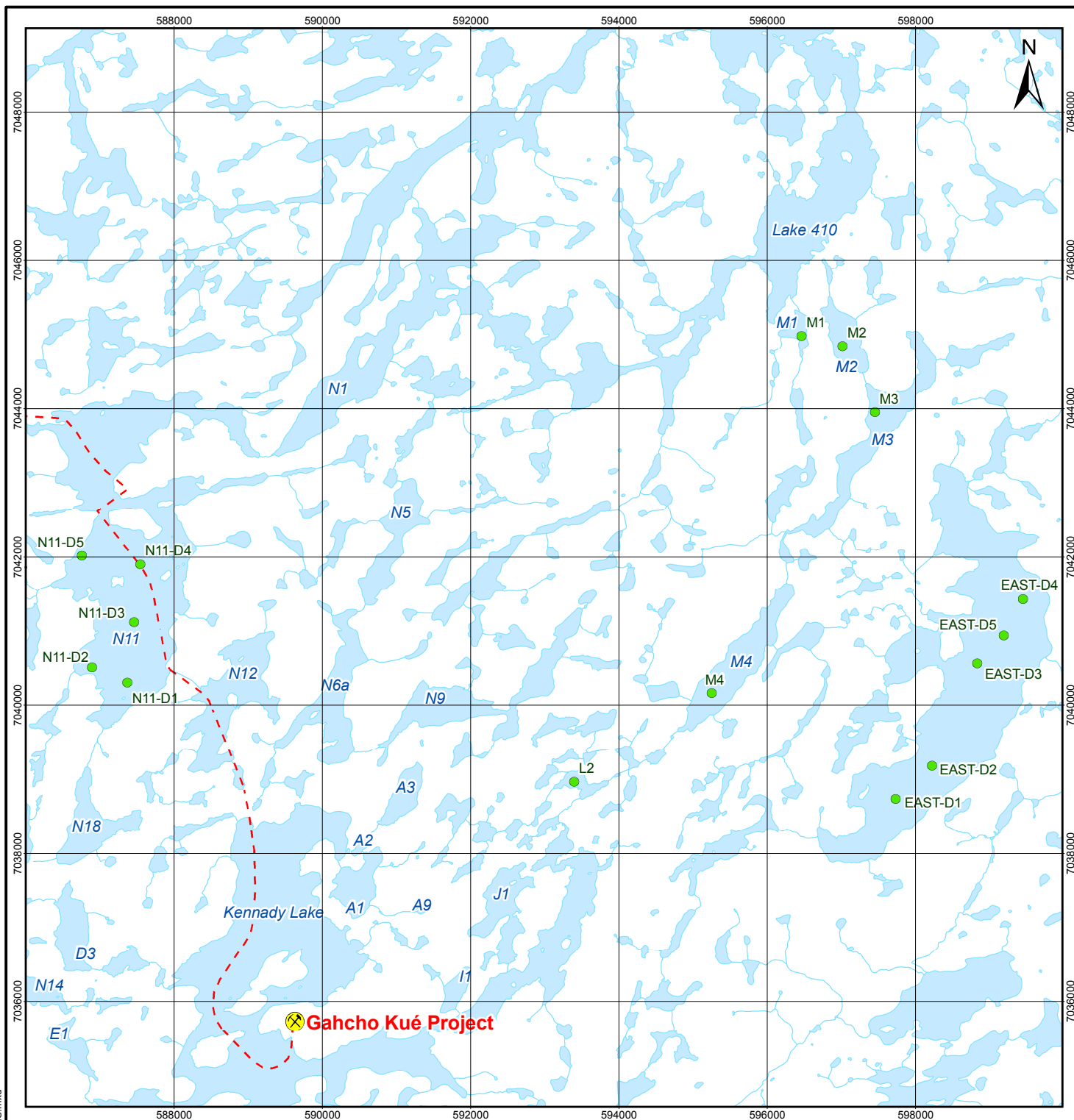
3.1.4 Benthic Invertebrates

3.1.4.1 2011 Supplemental Sampling

In 2011, benthic invertebrate sampling was conducted to enhance available baseline data, including information on among-station variation in lakes within a habitat type, reference lake data, and information for the chain of L and M lakes located downstream of Kennady Lake.

Benthic invertebrate samples were collected in the open water areas of East (Reference) Lake, Lake N11 and the L and M Lakes (Lakes L2, M1, M2, M3, and M4) (Figure 3). In East Lake and Lake N11, five benthic invertebrate stations were sampled in open-water areas (i.e., non-littoral areas). In the L and M Lakes, a single open-water station was sampled for benthic invertebrates. Five replicate samples were collected at each station.

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LEGEND

- Gahcho Kué Project
- Winter Access Road
- Watercourse
- Waterbody
- Benthic Invertebrate Sampling Location

NOTES

Base data source: National Topographic Base Data (NTDB) 1:50,000

GAHCHO KUÉ PROJECT

Benthic Invertebrate Sampling Locations, 2011

PROJECTION:

UTM Zone 12

DATUM:

NAD83

Scale: 1:75,000

1,000 500 0 1,000

Metres



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Figure 3



Field Methods

Benthic invertebrate samples were collected using a stainless steel Ekman Grab (15 x 15 x 15 cm) with a bottom sampling area of 0.0232 square metre (m²), where bottom sediments were suitable (i.e., fine-grained). Each sample was sieved through a sieve bucket with a 250 µm mesh bottom. The material retained in the bucket was placed into individually labelled 1 litre (L) polyethylene jars and preserved with 10% neutral buffered formalin. Samples were shipped to a qualified taxonomist (J. Zloty, Ph.D., Summerland, British Columbia) for taxonomic identification and enumeration of invertebrates.

At each station, a sediment sample was collected and sent to Maxxam Analytics for determination of sediment particle size, total organic carbon (TOC) and moisture content. However, sediment samples were not collected at M1, M2 and L2 during the August 2011 sampling.

During the benthic invertebrate survey, the following supporting environmental information was recorded at each sampling station:

- Sampling date and time;
- Weather conditions (air temperature, wind velocity and wind direction);
- Global positioning system (GPS) coordinates as universal transverse Mercator (UTM) for each station;
- Water depth (m); and
- Vertical profiles of water temperature (°C), dissolved oxygen (DO; milligrams per litre [mg/L]), pH, and specific conductivity (microSiemens per centimetre [µS/cm]) at 1 m intervals

Station UTM coordinates were recorded using a hand-held Garmin GPS unit. Temperature, dissolved oxygen concentration, and specific conductivity were measured at each benthic invertebrate sampling site with a YSI-556 multi-parameter meter.

Laboratory Methods

Samples were processed according to standard protocols based on recommendations in Environment Canada (2002) and Gibbons et al. (1993). Benthic invertebrate samples were first washed through a sieve with a 250 µm mesh opening to remove preservative and fine sediments remaining after field sieving. Organic material was separated from inorganic material using elutriation. Inorganic material was checked for any remaining shelled or cased benthic invertebrates, which were removed and added to the organic material. The organic material was split into coarse and fine fractions using a set of nested sieves of 1 millimetre (mm) and 250 µm mesh size.

Invertebrates were identified to the lowest practical taxonomic level, typically genus, using recognized taxonomic keys (Brinkhurst 1986, Clifford 1991, Coffman and Ferrington Jr. 1996, Epler 2001, Maschwitz and Cook 2000, McAlpine et al. 1981, Merritt et al. 2008, Oliver and Roussel 1983, Pennak 1989, Soptonis 1977, Wiederholm 1983). Organisms that could not be identified to the desired level, such as immature or damaged specimens, were reported as a separate category at the lowest taxonomic level possible, typically family. Organisms that required detailed microscopic examination for identification, such as midges (Chironomidae) and aquatic worms (Oligochaeta), were mounted on microscope slides using an appropriate mounting medium. Most common taxa



were distinguishable based on gross morphology and required only a few slide mounts for verification. All rare or less common taxa were slide mounted for identification.

Invertebrates removed from the samples, sorted organic material, and archived samples are being stored for six years to allow possible comparisons, if necessary, with samples collected during subsequent programs.

3.1.4.2 Data Analysis

Data Entry and Screening

Raw benthic invertebrate data were received from the taxonomist in Microsoft Excel® spreadsheet format, with data entry already verified. Non-benthic organisms, such as calanoid copepods (Calanoida), cyclopoid copepods (Cyclopoida), water fleas (Cladocera) and terrestrial invertebrates were removed from the data set prior to data analysis. True fly (Diptera) pupae were also removed prior to data analysis. Abundance data received as number of organisms per sample were converted to density data consisting of number of organisms per square meter (organisms/m²). Unusual abundance data were validated before data summary and analysis.

The following benthic invertebrate summary variables were calculated for each station:

- total invertebrate density;
- taxon richness;
- Simpson's index of diversity (diversity);
- evenness;
- densities of dominant taxa; and
- community composition (i.e., relative densities of major invertebrate taxa).

Richness is the total number of taxonomic groups within a station. It provides an indication of the diversity of benthic invertebrates in an area; a higher richness value usually indicates a more healthy and balanced community.

Simpson's index of diversity measures the proportional distribution of organisms in the community, given that not all organisms have the same success in the environment. Certain conditions may favour one organism over another (Simpson 1949). Simpson's index of diversity values range between 0 and 1, where lower values indicate a community dominated by fewer taxonomic groups (less diverse); these are often referred to as stressed communities. Values close to 1 indicate a community consisting of more taxa that are more evenly distributed among the taxonomic groups present. Simpson's index of diversity was calculated using the formula provided by Krebs (1999), as recommended by Environment Canada (2002) for environmental effects monitoring (EEM) programs:

$$D = 1 - \sum_{i=1}^S (p_i)^2$$

where:

D = Simpson's index of diversity;



S = the total number of taxa; and

p_i = the proportion of the i^{th} taxon.

Evenness is an index recommended by Environment Canada (2002) for analyzing EEM data. It is a measure of how evenly the total invertebrate density is distributed among the taxa present at the site. Evenness is also expressed as a value between one and zero, with one representing high evenness and zero representing low evenness. Evenness was calculated using the formula provided by Smith and Wilson(1996):

$$E = 1 / \sum_{i=1}^S (p_i)^2 / S$$

where:

E = Evenness;

p_i = the proportion of the i^{th} taxon; and

S = the total number of taxa.

Benthic invertebrate summary variables are presented in tabular and graphical format.

Spearman rank correlations were calculated between total benthic invertebrate density, richness, diversity and evenness, and selected habitat variables (total organic carbon, sediment particle size, and water depth). Statistically significant correlations were examined as scatter-plots to determine whether they represented consistent trends or resulted from one or a few atypical points with high leverage on the value of the correlation coefficient. SYSTAT 13 (SYSTAT 2009) was used to calculate Spearman rank correlations.



4.0 RESULTS

4.1 Lower Trophic Communities

4.1.1 Phytoplankton

4.1.1.1 Taxonomic Richness

Six major taxonomic groups (Chlorophyceae, Chrysophyceae, Cyanobacteria, Cryptophyceae, Bacillariophyceae, and Dinophyceae) were represented in the samples collected from the L, M, and N11 lakes and East Lake (Table 3; Appendix I, Tables I-2 and I-3). In general, phytoplankton taxonomic richness was similar among the lakes, with a range of 31 to 37 taxa. The greatest taxonomic richness was observed in East Lake and Lake M2 (37 taxa). Within these lakes, the greatest taxonomic richness was observed in the Chlorophyceae and Chrysophyceae (Table 3). The lowest taxonomic richness was observed in Lakes M1 and L2. In these lakes, the greatest taxonomic richness was observed in the Chrysophyceae (Table 3). Overall, the Chlorophyceae and Chrysophyceae were the most diverse groups, while the other groups appeared to have low taxonomic richness (Table 3).

Table 3 Total Number of Taxa Identified in Each Major Phytoplankton Group in each Lake, August 2011

Lake	Station	Chlorophyceae	Chrysophyceae	Cyanobacteria	Cryptophyceae	Bacillariophyceae	Euglenophyceae	Dinophyceae	Total Taxa
East Lake (Reference Lake)	D1	13	12	6	2	4	0	1	39
	D2	12	11	6	1	5	0	1	36
	D3	13	12	6	2	5	0	1	39
	D4	11	12	5	2	6	0	1	37
	D5	12	10	5	2	5	0	1	35
East Lake Mean		12	11	6	2	5	0	1	37
Lake L2	-	4	15	3	3	3	0	2	31
Lake M1	-	7	12	4	2	2	1	2	31
Lake M2	-	12	13	5	3	3	0	2	37
Lake M3	-	9	12	5	4	2	0	1	33
Lake M4	-	10	11	7	3	4	0	1	36
Lake N11	D1	14	9	7	2	2	0	1	34
	D2	13	9	6	2	2	0	1	33
	D3	14	9	7	2	2	0	1	36
	D4	14	10	5	1	2	0	0	32
	D5	14	12	6	2	2	0	1	37
Lake N11 Mean		14	10	6	2	2	0	1	34



4.1.1.2 *Abundance and Biomass*

Mean total phytoplankton abundance, biomass and community composition were similar among lakes (Figure 4; Figure 5; Figures 6 and 7). The highest mean abundance and variation around the mean was measured in Lake M1 (2,144,000 ind./L), while the lowest abundances were measured in East Lake (1,221,000 ind./L; Figure 4). The highest mean biomass and variation around the mean was observed in Lake L2 (782 mg/m³), while the lowest biomass was measured in Lake M2 (332 mg/m³) (Figure 5).

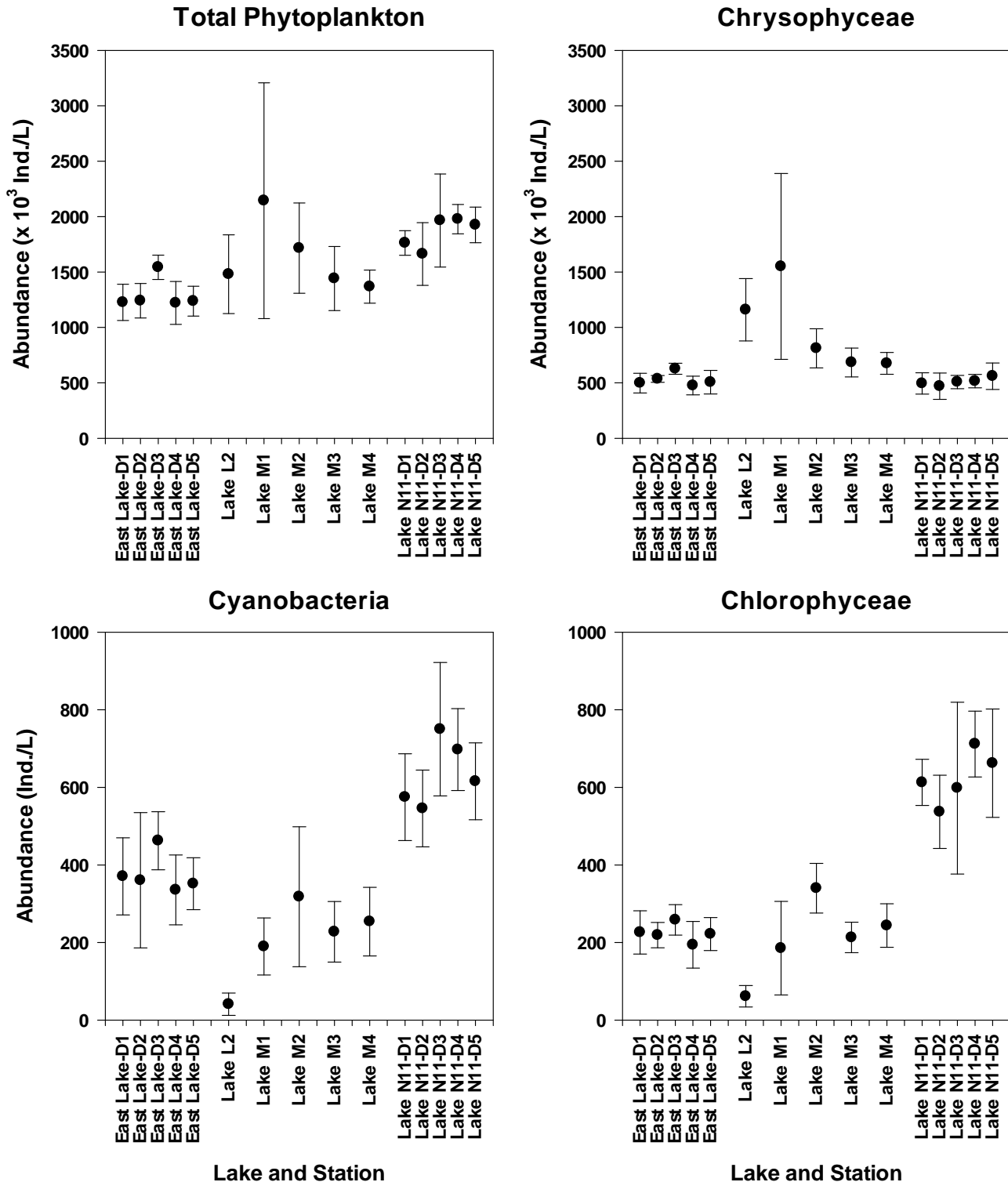
Abundances of the major phytoplankton groups were more variable among lakes than total abundance. Overall, phytoplankton abundance was dominated by Chlorophyceae (26 to 78%), followed closely by Chrysophyceae (4 to 36%) and cyanobacteria (3 to 38%), depending on lake (Figure 6). Chrysophycean abundance was high in Lakes M1, M2, M3, M4, L2 and East Lake, with the greatest abundance observed in Lake M1 (Figure 4). Abundance in Lake N11 was evenly divided amongst the Chrysophyceae, cyanobacteria and Chlorophyceae. Bacillariophyceae, Cryptophyceae, Euglenophyceae and Dinophyceae abundances were relatively low in all of the lakes (Figures 4 and 6).

Similar to abundance, biomass estimates of the major phytoplankton groups were more variable among lakes than total biomass. Overall, phytoplankton biomass was dominated by Chrysophyceae (15 to 57%), followed by Chlorophyceae (12 to 30%) and cyanobacteria (1 to 50%; Figure 7). Lakes L2, M1, M2, M3, and M4 were dominated by Chrysophycean biomass, while biomass in Lake N11 and East Lake were dominated by cyanobacteria and Chlorophyceae. Dinophyceae biomass was high in Lake L2 (304 mg/m³) and at Station D2 in Lake N11 (207 mg/m³) compared to the other lakes and stations (<100 mg/m³). Overall, Bacillariophyceae biomass was low, with the highest bacillariophycean biomass observed in East Lake (150 mg/m³; Figure 5). Euglenophycean and cryptophycean biomass were low in all lakes (Figures 5 and 7).



