

Jay Project

2014 Plankton Supplemental Baseline Report

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April 2015



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Abbreviations

Abbreviation	Definition			
AEMP	Aquatic Effects Monitoring Program			
CCME	Canadian Council of Ministers of the Environment			
DAR	Developers Assessment Report			
Diavik Mine	Diavik Diamond Mine			
DO	dissolved oxygen			
Dominion Diamond	Dominion Diamond Ekati Corporation			
e.g.	for example			
Ekati Mine	Ekati Diamond Mine			
et al.	and more than one additional author			
FF	far-field			
i.e.	that is			
No.	number			
NWT	Northwest Territories			
Project	Jay Project			
QA	quality assurance			
QA/QC	quality assurance and quality control			
QC	quality control			
RPD	relative percent difference			
sp.	species			
spp.	multiple species			
SRSi	soluble reactive silica			
TKN	total Kjeldahl nitrogen			
TN	total nitrogen			
TP	total phosphorus			
TSI	Trophic State Index			
UTM	Universal Transverse Mercator			
Х	times			



Units of Measure

Unit	Definition			
%	percent			
°C	degrees Celsius			
µg/L	micrograms per litre			
μm	micrometre or micron			
μm³	cubic micrometre			
µm³/L	cubic micrometres per litre			
cells/L	cells per litre			
cm	centimetre			
km	kilometre			
m	metre			
mg	milligram			
mg/m ³	milligrams per cubic metre			
mg/L	milligrams per litre			
mg-P/L	milligrams per litre as phosphorus			
mL	millilitre			
mm	millimetre			
mm ³	cubic millimetre			
m/s	metres per second			
org/L	organisms per litre			