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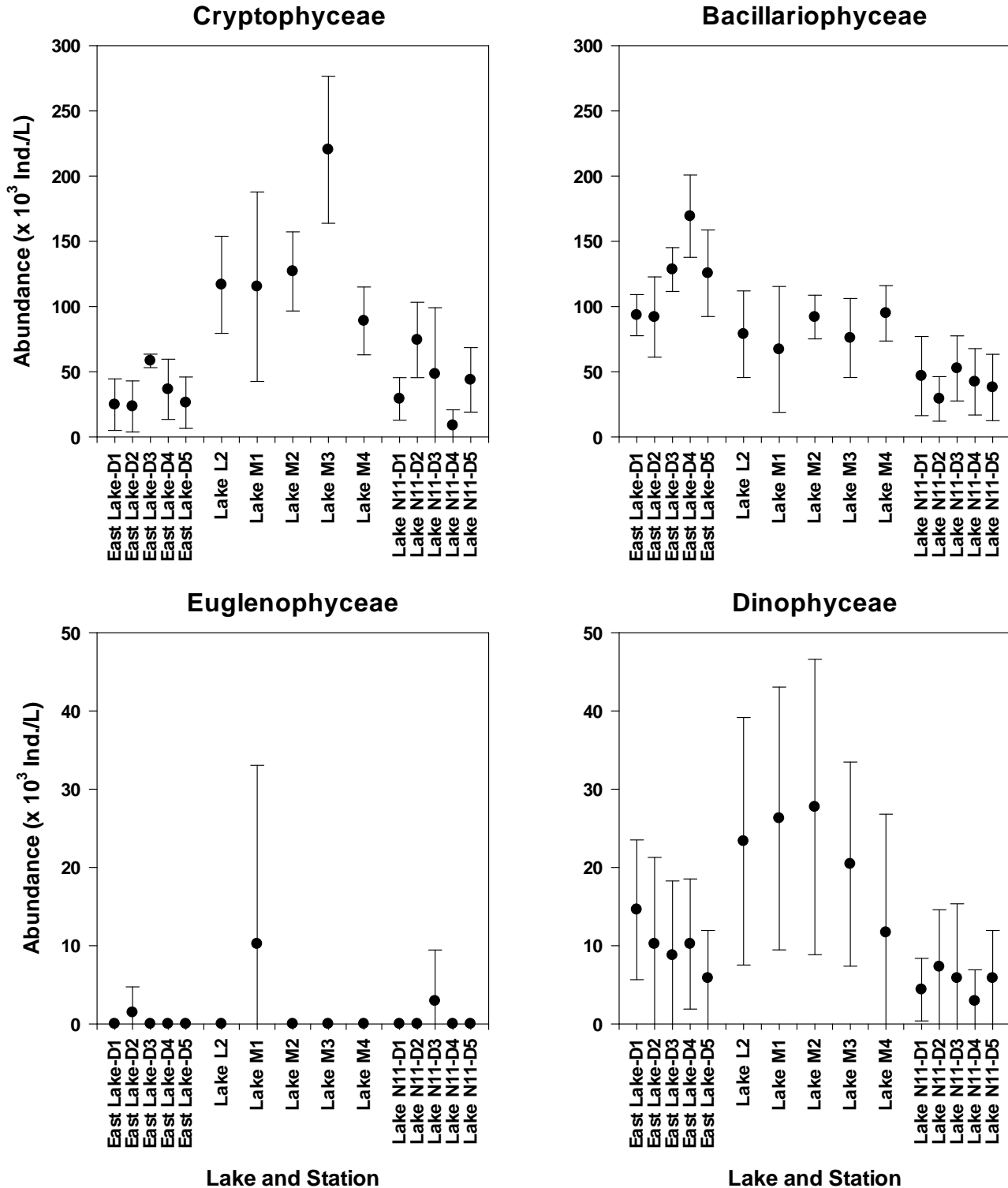
Figure 4 Total Phytoplankton Abundance (Mean \pm Standard Deviation) and Abundances of Major Phytoplankton Groups in each Lake, August 2011





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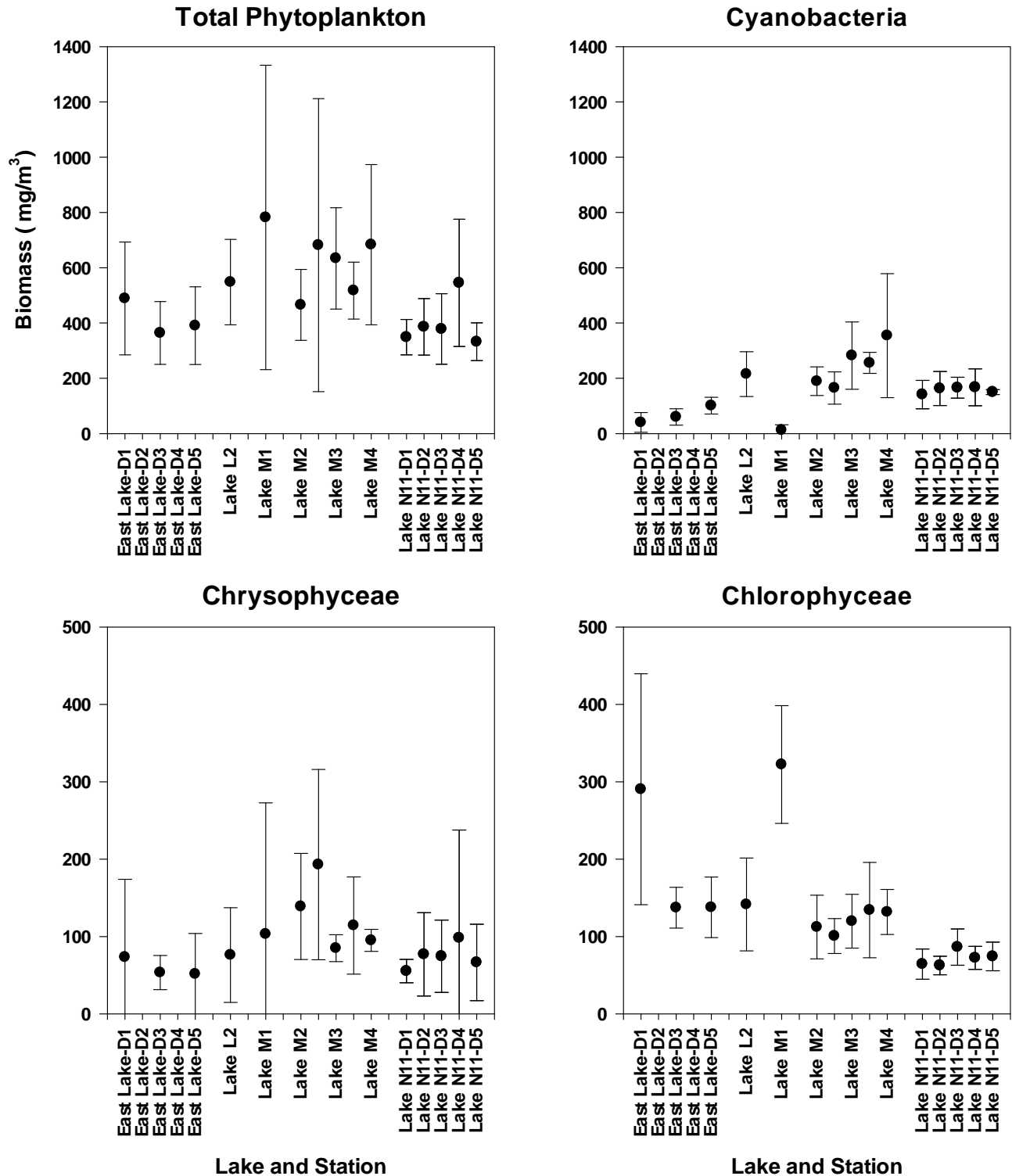
Figure 4 Total Phytoplankton Abundance (Mean \pm Standard Deviation) and Abundances of Major Phytoplankton Groups in each Lake, August 2011 (continued)





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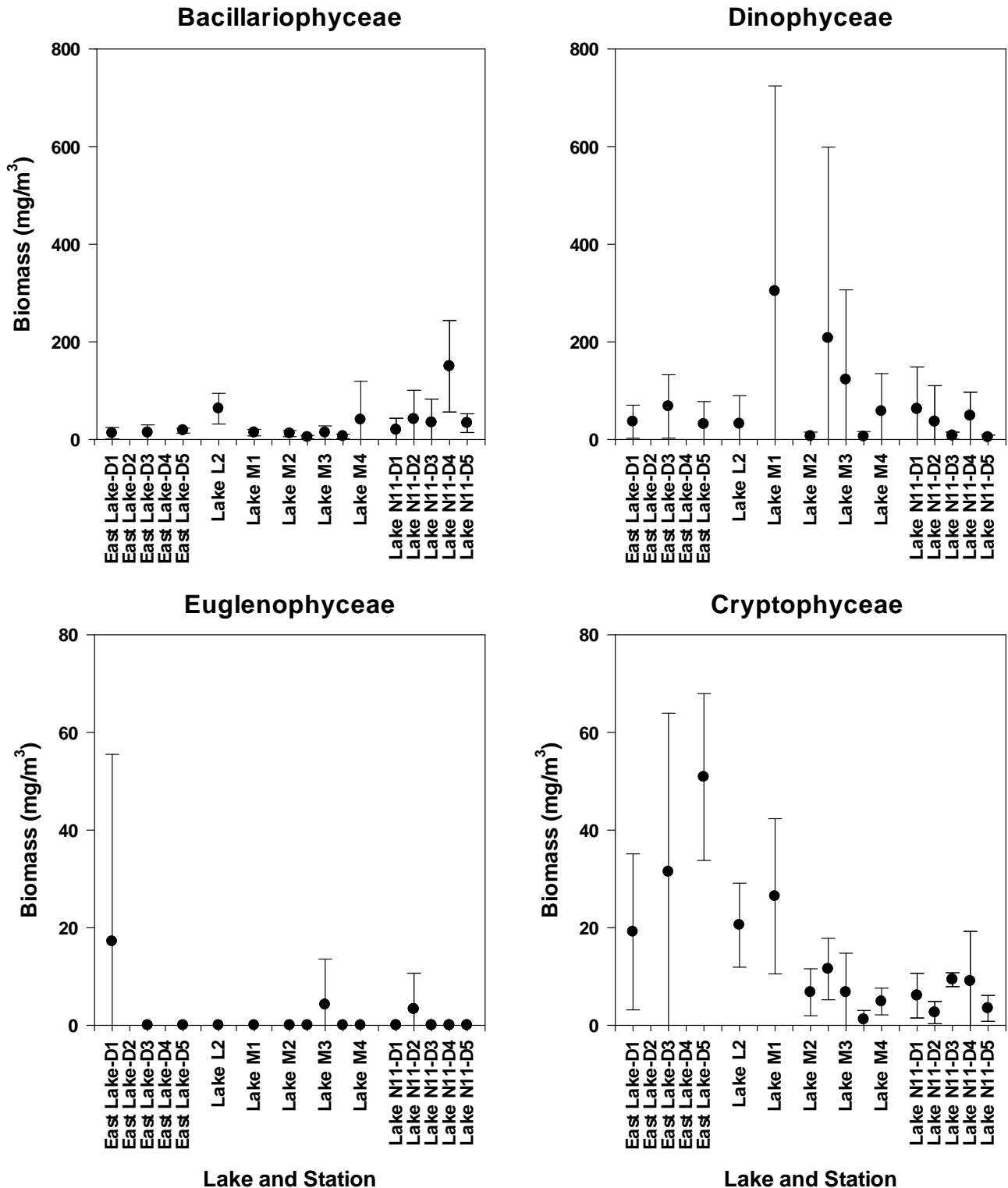
Figure 5 Total Phytoplankton Biomass (Mean \pm Standard Deviation) and Biomass of Major Phytoplankton Groups in each Lake, August 2011





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Figure 5 Total Phytoplankton Biomass (Mean \pm Standard Deviation) and Biomass of Major Phytoplankton Groups in each Lake, August 2011 (continued)





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Figure 6 Variation in Relative Abundances of Major Phytoplankton Groups in each Lake, August 2011

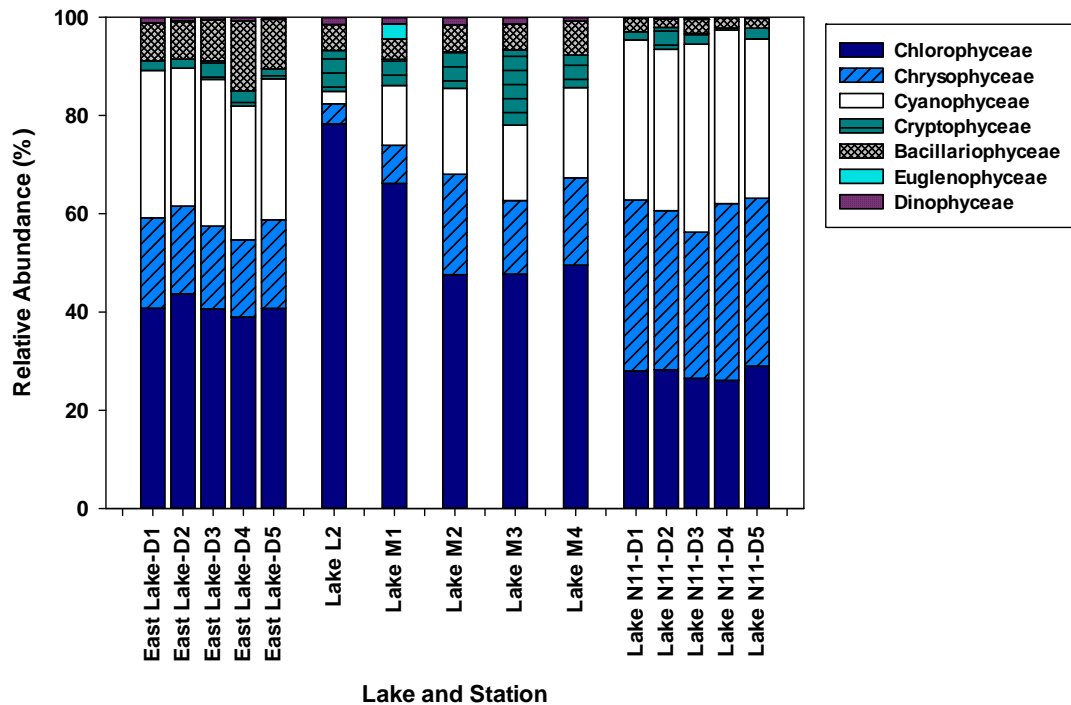
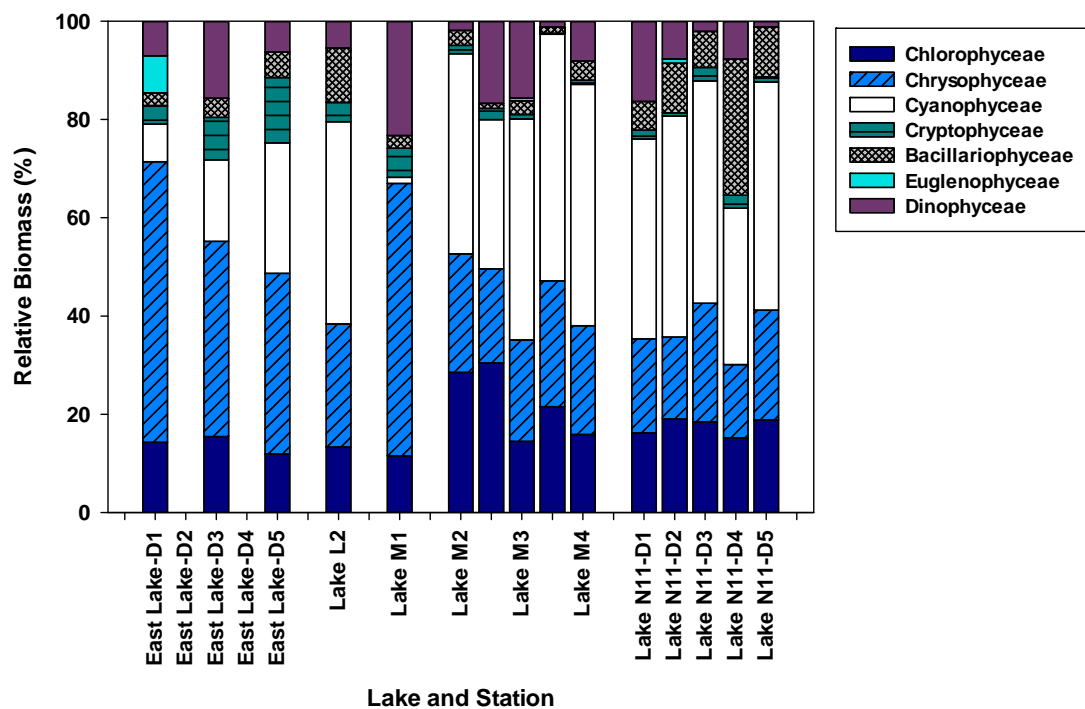


Figure 7 Variation in Relative Biomass of Major Phytoplankton Groups in each Lake, August 2011





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4.1.2 Zooplankton

4.1.2.1 Richness

Four major taxonomic groups (Cladocera, Calanoida, Cyclopoida, and Rotifera) were represented in the samples collected from the L, M, and N11 lakes and East Lake in August 2011 (Table 4). Taxonomic richness was similar among the lakes, with a range of 12 to 14. Taxonomic richness was generally evenly distributed among major groups, with Cyclopoida and Cladocera having the lowest diversity (2 to 3 taxa) and Calanoida and Rotifera having the greatest diversity (about 4 taxa) (Table 4).

Table 4 Total Number of Taxa Identified in the Major Zooplankton Groups in each Lake, August 2011

Lakes	Station	Calanoida	Cyclopoida	Cladocera	Rotifera	Total Taxa
East Lake	D1	3	2	3	4	12
East Lake	D2	4	2	3	4	13
East Lake	D3	4	2	3	4	12
East Lake	D4	3	2	2	3	9
East Lake	D5	4	2	3	3	12
East Lake Mean		4	2	3	4	12
Lake L2	-	4	3	2	3	13
Lake M1	-	4	2	3	4	13
Lake M2	-	5	2	3	4	14
Lake M3	-	4	2	3	4	14
Lake M4	-	3	2	3	4	12
Lake N11	D1	4	2	2	4	12
Lake N11	D2	3	2	2	4	12
Lake N11	D3	3	3	2	4	12
Lake N11	D4	3	2	2	4	11
Lake N11	D5	3	2	2	4	11
Lake N11 Mean		3	2	2	4	12

Note: Cladocera, Calanoida, and Cyclopoida were identified to species where they could be; Rotifera were identified to genus.

4.1.2.2 Abundance and Biomass

Abundance, biomass and community composition of zooplankton were more variable among the lakes than phytoplankton (Figures 8 to 11; Appendix I, Tables I-4 and I-5). Total zooplankton abundance ranged from 14 ind./L in East Lake Station D3 to 108 ind./L in Lake M2 (Figure JJ4.3-5). Total zooplankton biomass ranged from 80 mg/m³ in East Lake Station D4 to 1,931 mg/m³ in Lake N11 at Station D1 (Figure 8).

Zooplankton abundance was dominated by rotifers and cyclopoid copepods (Figures 8 and 10). Overall, rotifers accounted for 31 to 81% of the total abundance, and cyclopoid copepods accounted for 6 to 51%. Cladocerans and calanoid copepods accounted for a small proportion (<1 to 19% and 4 to 16%, respectively) of the zooplankton community based on abundance.

Relative biomass of major zooplankton groups was highly variable among the lakes (Figures 9 and 11). Cladocera biomass dominated in Lake N11 (80 to 91%), while calanoid copepod biomass dominated in East Lake (45 to 70%) and Lake L2 (78%; Figure 11). Lakes M1 and M2 were co-dominated by calanoid copepods and cladocerans, while Lake M3 and M4 were dominated by cladocerans and sub-dominated by calanoid and cyclopoid copepods (Figure 11).



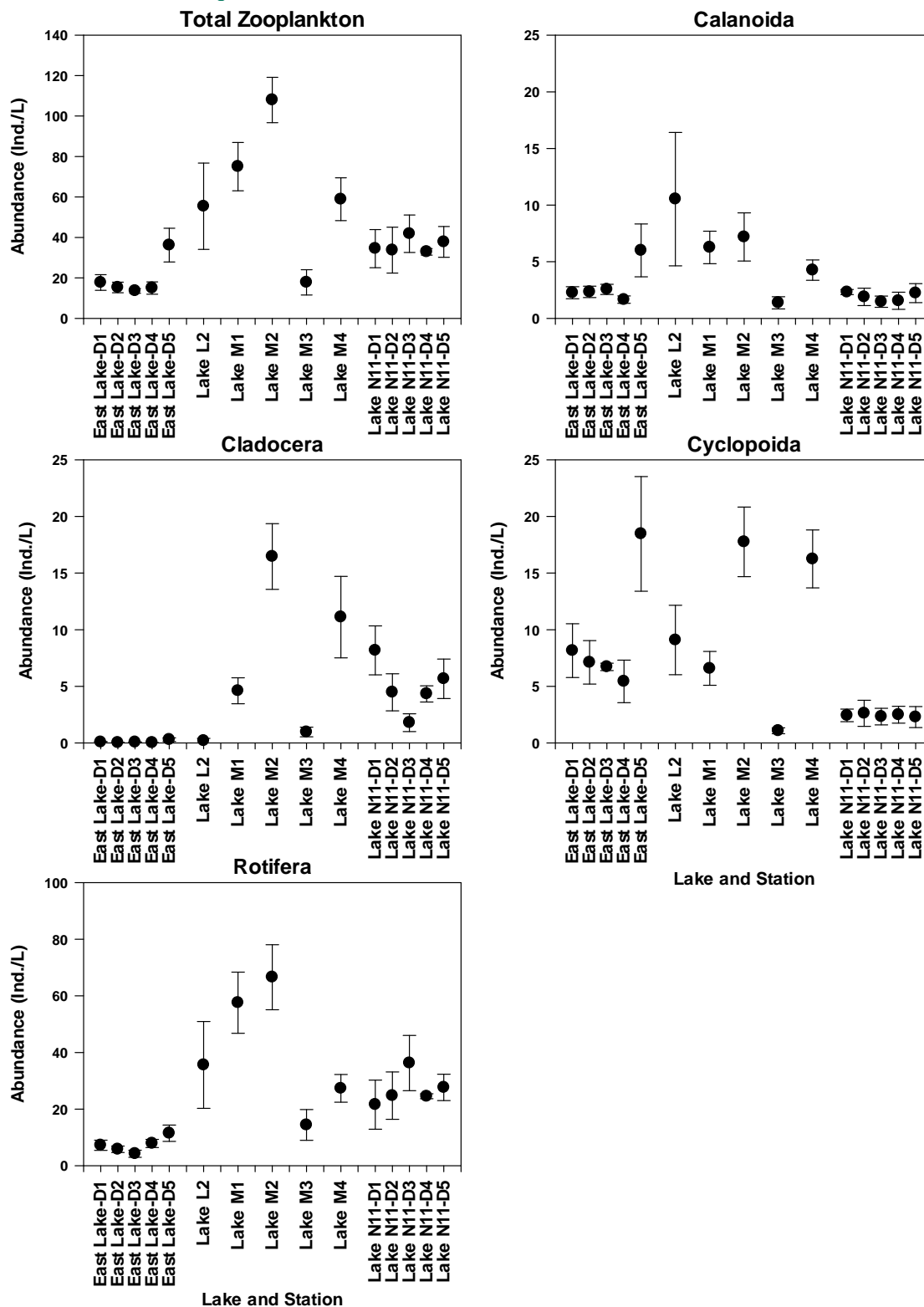
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Despite rotifers being the most abundant group in the lakes, their small body size explains their low relative biomass. In contrast, the large size of calanoid copepods and Cladocera account for their larger contributions to total biomass, despite their low relative abundance.



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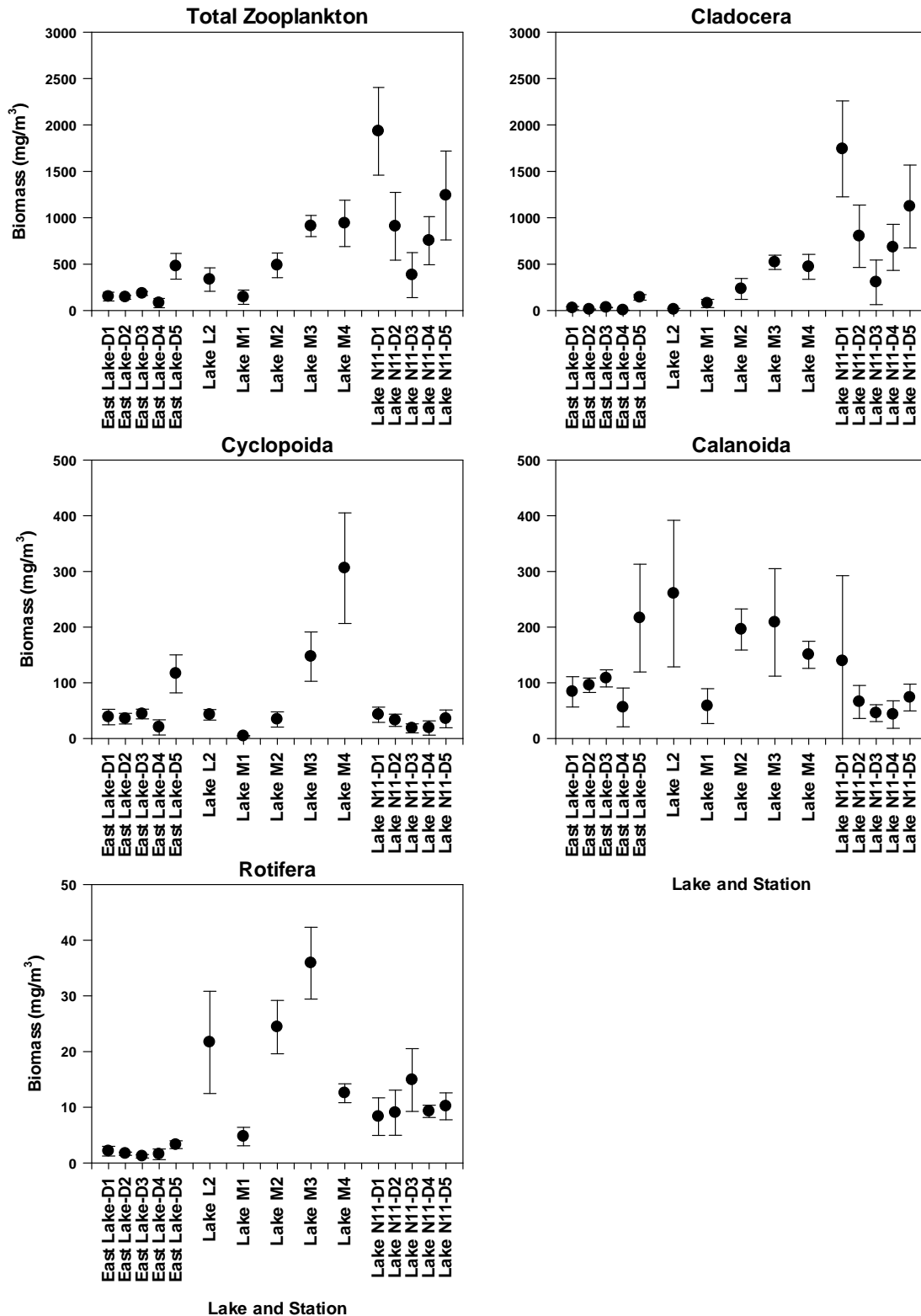
Figure 8 Total Zooplankton Abundance (Mean \pm Standard Deviation) and Abundance of Major Zooplankton Groups in each Lake and Station, August 2011





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Figure 9 Total Zooplankton Biomass (Mean \pm Standard Deviation) and Biomass of Major Zooplankton Groups in each Lake, August 2011





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Figure 10 Relative Abundance of Major Zooplankton Groups in each Lake, August 2011

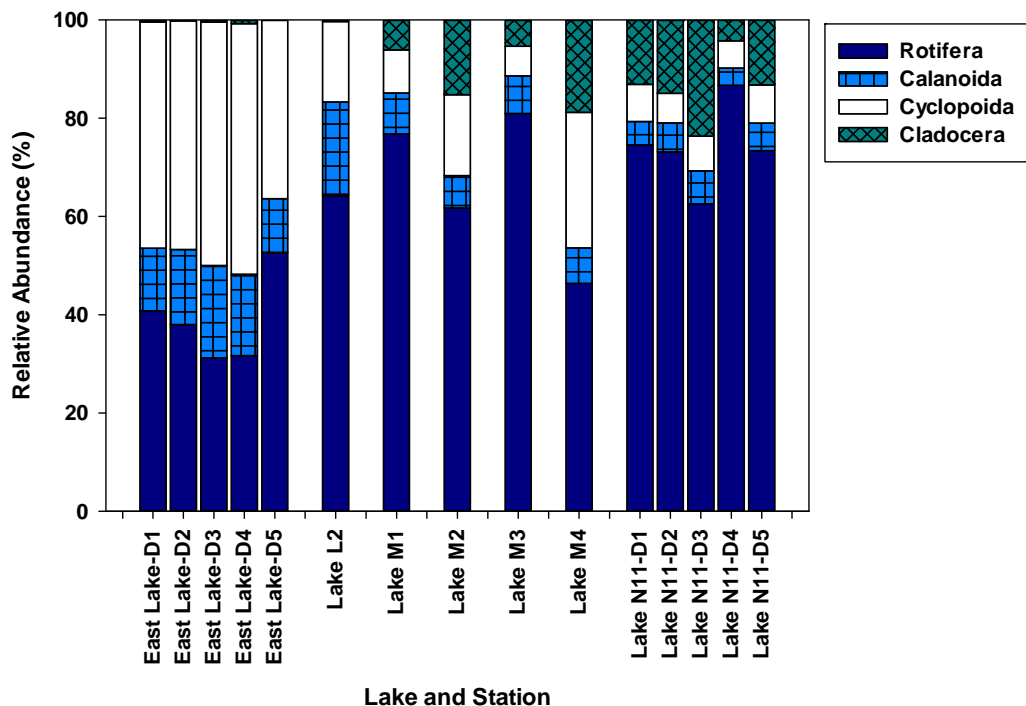
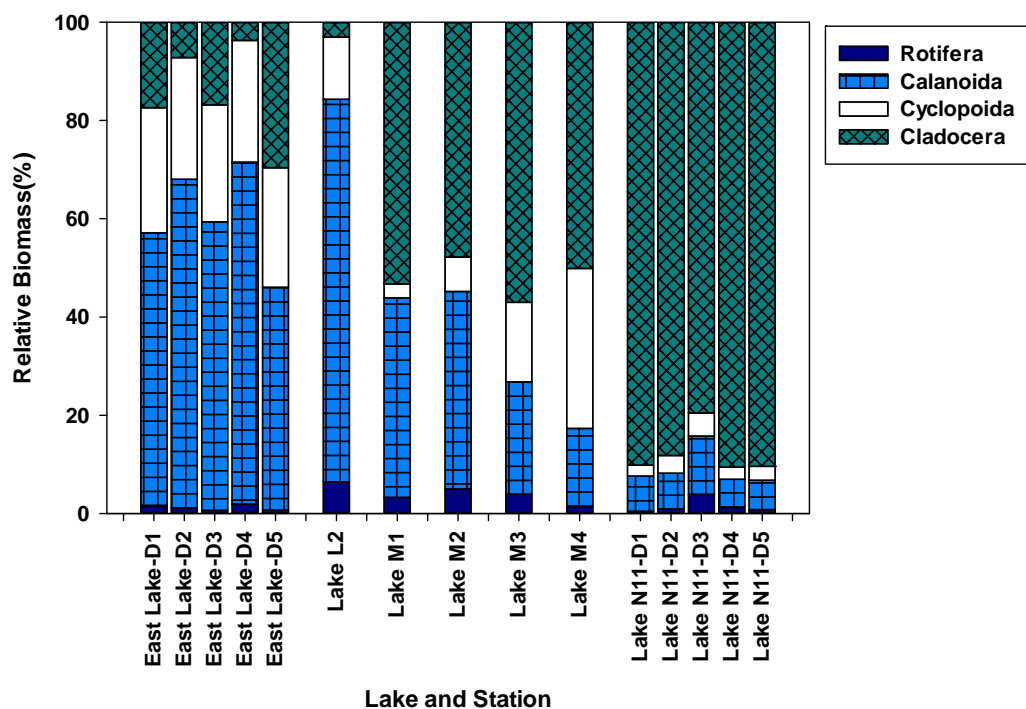


Figure 11 Relative Biomass of Major Zooplankton Groups in each Lake, August 2011





4.1.3 Benthic Invertebrates

4.1.3.1 Habitat Characteristics

Water depth at benthic invertebrate sites sampled in 2011 ranged from 7 to 15 m in East (Reference) Lake, from 2 to 12 m in the L and M Lakes, and from 6 to 7 m in Lake N11 (Appendix II, Table II-1). Water temperature, dissolved oxygen, specific conductivity and pH varied little with depth in the water column, indicating that the water column was well mixed in the lakes sampled. Surface water temperature ranged from 15.1 to 15.2°C in East Lake, from 9.6 to 13.6°C in the L and M lakes, and from 14.2 to 16.3°C in Lake N11. Surface DO ranged from 9.1 to 9.3 mg/L in East Lake, from 9.8 to 10.6 mg/L in the L and M lakes, and from 9.1 to 9.5 mg/L in Lake N11. Specific conductivity at the surface was low in all lakes, as expected in sub-arctic lakes. Specific conductivity was 15 µS/cm at all stations in East Lake, was 15 µS/cm in each of the L and M lakes, and ranged from 11 to 12 µS/cm in Lake N11. Surface pH ranged from 6.7 to 6.9 in East Lake, ranged from 6.6 to 6.9 in the L and M Lakes, and ranged from 6.1 to 6.8 in Lake N11.

Sediment characteristics were variable within and among lakes sampled in 2011 (Appendix II, Table II-1). Sediment data are available for two stations sampled in the L and M lakes (i.e., Lake M3 and Lake M4). Moisture content ranged from 71 to 92% in East Lake, from 87 to 97% in Lake N11, and from 90 to 92% in the L and M Lakes. Total organic carbon was variable in East Lake ranging from 3 to 13%. Total organic carbon was similar among stations in Lake N11, ranging from 9 to 16%, and in the L and M Lakes ranging from 13 to 15%. Bottom sediments in all lakes were dominated by sand, with sand content ranging from 62 to 87% in East Lake, from 63 to 67% in the L and M Lakes, and from 72 to 78% in Lake N11.

Spearman rank correlation analysis detected significant negative correlations between water depth, and total density and total richness ($P < 0.05$, $r_s > 0.521$, $n = 15$) (Table 5). The range in water depths (2 to 15 m) at benthic sampling locations moderately influenced the benthic invertebrate community. In general, both benthic invertebrate density and richness decreased with increasing depth, which is consistent with habitat associations of benthic invertebrates in lakes.

Table 5 Spearman Rank Correlations Between Benthic Invertebrate Variables and Habitat Variables

	Water	Total Organic	Percent Fines
Variable	Depth	Carbon	(silt + clay)
Total density	-0.545	0.086	-0.173
Total richness	-0.567	0.178	-0.074
Mean richness	-0.487	0.136	-0.088
Simpson's diversity index	-0.048	-0.473	-0.201
Evenness	0.195	-0.244	0.180

Note: Significant relationships ($P < 0.05$, $r_s > 0.521$, $n = 15$) are **bolded**.

4.1.3.2 Benthic Invertebrate Community

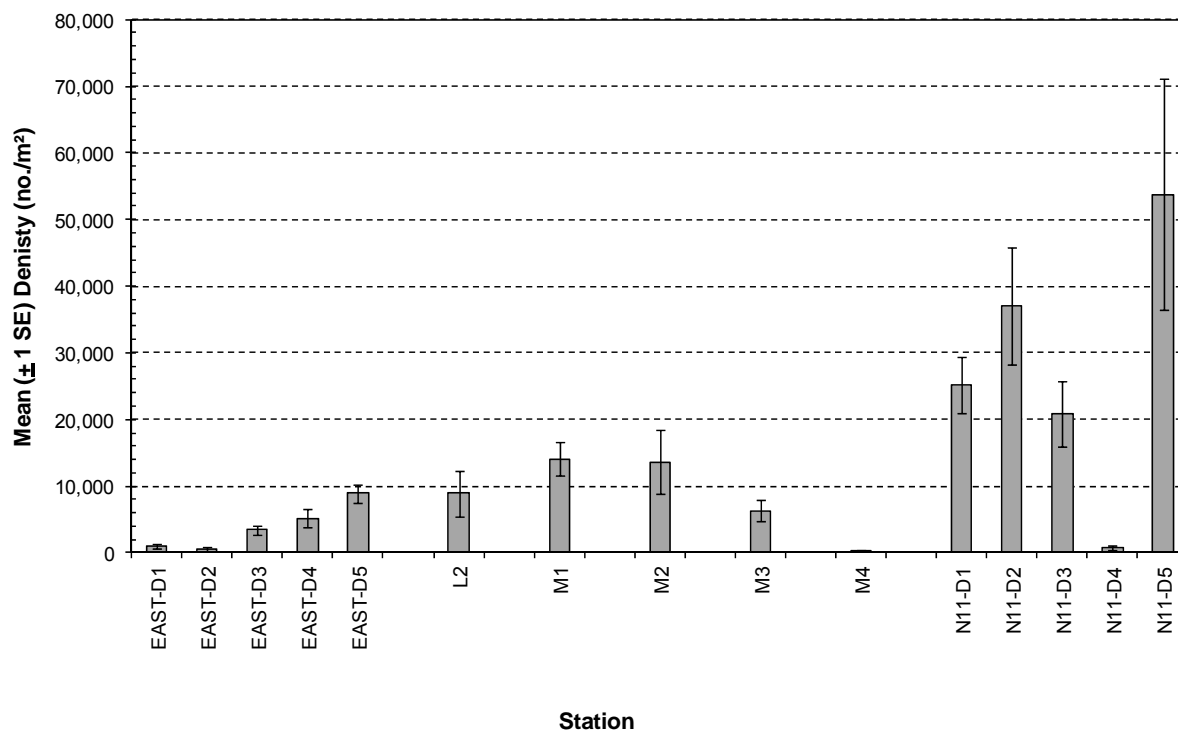
Total benthic invertebrate density was variable within and among lakes (Figure 12; Appendix II, Table II-2). Mean (± 1 SE) benthic invertebrate density ranged from 621 ± 325 to $8,914 \pm 1,418$ organisms/m² in East Lake, from 810 ± 316 to $53,776 \pm 17,303$ organisms/m² in Lake N11, and from 241 ± 84 to $14,095 \pm 2,554$ organisms/m² in the L and M lakes. In general, pooled means for stations in each lake were approximately 7 times greater in Lake N11 compared to East Lake, with only one station having a total density below 20,000 organisms/m² (Station N11-D4; 810 ± 316 organisms/m²). Mean density was approximately 2



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times greater in the L and M lakes compared to East Lake, with only one site having a density below 5,000 organism/m² (Lake M4; 241 ± 84 organisms/m²). Densities in Lake N11 were higher than expected for a sub-Arctic lake, particularly for stations N11-D1, N11-D2, N11-D3 and N11-D5. The lake variation in density was also influenced by the variation in sample depth.

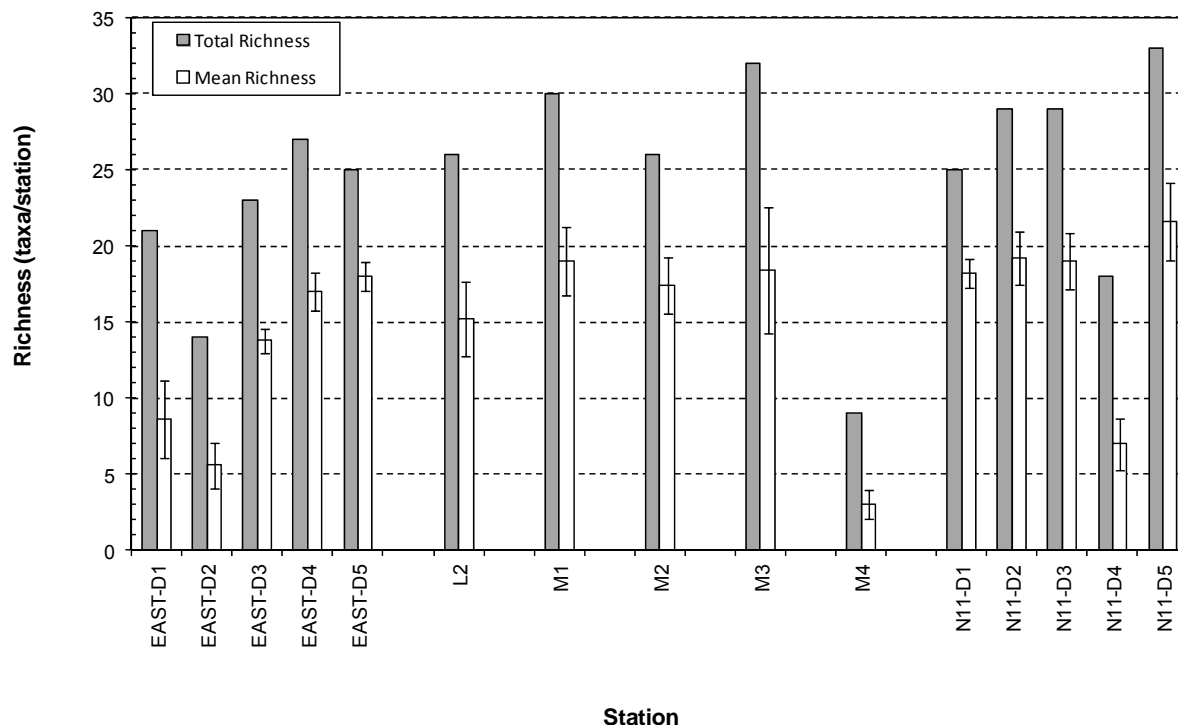
Figure 12 Mean Total Benthic Invertebrate Density at Lake Sampling Locations, Fall 2011



Richness ranged from low to moderate and varied within similar ranges among lakes. Total richness ranged from 14 to 27 taxa/station in East Lake, from 18 to 33 taxa/station in Lake N11, and from 9 to 30 taxa/station in the L and M Lakes (Figure 13; Appendix II, Figure II-2). Mean (± 1 SE) richness ranged from 6 ± 2 to 18 ± 1 taxa/station in East Lake, from 7 ± 2 to 22 ± 3 taxa/station in Lake N11, and from 3 ± 1 to 19 ± 2 taxa/station in the L and M Lakes. Overall, taxa richness was in the expected range for sub-Arctic lakes (Beaty et al. 2006).