1 INTRODUCTION

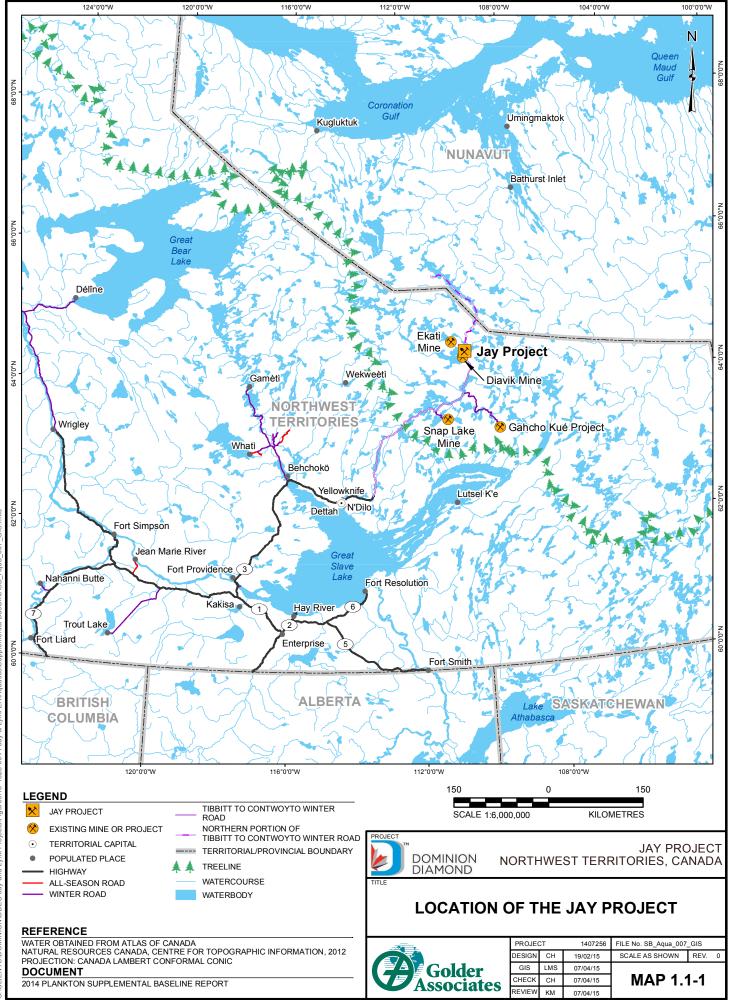
1.1 Background

Dominion Diamond Ekati Corporation (Dominion Diamond) is a Canadian-owned and Northwest Territories (NWT) based mining company that mines, processes, and markets Canadian diamonds from the Ekati Diamond Mine (Ekati Mine). Dominion Diamond also markets Canadian diamonds from its 40% ownership of the Diavik Diamond Mine. The existing Ekati Mine is located approximately 200 kilometres (km) south of the Arctic Circle and 300 km northeast of Yellowknife, NWT (Map 1.1-1).

Dominion Diamond is proposing to develop the Jay kimberlite pipe (Jay pipe) located beneath Lac du Sauvage. The proposed Jay Project (Project) will be an extension of the Ekati Mine, which is a large, stable, and successful mining operation that has been operating for 16 years. Most of the infrastructure required to support the development of the Jay pipe and to process the kimberlite currently exist at the Ekati Mine. The Project is located in the southeastern portion of the Ekati claim block approximately 25 km from the main facilities and approximately 7 km to the northeast of the Misery Pit, in the Lac de Gras watershed.

Plankton baseline field programs were completed in 2013 to support an environmental assessment. The Plankton Baseline Report for the Jay Project (Annex XII) of the Developer's Assessment Report (DAR; Dominion Diamond 2014) summarized the data collected during open-water conditions in 2013, and historical reference (pre-mining) and pre-2013 condition data, to characterize plankton in the lakes within the baseline study area.

The purpose of the 2014 plankton field program was to supplement existing plankton baseline data for lakes in the baseline study area. The 2014 plankton field program focused on Lac du Sauvage, Duchess Lake, and key regions of Lac de Gras that could potentially be influenced by the Jay Project. This report summarizes the supplemental baseline plankton data collected from lakes within the study area during the open-water season (late spring, summer, and fall) in 2014.



G:ICLIENTSIDOMINIONIDDEC Jay and Lynx Projects/Figures/13-1328-0041 Jay & Lynx EA/Aquatics/Supplemental Baseline/SB_Aqua_007_G



1.2 Baseline Study Area

The baseline study area for the 2014 plankton program is located within the headwaters of the Coppermine River drainage, and consists of sub-basins that flow directly into Lac du Sauvage and/or Lac de Gras.

The study area for the 2014 plankton baseline program included the following major basins:

- Lac du Sauvage and the Af sub-basin containing Duchess Lake; and,
- Lac de Gras basin, including Slipper Bay¹ and the upper East Bay² (Far-field 2 [FF2]) areas.

Further information regarding the physical setting of the Project, and the baseline study area, is available in Annex XII of the DAR (Dominion Diamond 2014).

The basin naming convention used for the 2014 baseline program remains unchanged from 2013. This convention was developed for use during the DAR process by all technical disciplines, including other aquatic components (i.e., hydrology, water quality, aquatic health, and fish and fish habitat). As part of the 2014 program, sampling was undertaken in two areas of Lac de Gras also sampled for the existing Ekati and Diavik Aquatic Effects Monitoring Programs (AEMPs). For the 2014 baseline program, samples were collected in Slipper Bay at the Slipper Bay stations established by Ekati Mine (ERM Rescan 2014) and in the upper East Bay at the FF2 stations established by Diavik Diamond Mine (Diavik Mine) (Golder 2014a). The same station identifiers as used in the Ekati and Diavik studies were also used in this baseline program.

¹ Slipper Bay is sampled as part of the Ekati AEMP. For the 2014 supplemental baseline sampling, the same stations were sampled, and station identifiers used by Ekati were applied to the samples collected from these stations. The results included in this report are independent of the Ekati AEMP.