Figure 13 Benthic Invertebrate Richness at Lake Sampling Locations, Fall 2011



Simpson's index of diversity values were generally high, with the exception of Lake N11, where diversity ranged from moderate to high. Diversity ranged from 0.83 to 0.90 in East Lake, from 0.81 to 0.91 in the L and M Lakes, and from 0.52 to 0.83 in Lake N11 (Figure 14; Appendix II Table II-2). Evenness was generally low to moderate in the lakes sampled, ranging from 0.26 to 0.53 in East Lake, from 0.29 to 0.59 in the L and M Lakes, and 0.06 to 0.33 in Lake N11. This indicated that a few taxa usually accounted for the majority of the total density observed at a station.

The benthic invertebrate community was dominated by the midges at all stations sampled in 2011, with the exception of station EAST-D2 in East Lake where the aquatic worms were dominant, and stations N11-D2 and N11-D3 in Lake N11, where the roundworms were co-dominant with the midges (Figure 15; Appendix II, Table II-3). Other abundant taxa included the roundworms, aquatic worms and clams (Pelecypoda). A combined total of 49 taxa were collected in the lakes sampled in Fall 2011, 30 of which were midges (Table 6). Dominance of the benthic invertebrate community by the midges is as expected for lakes in the sub-arctic region (Beaty et al. 2006, Danks 1981, Danks 2007).



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Figure 14 Simpson's Index of Diversity and Evenness at Lake Sampling Locations, Fall 2011

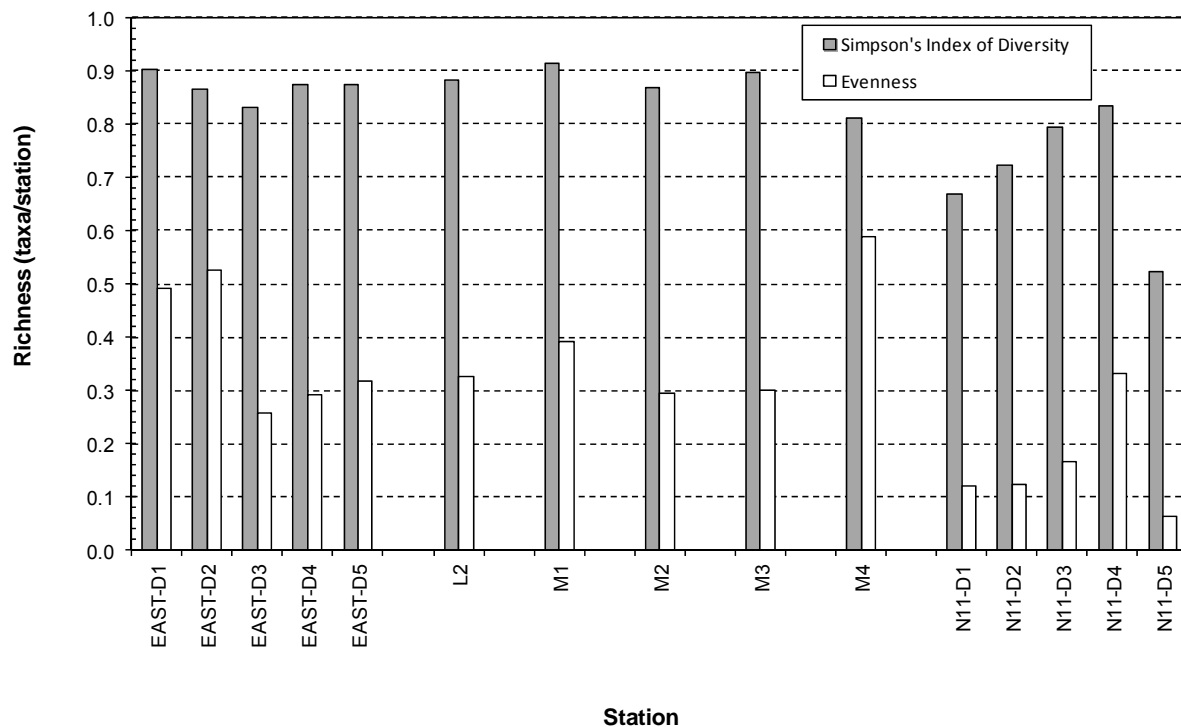
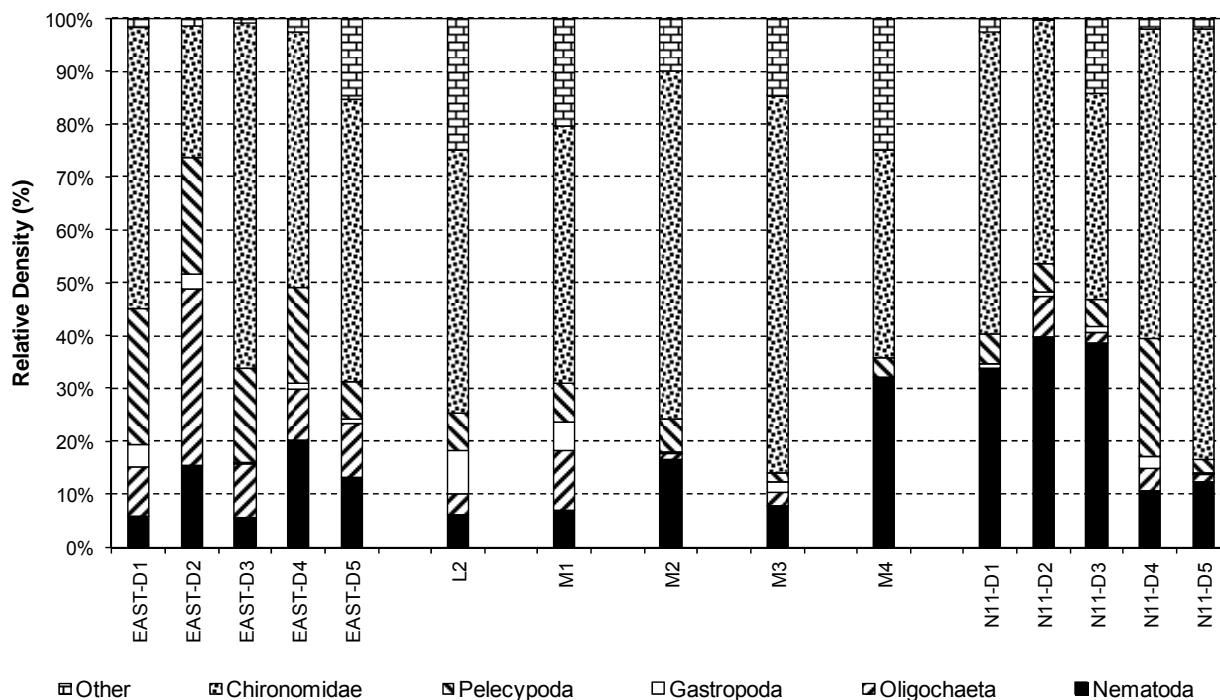


Figure 15 Benthic Invertebrate Community Composition at Lake Sampling Locations, Fall 2011





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Table 6 List of Benthic Invertebrate Taxa Collected at Lake Sampling Locations, Fall 2011

Major Taxon	Family	Subfamily	Tribe	Genus/Species
Microturbellaria	Typhloplanidae	-	-	<i>Mesostoma</i>
Nematoda	-	-	-	-
Oligochaeta	Enchytraeidae	-	-	-
	Lumbriculidae	-	-	-
	Naididae	Naidinae	-	-
	Naididae	Tubificinae	-	-
Gastropoda	Valvatidae	-	-	<i>Valvata sincera</i>
Bivalvia	Pisidiidae	-	-	<i>Sphaerium</i>
		-	-	<i>Pisidium</i>
Hydracarina	-	-	-	-
Copepoda - Harpacticoida	-	-	-	-
Ostracoda	-	-	-	-
Trichoptera	Hydroptilidae	-	-	<i>Agraylea</i>
	Leptoceridae	-	-	<i>Oecetis</i>
	Limnephilidae	-	-	<i>Grensia praeterica</i>
	Phryganeidae	-	-	<i>Phryganea</i>
Diptera	Chironomidae	Tanypodinae	Pentaneurini	<i>Ablabesmyia</i>
				<i>Thienemannimyia</i> group
		Diamesinae	Procladiini	<i>Procladius</i>
			Protanypini	<i>Protanypus</i>
		Prodiamesinae	-	<i>Monodiamesa</i>
		Orthoclaadiinae	Orthoclaidiini	<i>Abyskomyia</i>
				<i>Cricotopus / Orthocladus</i>
				<i>Heterotanytarsus</i>
				<i>Heterotrissocladius</i>
				<i>Parakiefferiella</i>
				<i>Psectrocladius</i>
				<i>Zalutschia</i>
		Chironominae	Chironomini	<i>Chironomus</i>
				<i>Cladopelma</i>
				<i>Cryptochironomus</i>
				<i>Dicrotendipes</i>
				<i>Microtendipes</i>
				<i>Pagastiella</i>
				<i>Parachironomus</i>
				<i>Polypedilum</i>
				<i>Sergentia</i>
				<i>Stictochironomus</i>
				<i>Pseudochironomus</i>
			Tanytarsini	<i>Cladotanytarsus</i>
				<i>Corynocera</i>
				<i>Micropsectra</i>
				<i>Micropsectra / Tanytarsus</i>
				<i>Paratanytarsus</i>
				<i>Stempellinella</i>
				<i>Tanytarsus</i>
	Ceratopogonidae	Ceratopogoninae	-	<i>Bezzia</i>
		Dasyheleinae	-	<i>Dasyhelea</i>
	Empididae	-	-	<i>Chelifera / Metachela</i>

- = not identified to this taxonomic level.



5.0 SUMMARY AND CONCLUSIONS

5.1 Lower Trophic Communities

5.1.1 Phytoplankton

In general, phytoplankton taxonomic richness, abundance, biomass and community composition were similar among the sampled lakes. The dominant algal groups by abundance and biomass were Chlorophyceae, Chrysophyceae and cyanobacteria. This is typical of lakes with low to moderate productivity (Wetzel 2001). Similarly, phytoplankton taxonomic richness was diverse in terms of the numbers of taxa present. This is often observed in low productivity lakes, where slower growth rates permit a greater number of species to coexist, compared to more productive waters (Wetzel 2001).

5.1.2 Zooplankton

Abundance, biomass and community composition of zooplankton were more variable among the lakes, compared to phytoplankton. Total zooplankton abundance ranged from 14 to 108 ind./L depending on the lake, and total zooplankton biomass ranged from 80 to 1,931 mg/m³. Zooplankton abundance was dominated by rotifers and cyclopoid copepods, while biomass was dominated by Cladocera and calanoid copepods. Taxonomic richness was similar among lakes (range 12-14). Taxonomic richness was evenly distributed among groups, with Cyclopoida and Cladocera having the lowest taxonomic diversity, and Calanoida and Rotifera having the greatest taxonomic diversity. The zooplankton communities documented in the lakes sampled are similar to those in other sub-Arctic lakes, such as Lac de Gras (Golder 2010) and Snap Lake (De Beers 2010b).

5.1.3 Benthic Invertebrates

The benthic invertebrate communities of lakes were characterized by low to moderate density and richness during the fall 2011 sampling program, consistent with the generally low productivity typical of sub-Arctic lakes on the Canadian Shield. Overall, Simpson's diversity was high and evenness was low to moderate indicating that a few taxa accounted for most of the organisms present in lakes. Midges were the dominant taxa, with the aquatic worms, roundworms and fingernail clams also representing a considerable proportion of the benthic invertebrate community at some stations.

The benthic invertebrate community in the lakes sampled in the Gahcho Kué study area, with the exception of Lake N11, is consistent with that expected in the sub-Arctic region where low productivity is common due to low nutrient levels, low temperatures, and long ice covered periods. Lake N11 is moderately productive compared to other lakes historically sampled in the area (De Beers 2010a, Annex J and Addendum JJ). Also, a moderately strong relationship existed between water depth, and both total density and richness in the lakes sampled in fall 2011.



6.0 CLOSURE

We trust the above meets your present requirements. If you have any questions or require additional details, please contact the undersigned.

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8.0 ABBREVIATIONS

De Beers	De Beers Canada Inc.
DO	dissolved oxygen
EEM	environmental effects monitoring
EIR	Environmental impact review
EIS	Environmental impact assessment
GPS	Global positioning system
LSA	local study area
n	number
NWT	Northwest Territories
P	probability
Project	Gahcho Kué Project
QA/QC	quality assurance / quality control
RPD	relative percent difference
RSA	regional study area
SE	standard error
TOC	total organic carbon
UTM	Universal Transverse Mercator

8.1 Units of Measure

%	percent
<	less than
>	greater than
°C	degrees Celsius
µg/L	micrograms per litre
µm	micrometre
µS/cm	microSiemens per centimetre
cm	centimetre
g	gram
g/cm ³	grams per cubic centimetre
ind/L	individuals per litre
km	kilometre
km ²	square kilometre
L	litre
m	metre
m ²	square metre
m ³	cubic metre
mg/L	milligrams per litre



%	percent
mg/ m ³	milligrams per cubic metre
mL	millilitre
mm	millimetre
mm ³	cubic millimetres
mm ³ / m ³	cubic millimetres per cubic metre wet weight
org/m ²	number of organisms per square metre

9.0 GLOSSARY

Benthic Invertebrates	Animals without backbones that live on river and lake bottoms. Benthic refers to the bottom, and these animals are also called zoobenthos.
Calanoida	An order of copepods; small planktonic animals that are a component of zooplankton.
Chlorophyta	Green algae; a component of phytoplankton.
Chrysophyta	Golden-brown algae; a component of phytoplankton.
Cladocera	A group of small planktonic animals (crustaceans) also known as water fleas; a component of zooplankton.
Colonial	Individuals of the same species clustered together to form a group.
Conductivity	A measure of the resistance of a solution to electrical flow; an indirect measure of the salinity of the water.
Copepoda	An order of planktonic crustacean; a component of zooplankton.
Cryptophyta	Flagellated algae also known as cryptomonads; a component of phytoplankton.
Cyanobacteria	Blue-green algae; a component of phytoplankton.
Cyclopoida	An order of copepods; small planktonic animals.
Diatom	A group of algae that are encased within a frustule made of silica; a component of phytoplankton.
Dissolved Oxygen	Oxygen dissolved within the water column.
Diversity	A numerical index that incorporates evenness and richness; the diversity index measures the proportional distribution of organisms in the community.
Enumeration	The act of counting individuals.
Euglenophyta	Euglena; a component of phytoplankton.
Evenness	A measure of how evenly the total invertebrate abundance is distributed among the different types of organisms present at the site.
Limnology Profiles	Refers to measurements of water temperature, conductivity, pH, and dissolved oxygen in the water column of a lake.
Littoral	The shallow, shoreline area of a lake.
Lower trophic	Organisms in an ecosystem that form the bottom of the food chain (benthic invertebrates, zooplankton, and phytoplankton) upon which fish depend as food.
Pelagic	Relating to fish or other aquatic organisms that live offshore in the middle or lower part of the water column.



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Periphyton	Algae and small crustaceans that live attached to rocks and other substrates projecting from the bottom of a stream or lake.
pH	A measure of the acidity or alkalinity of water.
Phytoplankton	Small, usually microscopic, plants that live in the water column of lakes and make their food through primary production.
Plankton	Small, often microscopic, plants (phytoplankton) and animals (zooplankton) that live in the open water column of lakes. They are an important food source for many larger animals.
Richness	The number of different types of animals present in a sample or at a location.
Rotifera	A large class of the pseudocoelomate phylum Aschelminthes; a component of zooplankton.
Secchi Depth	A measure of water clarity, measured by lowering a 20 cm diameter disk (Secchi disk) with alternating black and white coloured quadrants. The shallowest depth at which the disk is no longer visible is the Secchi depth.
Substrate	The bottom of a waterbody, usually consisting of sediments of various particle sizes (e.g., sand, silt, clay, gravel, cobble, boulder) and organic material (e.g., living or dead plant material).
Taxon	A group of organisms at the same level of the standard biological classification system; the plural of taxon is taxa.
Terrestrial	Living or growing on land.
Total Organic Carbon (TOC)	Total organic carbon is composed of both dissolved and particulate forms. Total organic carbon is often calculated as the difference between Total Carbon (TC) and Total Inorganic Carbon (TIC). Total organic carbon has a direct relationship with both biochemical and chemical oxygen demands, and varies with the composition of organic matter present in the water. Organic matter in soils, aquatic vegetation and aquatic organisms are major sources of organic carbon.
Watershed	The upstream land area drained by a river network.
Yellow Springs Instrument (YSI)	A meter that measures temperature, conductivity and dissolved oxygen in water.
Zooplankton	Small, sometimes microscopic, animals that live in the water column of lakes and mainly eat primary producers (phytoplankton).



APPENDIX I

Plankton - Supporting Data, 2011



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Table I-1: Water Quality Profiles at Plankton Sampling Locations in East Lake, Lake N11 and the L and M Lakes of the Gahcho Kué Project, Fall 2011

Lake	Station	Maximum Depth [m]	Secchi Depth [m]	Depth [m]	Temperature [°C]	Dissolved Oxygen [mg/L]	Specific Conductivity [µS/cm]	pH
Lake M1	-	2.0	2.0	0.3	9.6	10.6	14.0	6.8
				1.0	9.6	10.5	14.0	6.7
				1.5	9.6	10.6	14.0	6.0
Lake M2	-	4.5	3.0	0.3	13.5	9.8	14.0	6.7
				1.0	13.5	9.8	14.0	6.7
				1.5	13.5	9.8	14.0	6.6
				2.0	13.5	9.8	14.0	6.5
				2.5	13.5	9.8	14.0	6.5
				3.0	13.5	9.8	14.0	6.5
				3.8	13.6	9.8	14.0	6.5
				4.0	13.6	9.8	14.0	6.4
Lake M3	-	7.3	3.2	4.5	13.6	9.8	14.0	6.4
				0.3	14.3	9.8	14.0	6.6
				1.0	14.3	9.8	14.0	6.7
				2.0	14.3	9.8	14.0	6.7
				3.0	14.3	9.8	14.0	6.6
				4.0	14.3	9.8	14.0	6.7
				5.0	14.3	9.8	14.0	6.7
Lake M4	-	12.5	7.0	6.0	14.3	9.8	14.0	6.6
				0.5	13.6	9.8	14.0	6.7
				1.0	13.6	9.8	14.0	6.7
				2.0	13.6	9.8	14.0	6.7
				3.0	13.5	9.9	14.0	6.6
				4.0	13.6	9.8	14.0	6.7
				5.0	13.5	9.8	14.0	6.7
				6.0	13.5	9.8	14.0	6.7
				7.0	13.5	9.8	14.0	6.7
				8.0	13.5	9.8	14.0	6.6
				9.0	13.6	9.5	14.0	6.6
				10.0	13.6	9.5	14.0	6.6
				11.0	14.0	9.5	14.0	6.7



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Table I-1: Water Quality Profiles at Plankton Sampling Locations in East Lake, Lake N11 and the L and M Lakes of the Gahcho Kué Project, Fall 2011 (continued)

Lake	Station	Maximum Depth [m]	Secchi Depth [m]	Depth [m]	Temperature [°C]	Dissolved Oxygen [mg/L]	Specific Conductivity [µS/cm]	pH
Lake L2	-	3.8	3.8	0.3	12.2	10.5	14.0	6.9
				1.0	11.7	10.5	14.0	6.8
				1.5	11.6	10.5	14.0	6.7
				2.0	11.5	10.5	14.0	6.7
				2.5	11.4	10.5	14.0	6.7
				3.0	11.3	10.5	14.0	6.7
				3.5	11.4	9.7	15.0	6.4
Lake N11	D1	6.5	6.0	0.3	16.1	9.4	11.0	6.5
				1.0	16.1	9.3	11.0	6.6
				2.0	16.1	9.3	11.0	6.6
				3.0	16.1	9.3	11.0	6.6
				4.0	16.1	9.3	11.0	6.6
				5.0	16.0	9.3	11.0	6.6
				6.0	15.9	9.3	11.0	6.6
	D2	6.2	5.8	0.3	14.2	9.5	12.0	6.7
				1.0	14.2	9.5	12.0	6.7
				2.0	14.2	0.9	12.0	6.7
				3.0	14.2	0.9	12.0	6.7
				4.0	14.2	0.9	12.0	6.7
				5.0	14.2	0.9	12.0	6.7
	D3	6.5	Bottom	0.3	16.3	9.1	11.0	6.8
				1.0	16.3	9.1	11.0	6.8
				2.0	16.3	9.1	11.0	6.8
				3.0	16.3	9.1	11.0	6.8
				4.0	16.3	9.1	11.0	6.8
				5.0	16.3	9.1	11.0	6.8
				6.0	16.3	9.0	11.0	6.8



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Table I-1: Water Quality Profiles at Plankton Sampling Locations in East Lake, Lake N11 and the L and M Lakes of the Gahcho Kué Project, Fall 2011 (continued)

Lake	Station	Maximum Depth [m]	Secchi Depth [m]	Depth [m]	Temperature [°C]	Dissolved Oxygen [mg/L]	Specific Conductivity [µS/cm]	pH
Lake N11	D4	6.0	5.8	0.3	15.9	9.2	12.0	6.1
				1.0	15.9	9.1	12.0	6.2
				2.0	15.9	9.1	12.0	6.2
				3.0	15.9	9.1	12.0	6.3
				4.0	15.9	9.1	12.0	6.3
				5.0	15.9	9.1	12.0	6.3
	D5	5.6	5.0	0.3	15.3	9.3	12.0	6.3
				1.0	15.3	9.3	12.0	6.3
				2.0	15.3	9.3	12.0	6.3
				3.0	15.3	9.3	12.0	6.3
				4.0	15.3	9.2	12.0	6.3
				5.0	15.3	9.2	12.0	6.3
East Lake (REF)	D1	6.9	Bottom	0.3	15.2	9.3	15.0	6.9
				1.0	15.1	9.3	15.0	6.9
				2.0	15.1	9.3	15.0	6.9
				3.0	15.1	9.3	15.0	6.9
				4.0	15.1	9.3	15.0	6.9
				5.0	15.1	9.3	15.0	6.9
				6.0	15.0	9.3	15.0	6.9
	D2	14.2	7.8	0.3	15.2	9.3	15.0	6.8
				1.0	15.2	9.2	15.0	6.8
				2.0	15.2	9.2	15.0	6.8
				3.0	15.2	9.2	15.0	6.8
				4.0	15.2	9.2	15.0	6.8
				5.0	15.2	9.2	15.0	6.8
				6.0	15.2	9.2	15.0	6.8
				7.0	15.2	9.2	15.0	6.8
				8.0	15.2	9.2	15.0	6.8
				9.0	15.2	9.2	15.0	6.8
				10.0	15.2	9.2	15.0	6.8



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Table I-1: Water Quality Profiles at Plankton Sampling Locations in East Lake, Lake N11 and the L and M Lakes of the Gahcho Kué Project, Fall 2011 (continued)

Lake	Station	Maximum Depth [m]	Secchi Depth [m]	Depth [m]	Temperature [°C]	Dissolved Oxygen [mg/L]	Specific Conductivity [µS/cm]	pH
East Lake (REF)	D2	14.2	7.8	11.0	15.2	9.2	15.0	6.8
				12.0	15.1	9.2	15.0	6.8
				13.0	15.1	9.2	15.0	6.8
				14.0	15.1	9.2	15.0	6.8
	D3	15.3	7.4	0.3	15.1	9.3	15.0	6.8
				1.0	15.1	9.3	15.0	6.8
				2.0	15.1	9.3	15.0	6.8
				3.0	15.1	9.3	15.0	6.8
				4.0	15.1	9.2	15.0	6.8
				5.0	15.1	9.2	15.0	6.8
				6.0	15.1	9.2	15.0	6.8
				7.0	15.1	9.2	15.0	6.8
				8.0	15.1	9.2	15.0	6.8
				9.0	15.1	9.2	15.0	6.8
				10.0	15.1	9.2	15.0	6.8
				11.0	15.1	9.2	15.0	6.8
				12.0	15.1	9.2	15.0	6.8
				13.0	15.0	9.2	15.0	6.8
				14.0	15.0	9.2	15.0	6.8
				15.0	15.0	9.2	15.0	6.8
	D4	9.0	7.5	0.3	15.1	9.1	15.0	6.8
				1.0	15.1	9.1	15.0	6.8
				2.0	15.1	9.1	15.0	6.8
				3.0	15.1	9.1	15.0	6.8
				4.0	15.1	9.1	15.0	6.8
				5.0	15.1	9.1	15.0	6.8
				6.0	15.1	9.1	15.0	6.8
				7.0	15.1	9.1	15.0	6.8
				8.0	15.1	9.1	15.0	6.8



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Table I-1: Water Quality Profiles at Plankton Sampling Locations in East Lake, Lake N11 and the L and M Lakes of the Gahcho Kué Project, Fall 2011 (continued)

Lake	Station	Maximum Depth [m]	Secchi Depth [m]	Depth [m]	Temperature [°C]	Dissolved Oxygen [mg/L]	Specific Conductivity [µS/cm]	pH
East Lake (REF)	D5	14.9	7.8	0.3	15.1	9.2	15.0	6.7
				1.0	15.1	9.2	15.0	6.7
				3.0	15.1	9.2	15.0	6.7
				5.0	15.1	9.1	15.0	6.8
				7.0	15.1	9.1	15.0	6.8
				9.0	15.1	9.1	15.0	6.8
				11.0	15.1	9.1	15.0	6.8
				13.0	15.1	9.1	15.0	6.8
				14.0	15.1	9.1	15.0	6.8

Notes: m = metre; °C = degrees Celsius; mg/L = milligrams per litre; µS/cm = microSiemens per centimetre.

Table 1.1: Phylogenetic Analysis (Fig. 2) (Continued). All 1000 bootstrap values are shown for all nodes.

Phylogenetic Analysis (Fig. 2)	Locus 1C1										Locus 1C2										Locus 1C3										Locus 1C4										Locus 1C5										Locus 1C6										Locus 1C7										Locus 1C8										Locus 1C9										Locus 1C10										Locus 1C11										Locus 1C12										Locus 1C13										Locus 1C14										Locus 1C15										Locus 1C16										Locus 1C17										Locus 1C18										Locus 1C19										Locus 1C20										Locus 1C21										Locus 1C22										Locus 1C23										Locus 1C24										Locus 1C25										Locus 1C26										Locus 1C27										Locus 1C28										Locus 1C29										Locus 1C30										Locus 1C31										Locus 1C32										Locus 1C33										Locus 1C34										Locus 1C35										Locus 1C36										Locus 1C37										Locus 1C38										Locus 1C39										Locus 1C40										Locus 1C41										Locus 1C42										Locus 1C43										Locus 1C44										Locus 1C45										Locus 1C46										Locus 1C47										Locus 1C48										Locus 1C49										Locus 1C50										Locus 1C51										Locus 1C52										Locus 1C53										Locus 1C54										Locus 1C55										Locus 1C56										Locus 1C57										Locus 1C58										Locus 1C59										Locus 1C60										Locus 1C61										Locus 1C62										Locus 1C63										Locus 1C64										Locus 1C65										Locus 1C66										Locus 1C67										Locus 1C68										Locus 1C69										Locus 1C70										Locus 1C71										Locus 1C72										Locus 1C73										Locus 1C74										Locus 1C75										Locus 1C76										Locus 1C77										Locus 1C78										Locus 1C79										Locus 1C80										Locus 1C81										Locus 1C82										Locus 1C83										Locus 1C84										Locus 1C85										Locus 1C86										Locus 1C87										Locus 1C88										Locus 1C89										Locus 1C90										Locus 1C91										Locus 1C92										Locus 1C93										Locus 1C94										Locus 1C95										Locus 1C96										Locus 1C97										Locus 1C98										Locus 1C99										Locus 1C100										Locus 1C101										Locus 1C102										Locus 1C103										Locus 1C104										Locus 1C105										Locus 1C106										Locus 1C107										Locus 1C108										Locus 1C109										Locus 1C110										Locus 1C111										Locus 1C112										Locus 1C113										Locus 1C114										Locus 1C115										Locus 1C116										Locus 1C117										Locus 1C118										Locus 1C119										Locus 1C120										Locus 1C121										Locus 1C122										Locus 1C123										Locus 1C124										Locus 1C125										Locus 1C126										Locus 1C127										Locus 1C128										Locus 1C129										Locus 1C130										Locus 1C131										Locus 1C132										Locus 1C133										Locus 1C134										Locus 1C135										Locus 1C136										Locus 1C137										Locus 1C138										Locus 1C139										Locus 1C140										Locus 1C141										Locus 1C142										Locus 1C143										Locus 1C144										Locus 1C145										Locus 1C146										Locus 1C147										Locus 1C148										Locus 1C149										Locus 1C150										Locus 1C151										Locus 1C152										Locus 1C153										Locus 1C154										Locus 1C155										Locus 1C156										Locus 1C157										Locus 1C158										Locus 1C159										Locus 1C160										Locus 1C161										Locus 1C162										Locus 1C163										Locus 1C164										Locus 1C165										Locus 1C166										Locus 1C167										Locus 1C168										Locus 1C169										Locus 1C170										Locus 1C171										Locus 1C172										Locus 1C173										Locus 1C174										Locus 1C175										Locus 1C176										Locus 1C177										Locus 1C178										Locus 1C179										Locus 1C180										Locus 1C181										Locus 1C182										Locus 1C183										Locus 1C184										Locus 1C185										Locus 1C186										Locus 1C187										Locus 1C188										Locus 1C189										Locus 1C190										Locus 1C191										Locus 1C192										Locus 1C193										Locus 1C194										Locus 1C195										Locus 1C196										Locus 1C197										Locus 1C198										Locus 1C199										Locus 1C200										Locus 1C201										Locus 1C202										Locus 1C203										Locus 1C204										Locus 1C205										Locus 1C206										Locus 1C207										Locus 1C208										Locus 1C209										Locus 1C210										Locus 1C211										Locus 1C212										Locus 1C213										Locus 1C214										Locus 1C215										Locus 1C216										Locus 1C217										Locus 1C218										Locus 1C219										Locus 1C220										Locus 1C221										Locus 1C222										Locus 1C223										Locus 1C224										Locus 1C225										Locus 1C226										Locus 1C227										Locus 1C228										Locus 1C229										Locus 1C230										Locus 1C231										Locus 1C232										Locus 1C233										Locus 1C234										Locus 1C235										Locus 1C236										Locus 1C237										Locus 1C238										Locus 1C239										Locus 1C240										Locus 1C241										Locus 1C242										Locus 1C243										Locus 1C244										Locus 1C245										Locus 1C246										Locus 1C247										Locus 1C248										Locus 1C249										Locus 1C250										Locus 1C251										Locus 1C252										Locus 1C253										Locus 1C254										Locus 1C255										Locus 1C256										Locus 1C257										Locus 1C258										Locus 1C259										Locus 1C260										Locus 1C261										Locus 1C262										Locus 1C263										Locus 1C264										Locus 1C265										Locus 1C266										Locus 1C267										Locus 1C268										Locus 1C269										Locus 1C270										Locus 1C271										Locus 1C272										Locus 1C273										Locus 1C274										Locus 1C275										Locus 1C276										Locus 1C277										Locus 1C278										Locus 1C279										Locus 1C280										Locus 1C281										Locus 1C282										Locus 1C283										Locus 1C284										Locus 1C285										Locus 1C286										Locus 1C287										Locus 1C288										Locus 1C289										Locus 1C290										Locus 1C291										Locus 1C292										Locus 1C293										Locus 1C294										Locus 1C295										Locus 1C296										Locus 1C297										Locus 1C298										Locus 1C299										Locus 1C300										Locus 1C301										Locus 1C302										Locus 1C303										Locus 1C304										Locus 1C305										Locus 1C306										Locus 1C307										Locus 1C308										Locus 1C309										Locus 1C310										Locus 1C311										Locus 1C312										Locus 1C313										Locus 1C314										Locus 1C315										Locus 1C316										Locus 1C317										Locus 1C318										Locus 1C319										Locus 1C320										Locus 1C321										Locus 1C322										Locus 1C323										Locus 1C324										Locus 1C325										Locus 1C326										Locus 1C327										Locus 1C328										Locus 1C329										Locus 1C330										Locus 1C331										Locus 1C332										Locus 1C333										Locus 1C334										Locus 1C335										Locus 1C336										Locus 1C337										Locus 1C338										Locus 1C339										Locus 1C340										Locus 1C341										Locus 1C342										Locus 1C343										Locus 1C344										Locus 1C345										Locus 1C346										Locus 1C347										Locus 1C348										Locus 1C349										Locus 1C350										Locus 1C351										Locus 1C352										Locus 1C353										Locus 1C354										Locus 1C355										Locus 1C356										Locus 1C357										Locus 1C358										Locus 1C359										Locus 1C360										Locus 1C361										Locus 1C362										Locus 1C363										Locus 1C364										Locus 1C365										Locus 1C366										Locus 1C367										Locus 1C368										Locus 1C369										Locus 1C370										Locus 1C371										Locus 1C372										Locus 1C373										Locus 1C374										Locus 1C375										Locus 1C376										Locus 1C377										Locus 1C378										Locus 1C379										Locus 1C380										Locus 1C381										Locus 1C382										Locus 1C383										Locus 1C384										Locus 1C385										Locus 1C386										Locus 1C387										Locus 1C388										Locus 1C389										Locus 1C390										Locus 1C391										Locus 1C392										Locus 1C393										Locus 1C394										Locus 1C395										Locus 1C396										Locus 1C397										Locus 1C398										Locus 1C399										Locus 1C400										Locus 1C401										Locus 1C402										Locus 1C403										Locus 1C404										Locus 1C405										Locus 1C406										Locus 1C407										Locus 1C408										Locus 1C409										Locus 1C410										Locus 1C411										Locus 1C412										Locus 1C413										Locus 1C414										Locus 1C415										Locus 1C416										Locus 1C417										Locus 1C418										Locus 1C419										Locus 1C420										Locus 1C421										Locus 1C422										Locus 1C423										Locus 1C424										Locus 1C425										Locus 1C426										Locus 1C427										Locus 1C428										Locus 1C429										Locus 1C430										Locus 1C431										Locus 1C432										Locus 1C433										Locus 1C434										Locus 1C435										Locus 1C436										Locus 1C437										Locus 1C438										Locus 1C439										Locus 1C440										Locus 1C441										Locus 1C442										Locus 1C443										Locus 1C444										Locus 1C445										Locus 1C446										Locus 1C447										Locus 1C448										Locus 1C449										Locus 1C450										Locus 1C451										Locus 1C452										Locus 1C453										Locus 1C454										Locus 1C455										Locus 1C456										Locus 1C457										Locus 1C458										Locus 1C459										Locus 1C460										Locus 1C461										Locus 1C462										Locus 1C463										Locus 1C464										Locus 1C465										Locus 1C466										Locus 1C467										Locus 1C468										Locus 1C469										Locus 1C470										Locus 1C471										Locus 1C472										Locus 1C473										Locus 1C474										Locus 1C475										Locus 1C476										Locus 1C477										Locus 1C478										Locus 1C479										Locus 1C480										Locus 1C481										Locus 1C482										Locus 1C483										Locus 1C484										Locus 1C485										Locus 1C486										Locus 1C487										Locus 1C488										Locus 1C489										Locus 1C490										Locus 1C491										Locus 1C492										Locus 1C493										Locus 1C494										Locus 1C495										Locus 1C496										Locus 1C497										Locus 1C498										Locus 1C499										Locus 1C500										Locus 1C501										Locus 1C502										Locus 1C503										Locus 1C504										Locus 1C505										Locus 1C506										Locus 1C507										Locus 1C508										Locus 1C509										Locus 1C510										Locus 1C511										Locus 1C512										Locus 1C513										Locus 1C514										Locus 1C515										Locus 1C516										Locus 1C517										Locus 1C518										Locus 1C519										Locus 1C520										Locus 1C521										Locus 1C522										Locus 1C523										Locus 1C524										Locus 1C525										Locus 1C526										Locus 1C527										Locus 1C528										Locus 1C529										Locus 1C530										Locus 1C531										Locus 1C532										Locus 1C533										Locus 1C534										Locus 1C535										Locus 1C536										Locus 1C537										Locus 1C538										Locus 1C539										Locus 1C540										Locus 1C541										Locus 1C542										Locus 1C543										Locus 1C544										Locus 1C545										Locus 1C546										Locus 1C547										Locus 1C548										Locus 1C549										Locus 1C550										Locus 1C551										Locus 1C552										Locus 1C553										Locus 1C554										Locus 1C555										Locus 1C556										Locus 1C557										Locus 1C558										Locus 1C559										Locus 1C560										Locus 1C561										Locus 1C562										Locus 1C563										Locus 1C564										Locus 1C565										Locus 1C566										Locus 1C567										Locus 1C568										Locus 1C569										Locus 1C570										Locus 1C571										Locus 1C572										Locus 1C573										Locus 1C574										Locus 1C575										Locus 1C576										Locus 1C577										Locus 1C578										Locus 1C579										Locus 1C580										Locus 1C581										Locus 1C582										Locus 1C583										Locus 1C584										Locus 1C585										Locus 1C586										Locus 1C587										Locus 1C588										Locus 1C589										Locus 1C590										Locus 1C591										Locus 1C592										Locus 1C593										Locus 1C594										Locus 1C595										Locus 1C596										Locus 1C597										Locus 1C598										Locus 1C599										Locus 1C600										Locus 1C601										Locus 1C602										Locus 1C603										Locus 1C604										Locus 1C605										Locus 1C606										Locus 1C607										Locus 1C608										Locus 1C609										Locus 1C610										Locus 1C611										Locus 1C612										Locus 1C613										Locus 1C614										Locus 1C615										Locus 1C616										Locus 1C617										Locus 1C618										Locus 1C619										Locus 1C620										Locus 1C621										Locus 1C622										Locus 1C623										Locus 1C624										Locus 1C625										Locus 1C626										Locus 1C627										Locus 1C628										Locus 1C629										Locus 1C630										Locus 1C631										Locus 1C632										Locus 1C633										Locus 1C634										Locus 1C635										Locus 1C636										Locus 1C637										Locus 1C638										Locus 1C639										Locus 1C640										Locus 1C641										Locus 1C642										Locus 1C643										Locus 1C644										Locus 1C645										Locus 1C646										Locus 1C647										Locus 1C648										Locus 1C649										Locus 1C650										Locus 1C651										Locus 1C652										Locus 1C653										Locus 1C654										Locus 1C655										Locus 1C656										Locus 1C657										Locus 1C658										Locus 1C659										Locus 1C660										Locus 1C661										Locus 1C662										Locus 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Table 1. Supplemental Monitoring Data for Liver Trends and Risk Assessment of the Hudson River Fishery

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[illegible]



Table I-6: Zooplankton Quality Control Samples Base on Abundance (ind./L) in East Lake and Lake N11, Fall 2011

Lake	Lake N11			Lake N11			Lake N11			East Lake (Reference)			East Lake (Reference)			East Lake (Reference)			East Lake (Reference)		
Date	15-Aug-11			15-Aug-11			15-Aug-11			17-Aug-11			17-Aug-11			17-Aug-11			17-Aug-11		
Station Replicates	N11D1A			N11D3C			N11D3E			REFD3C			REFD3D			REFD4A			REFD5A		
Stations	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)
COPEPODA																					
Calanoida																					
<i>Hetercope septentrionalis</i> Juday & Muttkowski																					
H.s. adult female	0	0	-	0	0	-	0	0	-	0.03	0.03	0	0.02	0.02	0	0.01	0.01	0	0.03	0.03	0
H.s. adult male	0	0	-	0	0	-	0.01	0.01	0	0.01	0.01	0	0.01	0.01	0	0.01	0.01	0	0.01	0.01	0
H.s. 4.0 mm F & M	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
H.s. 3.0mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
H.s. 2.0 mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
H.s. 1.0 mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
Total <i>H. septentrionales</i>	0	0	-	0	0	-	0.01	0.01	0	0.04	0.04	0	0.03	0.03	0	0.02	0.02	0	0.04	0.04	0
<i>Epischura lacustris</i> S.A. Forbes																					
E.l. adult female	0.00	0.00	0	0.00	0.00	0	0	0	-	0.01	0.01	0.0	0.04	0.01	38.9	0	0.00	50.0	0.05	0.02	16.7
E.l. adult male	0	0	-	0.00	0.00	0	0.00	0.00	0	0.00	0.00	0.2	0	0.01	50.0	0	0	-	0.05	0.02	16.7
E.l. immature 0.5-1.0 mm	0	0	-	0	0	-	0	0	-	0		-	0	0	-	0	0	-	0	0	-
Total <i>E. lacustris</i>	0.00	0.00	0	0.01	0.01	0	0.00	0.00	0	0.01	0.01	0.1	0.04	0.01	24.5	0	0.00	50.0	0.10	0.05	16.7
<i>Diaptomus pribilofensis</i> Juday & Muttkowski																					
D.p. adult female	0.10	0.10	0	0.12	0.17	8.3	0.02	0.11	32.1	0.69	0.49	8.6	0.49	0.53	2.0	0.64	0.57	2.9	1.47	1.13	6.5
D.p. gravid female	0.01	0.01	0	0.00	0.00	0	0.00	0.00	0	0.04	0.02	16.7	0.02	0.02	0	0.02	0.02	2.0	0.05	0.02	16.7
D.p. adult male	0.31	0.41	7.1	0.05	0.07	10.0	0.20	0.07	22.7	0.65	0.32	16.7	0.61	0.53	3.6	0.14	0.64	31.8	0.45	1.13	21.4
D.p. immature 2.0 mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
D.p. immature 1.0 mm	0.21	0	50.0	0.57	0.11	33.3	0.11	0.11	0	0.04	0.04	0	0	0.12	50.0	0	0.07	50.0	0	0.11	50.0
D.p. immature 0.75 mm	0	0	-	0	0	-	0	0	-	0	0.04	50.0	0	0	-	0	0	-	0	0	-
D.p. immature 0.5 mm	0	0	-	0	0	-	0	0	-	0	0	-	0.04	0	50.0	0.07	0	50.0	0	0	-
Total <i>D. pribilofensis</i>	0.63	0.52	4.5	0.74	0.36	17.2	0.34	0.30	2.7	1.41	0.91	10.8	1.15	1.19	0.9	0.87	1.30	9.8	1.97	2.40	4.9
<i>Diaptomus minutus</i> Lilljeborg																					
D.m. adult female	0.10	0.41	30.0	0.45	0	-	0.11	0.11	0	0	0	-	0	0	-	0	0	-	0	0.11	50.0
D.m. gravid female	0.13	0.07	16.7	0.02	0.02	0	0.10	0.22	19.2	0	0	-	0	0	-	0	0	-	0.02	0	50.0
D.m. adult male	0.62	0.93	10.0	0.57	0.57	0	0.34	0.23	10.0	0	0	-	0	0	-	0	0	-	0	0	-
D.m. immature 2.0 mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
D.m. immature 1.0 mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
D.m. immature 0.75 mm	0.51	0.51	0	0.34	0.79	20.0	0	0.57	50.0	1.50	1.13	6.9	1.37	1.09	5.7	0.78	1.06	7.7	2.49	2.04	5.0



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Table I-6: Zooplankton Quality Control Samples Base on Abundance (ind./L) in East Lake and Lake N11, Fall 2011 (continued)

Lake	Lake N11			Lake N11			Lake N11			East Lake (Reference)			East Lake (Reference)			East Lake (Reference)			East Lake (Reference)		
Date	15-Aug-11			15-Aug-11			15-Aug-11			17-Aug-11			17-Aug-11			17-Aug-11			17-Aug-11		
Station Replicates	N11D1A			N11D3C			N11D3E			REFD3C			REFD3D			REFD4A			REFD5A		
Stations	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)
D.m. immature 0.5 mm	0.10	0	50.0	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
Total <i>D. minutus</i>	1.47	1.92	6.6	1.38	1.38	0	0.55	1.13	17.2	1.50	1.13	6.9	1.37	1.09	5.7	0.78	1.06	7.7	2.52	2.15	3.9
Calanoid nauplius	0.21	0.10	16.7	0.11	0.11	0	0	0	-	0	0.04	50.0	0	0	-	0	0	-	0	0.11	50.0
Total Calanoida ind/L	2.31	2.55	2.5	2.24	1.86	4.6	0.90	1.44	11.6	2.96	2.13	8.1	2.60	2.33	2.7	1.67	2.38	8.7	4.63	4.76	0.7
Cyclopoida																					
<i>Cyclops scutifer</i> Sars																					
C. s. adult female	0.21	0.10	16.7	0	0.11	50.0	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
C. s. gravid female	0.01	0.01	0	0.00	0.00	0	0.02	0.02	0	0	0	-	0	0	-	0	0	-	0	0	-
C. s. adult male	0	0	-	0.11	0	50.0	0	0.11	50.0	0	0	-	0	0	-	0	0	-	0	0	-
C. s. immature 2.0 mm	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
C. s. immature 1.0 mm	0.31	0.41	7.1	0	0	-	0	0	-	0.32	0	50.0	0	0	-	0	0	-	0	0.11	50.0
C. s. immatue 0.75 mm	0.82	1.65	16.7	0.34	0.11	25.0	0.11	0.23	16.7	0.97	0.69	8.5	1.29	0.57	19.6	0.57	0.35	11.5	2.49	1.81	7.9
C. s. immature 0.5 mm	0.10	0	50.0	0	0	-	0	0	-	0.36	0.40	2.6	0.32	0.20	11.5	0.14	0.14	0.0	0.57	0.57	0
Total <i>C. scutifer</i>	1.45	2.17	10.0	0.46	0.23	16.5	0.14	0.36	22.5	1.66	1.09	10.3	1.62	0.77	17.8	0.71	0.50	8.8	3.06	2.49	5.1
Small unidentified cyclopoids	1.45	0.21	37.5	0.11	0.11	0			-			-			-			-			-
Cyclopoid nauplius	1.34	2.37	13.9	2.94	2.15	7.8	1.81	1.59	3.3	5.30	3.92	7.5	4.69	4.73	0.2	6.23	6.23	0	6.91	8.27	4.5
Total Cyclopoida ind/L	2.78	4.74	13.0	3.51	2.49	8.5	1.95	1.95	0.0	6.96	5.01	8.1	6.31	5.50	3.4	6.94	6.72	0.8	9.96	10.76	1.9
CLADOCERA																					
<i>Daphnia middendorffiana</i> Fischer																					
D. m. 3.0	0	0	-	0	0	-	0	0	-	0	0	-	0.00	0.00	0	0	0	-	0	0	-
D. m. 2.5	0	0	-	0	0	-	0	0	-	0.00	0.00	0	0.01	0.01	3.3	0.00	0.00	0	0.03	0.02	2.9
D. m. 2.0	0	0	-	0	0	-	0	0	-	0.01	0.01	1.9	0.01	0.01	8.3	0	0	-	0.01	0.01	0
D. m. 1.5	0	0	-	0	0	-	0	0	-	0.01	0.02	16.7	0.02	0.01	16.7	0.00	0.00	0	0.01	0.01	10.0
D. m. 1.0	0	0	-	0	0	-	0	0	-	0.01	0.01	0	0	0	-	0.00	0.00	0	0.02	0	50.0
D. m. 0.5	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
Total <i>D. middendorfianna</i>	0	0	-	0	0	-	0	0	-	0.03	0.04	6.5	0.03	0.02	7.0	0.01	0.01	0	0.07	0.05	10.7
<i>Holopedium gibberum</i> Zaddach																		-			
H. g. 3.0	0	0	-	0	0	-	0	0	-	0.01	0.01	0	0.01	0.01	0	0	0	-	0.06	0.06	2.4
H. g. 2.0	4.43	4.01	2.4	0.32	0.44	8.1	0.17	0.27	11.1	0.01	0.01	0	0.01	0.01	2.9	0	0	-	0.05	0.04	3.3
H. g. 1.0	0.82	1.24	10.0	0.10	0.32	26.5	0.05	0.10	16.7	0	0	-	0	0	-	0	0	-	0	0	-
H. g. 0.5	1.54	1.44	1.7	0.91	1.13	5.6	0.57	1.25	18.8	0	0	-	0	0	-	0	0	-	0	0	-
Total <i>H. gibberum</i>	6.79	6.69	0.4	1.32	1.90	8.9	0.79	1.62	17.2	0.02	0.02	0	0.02	0.01	1.7	0	0	-	0.11	0.10	2.8



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Table I-6: Zooplankton Quality Control Samples Base on Abundance (ind./L) in East Lake and Lake N11, Fall 2011 (continued)

Lake	Lake N11			Lake N11			Lake N11			East Lake (Reference)			East Lake (Reference)			East Lake (Reference)			East Lake (Reference)		
Date	15-Aug-11			15-Aug-11			15-Aug-11			17-Aug-11			17-Aug-11			17-Aug-11			17-Aug-11		
Station Replicates	N11D1A			N11D3C			N11D3E			REFD3C			REFD3D			REFD4A			REFD5A		
Stations	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)	A	B	RPD (%)
<i>Eubosmina longispina</i>																					
E. l. 1.0	0.02	0	50.0	0.00	0.00	0	0	0	-	0	0	-	0.00	0.00	0	0	0	-	0	0	-
E. l. 0.75	0.62	0.62	0	0.05	0.02	16.7	0	0	-	0.00	0.00	16.7	0	0.01	50.0	0	0	-	0.01	0.02	13.5
E. l. 0.5	3.81	2.16	13.8	0	1.13	50.0	0.23	1.25	34.6	0.00	0.00	0	0.01	0.01	0	0.00	0.00	-	0.00	0	50.0
E. l. 0.25	0.62	1.44	20.0	0	0.45	50.0	0.11	0.23	16.7	0	0	-	0	0	-	0	0	-	0	0	-
E. l. male	0	0.10	50.0	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-	0	0	-
Total <i>E. longispina</i>	5.07	4.32	4.0	0.05	1.61	46.9	0.34	1.47	31.3	0.00	0.00	7.1	0.01	0.02	15.5	0.00	0.00	0	0.02	0.02	9.2
Total Cladocera ind/L	11.86	11.01	1.8	1.38	3.51	21.8	1.13	3.09	23.2	0.06	0.07	4.4	0.06	0.06	0.0	0.01	0.01	0	0.19	0.17	3.9
ROTIFERA																					
<i>Kellicottia</i> spp.	8.75	8.24	1.5	13.02	13.81	1.5	10.42	9.85	1.4	2.02	2.67	6.9	1.94	3.11	11.6	6.16	5.02	5.1	3.96	3.85	0.7
<i>Keratella</i> spp.	3.50	3.91	2.8	7.02	8.38	4.4	5.77	5.21	2.6	0.49	0.57	3.8	0.65	0.69	1.5	0.35	0.71	16.7	0.91	1.13	5.6
<i>Polyarthra</i> spp.	0.21	0	50.0	0.68	0.91	7.1	0.79	0.34	20.0	0	0	-	0	0	-	0.07	0.07	0	0	0	-
<i>Conochilus</i> spp.	6.38	6.79	1.6	15.51	18.23	4.0	13.14	15.97	4.9	1.09	0.77	8.7	0.57	0.77	7.6	1.42	0.78	14.5	3.74	4.30	3.5
<i>Lecane</i> spp.	0	0	-	0	0	-	0	0.11	50.0	0	0	-	0	0	-	0	0	-	0	0	-
<i>Pleosoma</i> spp.	0	0.10	50.0		0.11	50.0			-			-			-			-			-
<i>Synchaeta</i> spp.			-			-			-	0.04		50.0	0.04	0.04	0		0.07	50.0			-
Total Rotifera ind/L	18.84	19.04	0.3	36.23	41.44	3.4	30.12	31.48	1.1	3.64	4.00	2.4	3.19	4.61	9.1	8.00	6.65	4.6	8.61	9.29	1.9
TOTAL ZOOPLANKTON	35.79	37.35	1.1	43.37	49.31	3.2	34.10	37.95	2.7	13.61	11.22	4.8	12.16	12.49	0.7	16.62	15.77	1.3	23.39	24.96	1.6

Notes: ind./L = Individuals per litre; sp. = one species in the genera; spp. = numerous species in the genera; mm = millimetres; RPD = Relative Percent Difference



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Table I-7: Zooplankton Biomass Length-Weight Regression Information for East Lake, Lake N11 and the L and M Lakes of the Gahcho Kue Project, Fall 2011

Instar Identification and Size Classes	Mean Length [mm]	Ln ^(a) (Length) [mm]	Regression Equation Number ^(b)	Calculated Ln (Weight) [µg]	Calculated Dry Weight [µg]	Calculated Wet Weight [µg]
<i>Heterocope septentrionalis</i> Juday and Muttkowski						
H.s. adult female	2.7	1.0	R30	3.5	34.2	488.8
H.s. adult male	3.1	1.1	R30	3.8	45.4	649.0
H.s. 4.0 mm (male and female)	3.3	1.2	R30	4.0	52.9	756.2
H.s. 3.0mm	3.0	1.1	R30	3.8	43.2	617.2
H.s. 2.0 mm	1.8	0.6	R30	2.4	11.0	157.1
H.s. 1.0 mm	0.8	-0.3	R30	0.2	1.3	18.3
<i>Epischura lacustris</i> S.A. Forbes						
E.l. adult female	1.5	0.4	R32	1.9	6.6	94.2
E.l. adult male	1.4	0.3	R32	2.1	7.9	113.4
E. l. immature	1.0	0.0	R32	1.7	5.2	74.7
<i>Diaptomus pribilofensis</i> Juday and Muttkowski						
D.p. adult female	1.2	0.2	R30	1.4	4.0	57.8
D.p. gravid female	1.2	0.2	R30	1.4	4.0	57.8
D.p. adult male	1.2	0.2	R30	1.4	3.9	55.6
D.p. immature 2.0	1.2	0.1	R30	1.3	3.8	54.7
D.p. immature 1.0	1.0	0.0	R30	1.0	2.6	37.0
D.p. immature 0.75	0.8	-0.3	R30	0.2	1.3	18.3
D.p. immature 0.5	0.5	-0.7	R30	-0.8	0.5	6.5
<i>Diaptomus minutus</i> Lilljeborg						
D.m. adult female	0.9	-0.1	R27	0.9	2.5	35.1
D.m. gravid female	0.9	-0.1	R27	0.9	2.4	34.6
D.m. adult male	0.9	-0.1	R27	0.7	2.1	29.9
D.m. immature 2.0	1.1	0.0	R27	1.2	3.3	46.9
D.m. immature 1.0	1.0	0.0	R27	1.0	2.8	39.9
D.m. immature 0.75	0.8	-0.3	R27	0.3	1.3	18.6
D.m. immature 0.5	0.5	-0.7	R27	-0.9	0.4	6.1
Calanoid nauplius	0.3	-1.2	R6	-1.6	0.2	2.9
<i>Cyclops scutifer</i> Sars						
C. s. adult female	1.3	0.2	R94	2.0	7.5	107.0
C. s. gravid female	1.3	0.3	R94	2.1	8.3	118.2
C. s. adult male	1.0	0.0	R94	1.4	3.9	55.6



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Table I-7: Zooplankton Biomass Length-Weight Regression Information for East Lake, Lake N11 and the L and M Lakes of the Gahcho Kue Project, Fall 2011 (continued)

Instar Identification and Size Classes	Mean Length [mm]	Ln ^(a) (Length) [mm]	Regression Equation Number ^(b)	Calculated Ln (Weight) [µg]	Calculated Dry Weight [µg]	Calculated Wet Weight [µg]
C. s. immature 2.0	1.2	0.1	R94	1.7	5.6	80.0
C. s. immature 1.0	1.0	0.0	R94	1.3	3.6	51.8
C. s. immature 0.75	0.8	-0.3	R94	0.5	1.7	23.9
C. s. immature 0.5	0.5	-0.7	R94	-0.6	0.5	7.7
<i>Cyclops vernalis</i> Fischer						
C.v. immature 0.75 mm	0.5	-0.7	R92	-1.0	0.4	5.5
<i>Cyclops capillatus</i> Sars						
C.c. mature	1.6	0.5	R92	2.1	8.2	117.7
C. c. immature	0.5	-0.7	R92	-1.0	0.4	5.5
<i>Macrocyclops albidus</i> Jurine	1.7	0.5	R92	2.1	8.4	119.5
Immature cyclopoid	0.5	-0.7	R92	-1.0	0.4	5.5
Cyclopoid nauplius	0.3	-1.3	R49	-1.6	0.2	2.7
<i>Daphnia middendorffiana</i> Fischer						
D. m. 3.0	3.0	1.1	DsL885	4.7	110.1	1573.3
D. m. 2.5	2.5	0.9	DsL885	4.2	63.6	908.6
D. m. 2.0	2.0	0.7	DsL885	3.5	32.5	464.0
D. m. 1.5	1.5	0.4	DsL885	2.6	13.7	195.1
D. m. 1.0	1.0	0.0	DsL885	1.4	3.9	55.8
D. m. 0.5	0.5	-0.7	DsL885	-0.7	0.5	7.1
<i>Holopedium gibberum</i> Zaddach						
H. g. 3.0	2.5	0.9	L223Hg	4.6	98.3	1404.7
H. g. 2.0	1.8	0.6	L223Hg	3.6	37.6	536.8
H. g. 1.0	1.0	0.0	L223Hg	2.1	8.1	115.5
H. g. 0.5	0.5	-0.7	L223Hg	0.2	1.3	18.3
<i>Eubosmina longispina</i>						
E. l. 1.0	1.0	0.0	L223BI	2.4	11.5	164.1
E. l. 0.75	0.8	-0.3	L223BI	1.5	4.5	64.5
E. l. 0.5	0.5	-0.7	L223BI	0.1	1.2	16.5
E. l. 0.25	0.3	-1.4	L223BI	-2.2	0.1	1.6
E. l. male	0.5	-0.7	L223BI	0.1	1.2	16.5
<i>Daphnia longiremis</i> Sars						
D. l. 1.3mm	1.3	0.3	L302	2.5	12.2	174.5



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Table I-7: Zooplankton Biomass Length-Weight Regression Information for East Lake, Lake N11 and the L and M Lakes of the Gahcho Kue Project, Fall 2011 (continued)

Instar Identification and Size Classes	Mean Length [mm]	Ln ^(a) (Length) [mm]	Regression Equation Number ^(b)	Calculated Ln (Weight) [µg]	Calculated Dry Weight [µg]	Calculated Wet Weight [µg]
D.I. 1.0 mm	1.0	0.0	L302	1.6	4.9	70.3
D.I. 0.5 mm	0.5	-0.8	L302	-1.0	0.4	5.1
D.I.male 1.2 mm	1.2	0.2	L302	2.2	9.4	133.6
<i>Ophryoxus gracilis</i> Sars	1.1	0.1	L223BI	2.7	14.9	213.1
<i>Eurycerus lamellatus</i> (O.F. Muller)	2.1	0.7	L223Cs	5.5	243.4	3477.8
<i>Chydorus sphaericus</i> (O.F.Muller)	0.5	-0.8	L223Cs	-0.7	0.5	7.4
<i>Kellicottia</i> spp.	0.1	-2.1	L224	-	0.0	0.2
<i>Keratella</i> spp.	0.1	-2.3	L224	-	0.0	0.2
<i>Polyarthra</i> spp.	0.1	-2.0	L227*	-	0.1	0.8
<i>Conochilus</i> spp.	0.2	-1.7	L223	-	0.0	0.6
<i>Lecane</i> spp.	0.1	-2.3	L227	-	0.0	0.5
<i>Pleosoma</i> spp.	0.2	-1.5	L224*	-	0.1	1.0
<i>Synchaeta</i> spp.	0.1	-2.1	L227*	-	0.0	0.7

^(a) Length/DryWeight Regressions in form $\text{LnW} = \text{LnL} + b\text{LnL}$ from Lawrence et al. 1989.

R6 $\text{LnW} = 0.9926 - 2.0997 \text{ LnL}$.

R27 $\text{LnW} = 1.0542 - 2.748 \text{ LnL}$.

R30 $\text{LnW} = 0.9772 - 2.5384 \text{ LnL}$.

R32 $\text{LnW} = 1.1337 + 2.7882 \text{ LnL}$.

R49 $\text{LnW} = 1.6388 - 2.4474 \text{ LnL}$.

R92 $\text{LnW} = 0.8344 - 2.5760 \text{ LnL}$.

R94 $\text{LnW} = 1.3169 - 2.7197 \text{ LnL}$.

DsL885 $\text{LnW} = 1.3933 - 3.0114 \text{ LnL}$.

RL302 $\text{LnW} = 1.6274 - 3.3367 \text{ LnL}$.

RL223Hg $\text{LnW} = 2.1169 + 2.6972 \text{ LnL}$.

RL223BI $\text{LnW} = 2.4751 - 3.3614 \text{ LnL}$.

RL223Cs $\text{LnW} = 3.1270 - 3.3678 \text{ LnL}$.

^(b) * = interpolated.

Notes: mm= millimeters; µg/L = micrograms per litre; spp = numerous species in the genera; - = information not available.



APPENDIX II

Benthic Invertebrates - Supporting Data, 2011



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Table II-1: Habitat Data for Benthic Invertebrate Stations Sampled in East Lake, Lake N11, and the L and M Lakes of the Gahcho Kue Project, Fall 2011

Lake	Station	Date	UTM Coordinates (Zone 12V, NAD83)		Water Depth [m]	Field Water Quality Data								Sediment Chemistry Data					
						Water Temperature [°C]		Dissolved Oxygen [mg/L]		Specific Conductivity [µS/cm]		pH		Moisture Content [%]	Total Organic Carbon [%]	Sediment Particle Size			
			Easting	Northing		Surface	Bottom	Surface	Bottom	Surface	Bottom	Surface	Bottom			Sand [%]	Silt [%]	Clay [%]	Fines (silt + clay) [%]
East (Reference) Lake	EAST-D1	17-Aug-2011	597732	7038735	7.4 (reps A & B) 11.3 (reps C, D & E)	15.2	15.0	9.3	9.3	15.0	15.0	6.9	6.9	71	3	86	11	4	15
	EAST-D2	17-Aug-2011	598225	7039182	14.2	15.2	15.1	9.3	9.2	15.0	15.0	6.8	6.8	90	10	62	30	8	38
	EAST-D3	17-Aug-2011	598834	7040563	15.3	15.1	15.0	9.3	9.2	15.0	15.0	6.8	6.8	92	13	77	15	8	23
	EAST-D4	17-Aug-2011	599450	7041433	9.0	15.1	15.1	9.1	9.1	15.0	15.0	6.8	6.8	92	13	87	6	8	14
	EAST-D5	17-Aug-2011	599191	7040941	14.9	15.1	15.1	9.2	9.1	15.0	15.0	6.7	6.8	92	11	77	16	6	22
Lake N11	N11-D1	13-Aug-2011	587367	7040304	6.5	16.1	15.9	9.2	9.2	11.0	11.0	6.5	6.6	92	13	77	16	7	23
	N11-D2	16-Aug-2011	586893	7040510	6.2	14.2	14.2	9.5	9.4	12.0	12.0	6.7	6.7	91	9	74	22	4	26
	N11-D3	15-Aug-2011	587460	7041119	6.1	16.3	16.3	9.1	9.0	11.0	11.0	6.8	6.8	90	16	78	17	5	22
	N11-D4	14-Aug-2011	587543	7041902	6.1	15.9	15.9	9.2	9.1	12.0	12.0	6.1	6.3	87	16	75	18	8	26
	N11-D5	15-Aug-2011	586753	7042017	6.0	15.3	15.3	9.3	9.2	12.0	12.0	6.3	6.3	92	16	72	19	10	29
Lake M1	M1	20-Aug-2011	596462	7044978	2.0	9.6	9.6	10.6	10.6	14.0	14.0	6.8	6.7	-	-	-	-	-	-
Lake M2	M2	18-Aug-2011	597016	7044841	5.0	13.5	13.5	9.8	9.8	14.0	14.0	6.7	6.4	-	-	-	-	-	-
Lake M3	M3	18-Aug-2011	597452	7043954	7.0	13.6	14.3	9.8	9.8	14.0	14.0	6.6	6.7	90	13	63	24	14	38
Lake M4	M4	20-Aug-2011	595252	7040163	12.0	13.6	14.0	9.8	9.8	14.0	14.0	6.7	6.7	92	15	67	28	5	33
Lake L2	L2	20-Aug-2011	593396	7038967	3.8	12.2	11.4	10.5	9.7	14.0	15.0	6.9	6.4	-	-	-	-	-	-

Notes: - = not available.
Benthic invertebrates were sampled at two locations for station EAST-D1 because the first location did not have sufficient soft bottom substrate to sample more than two replicates.



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Table II-2: Benthic Invertebrate Summary Variables for East Lake, Lake N11, and the L and M Lake of the Gahcho Kue Project, Fall 2011

Lake	Habitat Type	Station	Total Density (no./m ²) (mean \pm 1 SE)			Total Richness (taxa/station)	Mean Total Richness (taxa/station) (mean \pm 1 SE)			Simpson's Diversity Index	Evenness
East (Reference) Lake	Deep Open-Water	EAST-D1	1,034	\pm	337	21	9	\pm	3	0.90	0.49
		EAST-D2	621	\pm	325	14	6	\pm	2	0.86	0.53
		EAST-D3	3,440	\pm	749	23	14	\pm	1	0.83	0.26
		EAST-D4	5,207	\pm	1,428	27	17	\pm	1	0.87	0.29
		EAST-D5	8,914	\pm	1,418	25	18	\pm	1	0.87	0.32
Lake N11	Deep Open-Water	N11-D1	25,190	\pm	4,158	25	18	\pm	1	0.67	0.12
		N11-D2	37,095	\pm	8,732	29	19	\pm	2	0.72	0.12
		N11-D3	20,819	\pm	4,857	29	19	\pm	2	0.79	0.17
		N11-D4	810	\pm	316	18	7	\pm	2	0.83	0.33
		N11-D5	53,776	\pm	17,303	33	22	\pm	3	0.52	0.06
Lake M1	Shallow Open-Water	M1	14,095	\pm	2,554	30	19	\pm	2	0.91	0.39
Lake M2	Shallow Open-Water	M2	13,595	\pm	4,769	26	17	\pm	2	0.87	0.29
Lake M3	Shallow Open-Water	M3	6,345	\pm	1,583	32	18	\pm	4	0.90	0.30
Lake M4	Shallow Open-Water	M4	241	\pm	84	9	3	\pm	1	0.81	0.59
Lake L2	Shallow Open-Water	L2	8,897	\pm	3,331	26	15	\pm	2	0.88	0.33

Notes: SE = standard error of the mean.

Deep Open-Water = open-water areas with water depths ranging from 6 to 10 m.

Shallow Open-Water = open-water areas with water depths less than 4 metres.



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Table II-3: Percent Mean Relative Density of Major Taxa in East Lake, Lake N11, and the L and M Lakes Sampled for the Gahcho Kue Project, Fall 2011

Taxa	Reference Lake					Lake N11					Lake M1	Lake M2	Lake M3	Lake M4	Lake L2
	East-D1	East-D2	East-D3	East-D4	East-D5	N11-D1	N11-D2	N11-D3	N11-D4	N11-D5	M1	M2	M3	M4	L2
	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]	[%]
Nematoda	6	15	6	20	13	33	40	38	11	12	7	16	8	32	6
Oligochaeta	9	33	10	10	10	0	8	2	4	1	11	1	2	0	4
Gastropoda	4	3	0	1	1	1	1	1	2	0	6	0	2	0	8
Pelecypoda	26	22	18	18	7	6	6	5	22	3	7	6	2	4	7
Chironomidae	53	25	66	49	53	57	46	39	59	81	49	66	71	39	50
Other	2	1	1	3	15	2	0	14	2	2	20	10	15	25	25
TOTAL	100	100	100	100	100	100	100	100	100	100	100	100	100	100	100

Note: % = percent.



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Table II-4: Raw Benthic Invertebrate Abundance Data (no./sample) Collected Using a Standard Ekman Grab Sampler, Fall 2011

Major Taxon	Family	Subfamily	Tribe	Genus/Species	East (Reference) Lake D1					East (Reference) Lake D2					East (Reference) Lake D3					East (Reference) Lake D4				
					EAST-D1-A	EAST-D1-B	EAST-D1-C	EAST-D1-D	EAST-D1-E	EAST-D2-A	EAST-D2-B	EAST-D2-C	EAST-D2-D	EAST-D2-E	EAST-D3-A	EAST-D3-B	EAST-D3-C	EAST-D3-D	EAST-D3-E	EAST-D4-A	EAST-D4-B	EAST-D4-C	EAST-D4-D	EAST-D4-E
Microturbellaria	Typhloplanidae	-	-	Mesostoma	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Nematoda	-	-	-	-	0	1	0	1	5	0	10	1	0	0	3	5	3	7	4	12	36	47	0	26
Oligochaeta	Enchytraeidae	-	-	-	1	0	0	0	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0
	Lumbriculidae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	1
	Naididae	Naidinae	-	-	2	2	0	0	3	2	12	3	0	1	3	4	8	7	8	4	1	10	2	6
	Naididae	Tubificinae	-	-	2	0	0	0	1	1	0	0	2	1	1	2	1	5	0	4	11	9	7	3
Gastropoda	Valvatidae	-	-	Valvata sincera	4	0	0	0	1	1	0	1	0	0	0	0	0	1	0	2	0	0	3	2
Bivalvia	Pisidiidae	-	-	(i/d)	5	2	5	0	4	2	5	0	3	0	3	3	4	9	2	3	10	19	12	7
		-	-	Sphaerium	1	0	0	0	0	0	1	0	0	0	0	1	0	0	1	1	1	4	0	0
		-	-	Pisidium	2	0	4	1	7	1	4	0	0	0	7	12	6	19	4	4	11	12	11	14
Hydracarina	-	-	-	-	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	1	3	0	1
Copepoda - Calanoida	-	-	-	-	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Copepoda - Cyclopoida	Cyclopidae	Cyclopinae	-	Acanthocyclops	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	0	10	0	0
	Ergasilidae	-	-	Ergasilus	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Copepoda - Harpacticoida	-	-	-	-	0	0	0	0	0	0	0	0	0	0	0	2	0	0	1	0	2	8	0	1
Ostracoda	-	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Cladocera	Bosminidae	-	-	Bosmina	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Chydoridae	-	-	Eurycercus	2	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Chydoridae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Daphnidae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Macrothricidae	-	-	-	7	16	0	0	0	1	0	0	1	0	1	0	1	0	1	0	2	11	1	1
	Sididae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Trichoptera	Hydroptilidae	-	-	Agraylea	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Leptoceridae	-	-	Oecetis	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Limnephilidae	-	-	Grensia praeterica	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Phryganeidae	-	-	Phryganea	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Diptera	Chironomidae	-	-	(pupa)	2	0	0	0	0	0	1	0	0	0	0	0	1	1	1	0	0	1	0	0
		Tanypodinae	Pentaneurini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Ablabesmyia	0	3	0	0	0	0	0	0	0	0	0	1	1	0	0	0	1	0	1	0
				Thienemannimyia group	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
			Procladiini	Procladius	0	3	1	0	1	0	0	0	0	0	1	4	2	3	2	1	2	6	3	1
		Diamesinae	Protanypini	Protanypus	0	0	0	0	0	0	0	0	0	0	0	3	0	0	0	0	0	0	0	0
		Prodiamesinae	-	Monodiamesa	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
		Orthocladiinae	Orthocladiini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Abyskomyia	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0
				Cricotopus / Orthocladius	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0
				Heterotanytarsus	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Heterotrissocladius	2	4	0	0	1	0	4	0	0	0	4	6	14	21	55	2	7	27	1	1
				Parakiefferiella	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Psectrocladius	1	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	3	0	0
				Zalutschia	0	0	0	0	0	0	0	0	0	0	1	0	3	1	3	0	1	0	1	0
				(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Chironominae	Chironomini	Chironomus	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	1	3	0	1
				Cladopelma	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	1
				Cryptochironomus	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0
				Dicrotendipes	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1



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Table II-4: Raw Benthic Invertebrate Abundance Data (no./sample) Collected Using a Standard Ekman Grab Sampler, Fall 2011 (continued)

Major Taxon	Family	Subfamily	Tribe	Genus/Species	East (Reference) Lake D1					East (Reference) Lake D2					East (Reference) Lake D3					East (Reference) Lake D4				
					EAST-D1-A	EAST-D1-B	EAST-D1-C	EAST-D1-D	EAST-D1-E	EAST-D2-A	EAST-D2-B	EAST-D2-C	EAST-D2-D	EAST-D2-E	EAST-D3-A	EAST-D3-B	EAST-D3-C	EAST-D3-D	EAST-D3-E	EAST-D4-A	EAST-D4-B	EAST-D4-C	EAST-D4-D	EAST-D4-E
Diptera (con't)	Chironomidae (con't)	Chironominae (con't)	Chironomini (con't)	<i>Microtendipes</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	1	1	0	1	1	0	1
				<i>Pagastiella</i>	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	3	2	2	1	1
				<i>Parachironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Polypedilum</i>	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Sergentia</i>	0	0	0	0	0	0	0	0	0	0	1	1	2	0	2	3	2	2	2	5
				<i>Stictochironomus</i>	4	5	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
			Pseudochironomini	<i>Pseudochironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
			Tanytarsini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Cladotanytarsus</i>	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1	0	1	0	0	0
				<i>Corynocera</i>	2	1	0	0	0	0	1	0	0	0	1	1	2	3	0	12	10	40	17	27
				<i>Micropsectra</i>	0	2	0	0	1	0	1	1	2	0	8	7	19	32	33	5	3	47	15	8
				<i>Micropsectra / Tanytarsus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Paratanytarsus</i>	17	5	0	0	1	1	2	1	1	0	2	0	0	2	1	3	1	4	1	3
				<i>Stempellinella</i>	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Tanytarsus</i>	1	0	1	0	4	3	0	0	0	0	0	3	2	4	4	0	0	0	0	0
	Ceratopogonidae	Ceratopogoninae	-	<i>Bezzia</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Dasyheleinae		<i>Dasyhelea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Empididae	-	-	<i>Chelifera / Metachela</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Terrestrial	-	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Total					57	49	12	2	30	12	45	7	9	2	37	55	69	119	126	61	108	270	80	112

Major Taxon	Family	Subfamily	Tribe	Genus/Species	East (Reference) Lake D5					Lake N11-D1					Lake N11-D2					Lake N11-D3				
					EAST-D5-A	EAST-D5-B	EAST-D5-C	EAST-D5-D	EAST-D5-E	N11-D1-A	N11-D1-B	N11-D1-C	N11-D1-D	N11-D1-E	N11-D2-A	N11-D2-B	N11-D2-C	N11-D2-D	N11-D2-E	N11-D3-A	N11-D3-B	N11-D3-C	N11-D3-D	N11-D3-E
Microturbellaria	Typhloplanidae	-	-	<i>Mesostoma</i>	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Nematoda	-	-	-	-	8	25	41	19	41	108	328	131	221	190	20	379	394	440	472	157	160	109	84	418
Oligochaeta	Enchytraeidae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0	0	0	0	0
	Lumbriculidae	-	-	-	0	0	0	0	0	1	1	1	0	0	0	3	4	1	8	0	0	0	0	4
	Naididae	Naidinae	-	-	19	22	27	8	19	0	0	3	1	0	1	64	65	74	56	4	8	4	4	17
Gastropoda	Naididae	Tubificinae	-	-	2	0	2	4	3	0	0	0	0	0	5	4	9	4	24	9	0	1	0	1
	Valvatidae	-	-	<i>Valvata sincera</i>	1	3	1	3	2	6	8	3	3	5	0	12	4	8	6	3	1	10	0	14
Bivalvia	Pisidiidae	-	-	(i/d)	6	12	4	2	3	0	29	8	25	18	0	53	27	43	59	12	20	6	7	25
		-	-	<i>Sphaerium</i>	0	0	0	0	0	4	1	2	0	4	1	3	1	0	0	0	0	2	0	0
		-	-	<i>Pisidium</i>	10	15	11	6	4	10	18	16	10	16	3	13	17	12	11	9	16	7	3	11
Hydracarina	-	-	-	-	0	2	8	0	10	0	8	0	0	4	0	0	0	0	0	0	0	16	0	0
Copepoda - Calanoida	-	-	-	-	0	1	1	1	1	0	8	0	0	0	1	0	0	8	0	4	4	4	2	8
Copepoda - Cyclopoida	Cyclopidae	Cyclopinae	-	<i>Acanthocyclops</i>	2	7	4	1	5	9	0	0	4	0	0	89	32	0	8	16	12	50	4	34
	Ergasilidae	-	-	<i>Ergasilus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0
Copepoda - Harpacticoida	-	-	-	-	1	44	21	19	52	16	8	11	20	4	4	0	0	0	0	108	48	105	18	48
Ostracoda	-	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0



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Table II-4: Raw Benthic Invertebrate Abundance Data (no./sample) Collected Using a Standard Ekman Grab Sampler, Fall 2011 (continued)

Major Taxon	Family	Subfamily	Tribe	Genus/Species	East (Reference) Lake D5					Lake N11-D1					Lake N11-D2					Lake N11-D3						
					EAST-D5-A	EAST-D5-B	EAST-D5-C	EAST-D5-D	EAST-D5-E	N11-D1-A	N11-D1-B	N11-D1-C	N11-D1-D	N11-D1-E	N11-D2-A	N11-D2-B	N11-D2-C	N11-D2-D	N11-D2-E	N11-D3-A	N11-D3-B	N11-D3-C	N11-D3-D	N11-D3-E		
Cladocera	Bosminidae	-	-	<i>Bosmina</i>	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
	Chydoridae	-	-	<i>Eurycercus</i>	0	0	0	0	0	7	0	0	1	0	0	3	0	0	0	0	2	12	1	3		
	Chydoridae	-	-	-	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0			
	Daphnidae	-	-	-	0	0	0	0	0	7	1	2	2	0	0	16	2	3	6	0	7	14	0	2		
	Macrothricidae	-	-	-	9	3	5	0	3	0	0	0	0	0	0	16	0	0	0	0	9	23	0	8		
	Sididae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
Trichoptera	Hydroptilidae	-	-	<i>Agraylea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
	Leptoceridae	-	-	<i>Oecetis</i>	0	0	0	0	0	0	1	0	0	0	0	0	2	1	0	0	0	0	0	0		
	Limnephilidae	-	-	<i>Grensia praeterica</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
	Phryganeidae	-	-	<i>Phryganea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
Diptera	Chironomidae	-	-	(pupa)	0	1	3	0	1	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0		
		Tanypodinae	Pentaneurini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0		
				<i>Ablabesmyia</i>	0	1	0	0	0	9	3	1	0	1	0	5	18	9	18	5	1	15	0	1		
				<i>Thienemannimyia</i> group	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
			Procladiini	<i>Procladius</i>	6	1	4	0	2	22	14	4	16	19	6	41	38	33	41	16	32	26	10	39		
		Diamesinae	Protanypini	<i>Protanypus</i>	1	2	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
		Prodiamesinae	-	<i>Monodiamesa</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
		Orthocladiinae	Orthocladiini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	0	0	
				<i>Abyskomyia</i>	3	7	3	5	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
				<i>Cricotopus</i> / <i>Orthocladius</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0	0	0	0	0	0	
				<i>Heterotanytarsus</i>	0	0	0	0	0	4	8	0	0	8	0	16	0	8	0	0	4	0	0	0	0	
				<i>Heterotrissocladius</i>	54	48	63	10	32	9	0	3	0	1	0	0	0	1	9	0	0	4	0	9	0	
				<i>Parakiefferiella</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Psectrocladius</i>	3	2	1	0	1	0	8	3	11	4	1	35	10	9	20	81	21	7	25	119	0	
				<i>Zalutschia</i>	1	3	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
		Chironominae	Chironomin	(i/d)	0	0	0	0	0	1	2	0	4	1	0	0	0	0	0	0	0	0	0	0	0	
				<i>Chironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	2	9	0	0	0	0	0	0	0	
				<i>Cladopelma</i>	0	0	0	0	2	4	17	4	5	2	2	8	0	9	8	14	15	5	3	8	0	
				<i>Cryptochironomus</i>	0	1	0	0	1	0	6	0	6	6	0	9	2	2	1	0	4	4	1	4	0	
				<i>Dicrotendipes</i>	0	0	0	0	0	0	0	4	1	0	0	0	0	0	1	0	0	1	0	0	0	
				<i>Microtendipes</i>	2	1	1	1	1	7	5	5	7	19	1	34	3	8	8	3	14	20	2	11	0	
				<i>Pagastiella</i>	0	0	0	1	0	5	0	0	0	1	0	0	0	0	9	4	4	15	1	17	0	
				<i>Parachironomus</i>	0	0	0	0	0	0	0	0	0	0	0	8	0	8	1	0	0	3	0	0	0	
				<i>Polypedilum</i>	0	0	0	0	0	0	16	1	8	2	3	8	2	15	3	2	4	1	5	5	0	
				<i>Sergentia</i>	5	2	4	0	1	0	0	0	0	0	0	0	0	0	0	1	6	0	0	8	0	
				<i>Stictochironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
		Pseudochironomini		<i>Pseudochironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	1	0	0		
				(i/d)	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0	0	0	0	8		
		Tanytarsini		<i>Cladotanytarsus</i>	0	1	0	1	0	0	11	2	11	12	3	16	22	10	13	0	1	0	2	18	0	
				<i>Corynocera</i>	6	22	21	3	15	214	395	139	310	241	13	283	491	228	272	35	68	57	23	69	0	
				<i>Micropsectra</i>	41	57	37	9	32	9	8	0	0	5	0	0	16	0	1	0	0	4	0	8	0	
				<i>Micropsectra</i> / <i>Tanytarsus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	8	0	
				<i>Paratanytarsus</i>	0	8	4	3	6	5	1	0	1	0	0	0	0	0	0	1	0	0	0	1	0	
				<i>Stempellinella</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Tanytarsus</i>	1	1	2	3	0	10	8	0	1	5	0	32	32	16	48	0	8	8	2	16	0	
	Ceratopogonidae	Ceratopogoninae	-	<i>Bezzia</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
		Dasyheleinae	-	<i>Dasyhelea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
	Empididae	-	-	<i>Chelifera</i> / <i>Metachela</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
	Terrestrial	-	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		
Total					182	292	269	99	240	475	914	343	668	568	64	1,152	1,193	958	1,121	488	473	538	197	942		



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Table II-4: Raw Benthic Invertebrate Abundance Data (no./sample) Collected Using a Standard Ekman Grab Sampler, Fall 2011 (continued)

Major Taxon	Family	Subfamily	Tribe	Genus/Species	Lake N11-D4					Lake N11-D5					Lake M1					Lake M2				
					N11-D4-A	N11-D4-B	N11-D4-C	N11-D4-D	N11-D4-E	N11-D5-A	N11-D5-B	N11-D5-C	N11-D5-D	N11-D5-E	M1-A	M1-B	M1-C	M1-D	M1-E	M2-A	M2-B	M2-C	M2-D	M2-E
Microturbellaria	Typhloplanidae	-	-	Mesostoma	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Nematoda	-	-	-	-	0	6	1	3	0	105	514	25	58	65	8	44	12	34	16	6	54	10	101	88
Oligochaeta	Enchytraeidae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Lumbriculidae	-	-	-	0	0	0	2	0	2	0	2	1	0	5	8	5	2	3	1	0	1	2	1
	Naididae	Naidinae	-	-	0	0	0	1	0	18	24	0	16	15	16	59	31	8	0	0	0	0	8	0
	Naididae	Tubificinae	-	-	0	0	0	1	0	0	1	0	2	1	0	19	9	0	16	0	0	0	5	0
Gastropoda	Valvatidae	-	-	Valvata sincera	0	1	0	0	1	9	8	1	3	1	14	28	13	21	16	0	1	2	2	1
Bivalvia	Pisidiidae	-	-	(i/d)	0	1	0	5	0	16	47	5	14	5	21	29	11	10	13	6	10	1	16	18
		-	-	Sphaerium	0	0	0	0	0	2	1	2	0	0	0	1	1	4	0	0	0	0	0	0
		-	-	Pisidium	3	4	0	5	3	23	18	4	14	9	15	7	1	2	3	9	7	1	19	12
Hydracarina	-	-	-	-	0	0	0	0	0	12	27	0	2	0	0	18	8	24	8	4	12	0	18	3
Copepoda - Calanoida	-	-	-	-	0	0	0	0	0	16	8	0	0	0	0	0	0	0	0	0	0	0	0	0
Copepoda - Cyclopoida	Cyclopidae	Cyclopinae	-	Acanthocyclops	0	1	0	0	0	28	34	0	0	0	0	16	4	0	8	2	1	0	20	41
	Ergasilidae	-	-	Ergasilus	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4
Copepoda - Harpacticoida	-	-	-	-	0	2	0	0	0	12	48	0	0	20	24	44	20	81	80	17	16	10	72	0
Ostracoda	-	-	-	-	0	0	0	0	0	0	0	0	0	0	8	1	0	8	0	1	0	0	0	1
Cladocera	Bosminidae	-	-	Bosmina	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	4
	Chydoridae	-	-	Eurycercus	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0
	Chydoridae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0
	Daphnidae	-	-	-	0	0	0	0	0	2	0	0	2	4	0	0	0	0	0	0	0	0	0	0
	Macrothricidae	-	-	-	0	0	0	0	0	0	0	0	0	0	8	6	0	0	0	0	2	0	5	19
	Sididae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4
Trichoptera	Hydroptilidae	-	-	Agraylea	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Leptoceridae	-	-	Oecetis	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Limnephilidae	-	-	Grensia praeterica	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Phryganeidae	-	-	Phryganea	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
Diptera	Chironomidae	-	-	(pupa)	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Tanypodinae	Pentaneurini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Ablabesmyia	0	0	0	0	0	5	8	0	2	0	17	16	5	21	1	1	4	0	9	3
				Thienemannimyia group	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
				Procladius	0	3	1	1	3	39	50	10	27	19	26	13	1	9	11	13	21	1	31	39
		Diamesinae	Protanypini	Protanypus	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Prodiamesinae	-	Monodiamesa	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Orthoclaadiinae	Orthoclaadiini	(i/d)	0	0	0	0	0	0	8	0	0	0	0	4	4	0	0	0	0	0	8	4
				Abyskomyia	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Cricotopus / Orthocladus	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0	0	0	0	0	0
				Heterotanytarsus	0	1	0	0	0	1	24	8	0	12	0	0	0	1	0	0	0	0	4	0
				Heterotrissocladius	0	0	0	0	0	0	10	0	0	1	0	0	0	0	0	0	2	0	0	0
				Parakiefferiella	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				Psectrocladius	0	1	0	0	0	28	27	1	17	1	59	60	4	59	16	0	24	0	45	46
				Zalutschia	0	0	0	0	0	0	0	0	0	5	1	0	0	0	0	0	0	0	0	0



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Table II-4: Raw Benthic Invertebrate Abundance Data (no./sample) Collected Using a Standard Ekman Grab Sampler, Fall 2011 (continued)

Major Taxon	Family	Subfamily	Tribe	Genus/Species	Lake N11-D4					Lake N11-D5					Lake M1					Lake M2				
					N11-D4-A	N11-D4-B	N11-D4-C	N11-D4-D	N11-D4-E	N11-D5-A	N11-D5-B	N11-D5-C	N11-D5-D	N11-D5-E	M1-A	M1-B	M1-C	M1-D	M1-E	M2-A	M2-B	M2-C	M2-D	M2-E
Diptera (con't)	Chironomidae (con't)	Chironominae	Chironomini	(i/d)	0	0	0	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0
				<i>Chironomus</i>	0	0	0	0	1	4	5	0	2	1	1	26	16	2	0	0	0	0	0	0
				<i>Cladopelma</i>	0	0	0	0	0	4	12	0	8	1	3	6	2	19	0	1	8	0	8	3
				<i>Cryptochironomus</i>	1	1	1	1	0	2	6	1	1	1	0	2	0	0	0	0	3	1	4	3
				<i>Dicrotendipes</i>	0	0	0	0	0	5	0	0	0	0	3	1	2	2	0	0	3	1	1	3
				<i>Microtendipes</i>	1	0	0	0	0	9	13	0	3	0	0	4	0	0	8	21	86	8	137	150
				<i>Pagastiella</i>	1	2	0	2	0	7	9	0	13	8	42	28	33	16	0	6	25	1	35	4
				<i>Parachironomus</i>	0	0	0	0	0	2	0	0	1	1	0	0	0	1	0	0	0	0	0	0
				<i>Polypedilum</i>	0	2	0	0	0	52	90	8	31	22	11	24	10	16	0	14	6	2	22	17
				<i>Sergenta</i>	0	0	0	1	0	2	0	1	0	1	1	0	0	1	0	0	0	0	0	3
				<i>Stictochironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
			Pseudochironomini	<i>Pseudochironomus</i>	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0
		Tanytarsini		(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	6	0	0	4
				<i>Cladotanytarsus</i>	0	0	0	0	0	19	89	0	26	8	1	0	0	1	0	2	11	0	13	7
				<i>Corynocera</i>	3	11	0	14	1	1,097	1,385	250	1,199	230	0	0	0	0	0	3	13	0	15	25
				<i>Micropsectra</i>	0	0	0	2	0	13	64	0	16	4	0	0	0	0	0	0	0	0	0	0
				<i>Micropsectra / Tanytarsus</i>	0	0	0	0	0	0	8	0	16	4	0	0	0	0	0	0	0	0	0	0
				<i>Paratanytarsus</i>	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Stempellinella</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Tanytarsus</i>	0	0	0	0	0	9	10	0	0	0	50	73	21	32	0	1	3	0	45	59
	Ceratopogonidae	Ceratopogoninae	-	<i>Bezzia</i>	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	1	0	0	1
		Dasyheleinae	-	<i>Dasyhelea</i>	0	0	0	0	0	0	0	0	0	0	0	3	0	0	0	0	0	0	0	0
	Empididae	-	-	<i>Chelifera / Metachela</i>	0	0	0	0	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0
Terrestrial	-	-	-	-	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
Total					9	36	3	38	9	1,545	2,551	318	1,477	443	334	540	221	383	199	109	321	39	655	567

Major Taxon	Family	Subfamily	Tribe	Genus/Species	Lake N11-D4					Lake N11-D5					Lake M1					Lake M2				
					N11-D4-A	N11-D4-B	N11-D4-C	N11-D4-D	N11-D4-E	N11-D5-A	N11-D5-B	N11-D5-C	N11-D5-D	N11-D5-E	M1-A	M1-B	M1-C	M1-D	M1-E	M2-A	M2-B	M2-C	M2-D	M2-E
Microturbellaria	Typhlopidae	-	-	<i>Mesostoma</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Nematoda	-	-	-	-	0	6	1	3	0	105	514	25	58	65	8	44	12	34	16	6	54	10	101	88
Oligochaeta	Enchytraeidae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Lumbriculidae	-	-	-	0	0	0	2	0	2	0	2	1	0	5	8	5	2	3	1	0	1	2	1
	Naididae	Naidinae	-	-	0	0	0	1	0	18	24	0	16	15	16	59	31	8	0	0	0	0	8	0
	Naididae	Tubificinae	-	-	0	0	0	1	0	0	1	0	2	1	0	19	9	0	16	0	0	0	5	0
Gastropoda	Valvatidae	-	-	<i>Valvata sincera</i>	0	1	0	0	1	9	8	1	3	1	14	28	13	21	16	0	1	2	2	1
Bivalvia	Pisidiidae	-	-	(i/d)	0	1	0	5	0	16	47	5	14	5	21	29	11	10	13	6	10	1	16	18
		-	-	<i>Sphaerium</i>	0	0	0	0	0	2	1	2	0	0	0	1	1	4	0	0	0	0	0	0
		-	-	<i>Pisidium</i>	3	4	0	5	3	23	18	4	14	9	15	7	1	2	3	9	7	1	19	12
Hydracarina	-	-	-	-	0	0	0	0	0	12	27	0	2	0	0	18	8	24	8	4	12	0	18	3
Copepoda - Calanoida	-	-	-	-	0	0	0	0	0	16	8	0	0	0	0	0	0	0	0	0	0	0	0	0
Copepoda - Cyclopoida	Cyclopidae	Cyclopinae	-	<i>Acanthocyclops</i>	0	1	0	0	0	28	34	0	0	0	0	16	4	0	8	2	1	0	20	41
	Ergasilidae	-	-	<i>Ergasilus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4
Copepoda - Harpacticoida	-	-	-	-	0	2	0	0	0	12	48	0	0	20	24	44	20	81	80	17	16	10	72	0
Ostracoda	-	-	-	-	0	0	0	0	0	0	0	0	0	0	8	1	0	8	0	1	0	0	0	1



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Table II-4: Raw Benthic Invertebrate Abundance Data (no./sample) Collected Using a Standard Ekman Grab Sampler, Fall 2011 (continued)

Major Taxon	Family	Subfamily	Tribe	Genus/Species	Lake N11-D4					Lake N11-D5					Lake M1					Lake M2				
					N11-D4-A	N11-D4-B	N11-D4-C	N11-D4-D	N11-D4-E	N11-D5-A	N11-D5-B	N11-D5-C	N11-D5-D	N11-D5-E	M1-A	M1-B	M1-C	M1-D	M1-E	M2-A	M2-B	M2-C	M2-D	M2-E
Cladocera	Bosminidae	-	-	<i>Bosmina</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0	0	4
	Chydoridae	-	-	<i>Eurycerus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	2	0
	Chydoridae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0
	Daphnidae	-	-	-	0	0	0	0	0	2	0	0	2	4	0	0	0	0	0	0	0	0	0	0
	Macrothricidae	-	-	-	0	0	0	0	0	0	0	0	0	0	8	6	0	0	0	0	2	0	5	19
Trichoptera	Sididae	-	-	-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	4
	Hydroptilidae	-	-	<i>Agraylea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Leptoceridae	-	-	<i>Oecetis</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
	Limnephilidae	-	-	<i>Grensia praeterica</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
Diptera	Phryganeidae	-	-	<i>Phryganea</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0
	Chironomidae	-	-	(pupa)	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Tanypodinae	Pentaneurini	(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Ablabesmyia</i>	0	0	0	0	0	5	8	0	2	0	17	16	5	21	1	1	4	0	9	3
				<i>Thienemannimyia</i> group	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
			Procladiini	<i>Procladius</i>	0	3	1	1	3	39	50	10	27	19	26	13	1	9	11	13	21	1	31	39
		Diamesinae	Protanypini	<i>Protanypus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Prodiamesinae	-	<i>Monodiamesa</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Orthoclaadiinae	Orthoclaidiini	(i/d)	0	0	0	0	0	0	8	0	0	0	0	4	4	0	0	0	0	0	8	4
				<i>Abyskomyia</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Cricotopus</i> / <i>Orthocladus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	8	0	0	0	0	0	0
				<i>Heterotanytarsus</i>	0	1	0	0	0	1	24	8	0	12	0	0	0	1	0	0	0	0	4	0
				<i>Heterotrissocladius</i>	0	0	0	0	0	0	10	0	0	1	0	0	0	0	0	0	2	0	0	0
				<i>Parakiefferiella</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Psectrocladius</i>	0	1	0	0	0	28	27	1	17	1	59	60	4	59	16	0	24	0	45	46
				<i>Zalutschia</i>	0	0	0	0	0	0	0	0	0	5	1	0	0	0	0	0	0	0	0	0
		Chironominae	Chironomini	(i/d)	0	0	0	0	0	1	1	0	1	0	0	0	0	0	0	0	0	0	0	0
				<i>Chironomus</i>	0	0	0	0	1	4	5	0	2	1	1	26	16	2	0	0	0	0	0	0
				<i>Cladopelma</i>	0	0	0	0	0	4	12	0	8	1	3	6	2	19	0	1	8	0	8	3
				<i>Cryptochironomus</i>	1	1	1	1	0	2	6	1	1	1	0	2	0	0	0	0	3	1	4	3
				<i>Dicrotendipes</i>	0	0	0	0	0	5	0	0	0	0	3	1	2	2	0	0	3	1	1	3
				<i>Microtendipes</i>	1	0	0	0	0	9	13	0	3	0	0	4	0	0	8	21	86	8	137	150
				<i>Pagastiella</i>	1	2	0	2	0	7	9	0	13	8	42	28	33	16	0	6	25	1	35	4
				<i>Parachironomus</i>	0	0	0	0	0	2	0	0	1	1	0	0	0	1	0	0	0	0	0	0
				<i>Polypedilum</i>	0	2	0	0	0	52	90	8	31	22	11	24	10	16	0	14	6	2	22	17
				<i>Sergenta</i>	0	0	0	1	0	2	0	1	0	1	1	0	0	1	0	0	0	0	0	3
				<i>Stictochironomus</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
		Pseudochironomini		<i>Pseudochironomus</i>	0	0	0	0	0	0	0	0	1	0	0	0	0	1	0	0	0	0	0	0
				(i/d)	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	6	0	0	4
		Tanytarsini		<i>Cladotanytarsus</i>	0	0	0	0	0	19	89	0	26	8	1	0	0	1	0	2	11	0	13	7
				<i>Corynocera</i>	3	11	0	14	1	1,097	1,385	250	1,199	230	0	0	0	0	0	3	13	0	15	25
				<i>Micropsectra</i>	0	0	0	2	0	13	64	0	16	4	0	0	0	0	0	0	0	0	0	0
				<i>Micropsectra</i> / <i>Tanytarsus</i>	0	0	0	0	0	0	8	0	16	4	0	0	0	0	0	0	0	0	0	0
				<i>Paratanytarsus</i>	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Stempellinella</i>	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
				<i>Tanytarsus</i>	0	0	0	0	0	9	10	0	0	0	50	73	21	32	0	1	3	0	45	59
	Ceratopogonidae	Ceratopogoninae	-	<i>Bezzia</i>	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	0	1	0	0	1
		Dasyheleinae		<i>Dasyhelea</i>	0	0	0	0	0	0	0	0	0	0	0	3	0	0	0	0	0	0	0	0
	Empididae	-	-	<i>Chelifera</i> / <i>Metachela</i>	0	0	0	0	0	0	0	0	0	4	0	0	0	0	0	0	0	0	0	0
Terrestrial	-	-	-	-	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0
Total					9	36	3	38	9	1,545	2,551	318	1,477	443	334	540	221	383	199	109	321	39	655	567

Notes: (i/d) = immature or damaged specimens not identified below the taxonomic level indicated; - not identified to this taxonomic level.
Samples were collected using a standard Ekman grab with a bottom sampling area of 0.0232 m².
Samples were sieved through a screen with a 250 µm mesh opening size.



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Table II-5: Sorting Efficiency for Benthic Invertebrate Samples for Lakes in the Gahcho Kue Project Area, Fall 2011

Site	Total Organisms in Initial Sort	Total Organisms in QA/QC Re-Sort	Sorting Efficiency [%]
Ref D1-A	57	0	100
Ref D3-C	69	0	100
Ref D5-E	240	4	98
N11-D3-B	473	10	98
N11-D4-D	38	0	100
M1-C	221	5	98
M3-B	226	7	97

Notes: % sorting efficiency = $[1 - (\# \text{ in QC re-sort} / (\# \text{ sorted originally} + \# \text{ QC resort}))] * 100$.

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