² The upper East Bay is the Far-field 2 (FF2) area sampled as part of the Diavik AEMP. For the 2014 supplemental baseline, the same stations were sampled, and the station identifiers used by Diavik were applied to the samples collected from these stations. The results included in this report are independent of the Diavik AEMP.



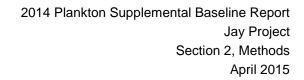
1.3 Objectives

The objectives of the 2014 plankton baseline program were to characterize:

- the trophic status of lakes in the baseline study area during open-water conditions;
- phytoplankton and zooplankton communities in lakes in the baseline study area during open-water conditions; and,
- spatial and seasonal variability in phytoplankton and zooplankton communities, where possible.

Section 2 describes the methods for the collection and analysis of plankton data in the 2014 baseline program. Results of the 2014 plankton sampling program are presented in Section 3.

Detailed descriptions of trophic status classification and phytoplankton and zooplankton community metrics are provided in Annex XII of the DAR.



2 METHODS

2.1.1 Sampling Locations and Timing

The study design for the program was developed to optimize the collection of samples during one-totwo-week field programs, while collecting sufficient data to characterize spatial and temporal variability in plankton communities. The 2014 plankton baseline program was completed during three open-water field programs:

- July 16 to 23 (late spring);
- August 6 to 18 (summer); and,
- September 3 to 16 (fall).

Chlorophyll *a* and depth-integrated nutrient samples were collected as part of the plankton program. Light levels were also measured in the field to generate estimates of light attenuation throughout the water column. Detailed nutrient and chlorophyll *a* results and water column light attenuation profiles are presented in the 2014 Water and Sediment Quality Supplemental Baseline Report (Dominion Diamond 2015).

Plankton samples were collected at stations established in the following lakes (Table 2.1-1; Map 2.1-1):

- Lac du Sauvage (eight stations) and the Af sub-basin containing Duchess Lake (one station);
- Lac de Gras Slipper Bay area (four stations); and,
- Lac de Gras FF2 area (five stations).

Not all stations were sampled during each field program. As the sites are located some distance from the camp at the main Ekati Mine site, access to the sites is via helicopter and boat. Weather conditions (e.g., fog, high winds) during the field programs occasionally resulted in the inability to access some planned locations.

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			UTM Coo	rdinates ^(a)		Phytoplankton and Zooplankton			
Basin	Waterbody Name	Station	Easting (m)	Northing (m)	Total Depth (m)	Late Spring (July)	Summer (August)	Fall (September)	
		Aa-1	552282	7165025	11.4	-	Х	Х	
		Ab-1	547766	7162266	10.1	Х	Х	Х	
		Ac-1	545339	7165138	14.8	Х	Х	X ^(b)	
		Ac-4	543695	7162938	10.2	Х	Х	Х	
Lac du Sauvage	Lac du Sauvage	Ac-7	544247	7165068	13.0	Х	Х	Х	
		Ad-1	539898	7168781	12.2	_(c)	Х	Х	
		Ad-5	540112	7168316	25.3	Х	_(c)	_(c)	
		Ae-1	542494	7170252	15.6	Х	Х	Х	
	Duchess Lake	Af-1	542155	7173731	15.1	-	Х	Х	
	Slipper Bay	S2	507638	7164468	6.5	Х	Х	Х	
		S3	505912	7164439	12.9	Х	X ^(b)	Х	
		S 5	503125	7161482	17.3	X ^(b)	Х	Х	
		S6	501976	7159857	25.5	Х	Х	-	
Lac de Gras		FF2-1	541500	7159522	21.5	Х	Х	Х	
	Far-field 2	FF2-2	541583	7158573	18.8	Х	Х	Х	
		FF2-3	543478	7159267	19.5	Х	Х	Х	
		FF2-4	543752	7158945	19.3	Х	X ^(b)	Х	
		FF2-5	544734	7158898	19.1	X ^(b)	Х	Х	

Table 2.1-1 Plankton Sampling Stations and Sampling Events in the Jay Project Area, 2014

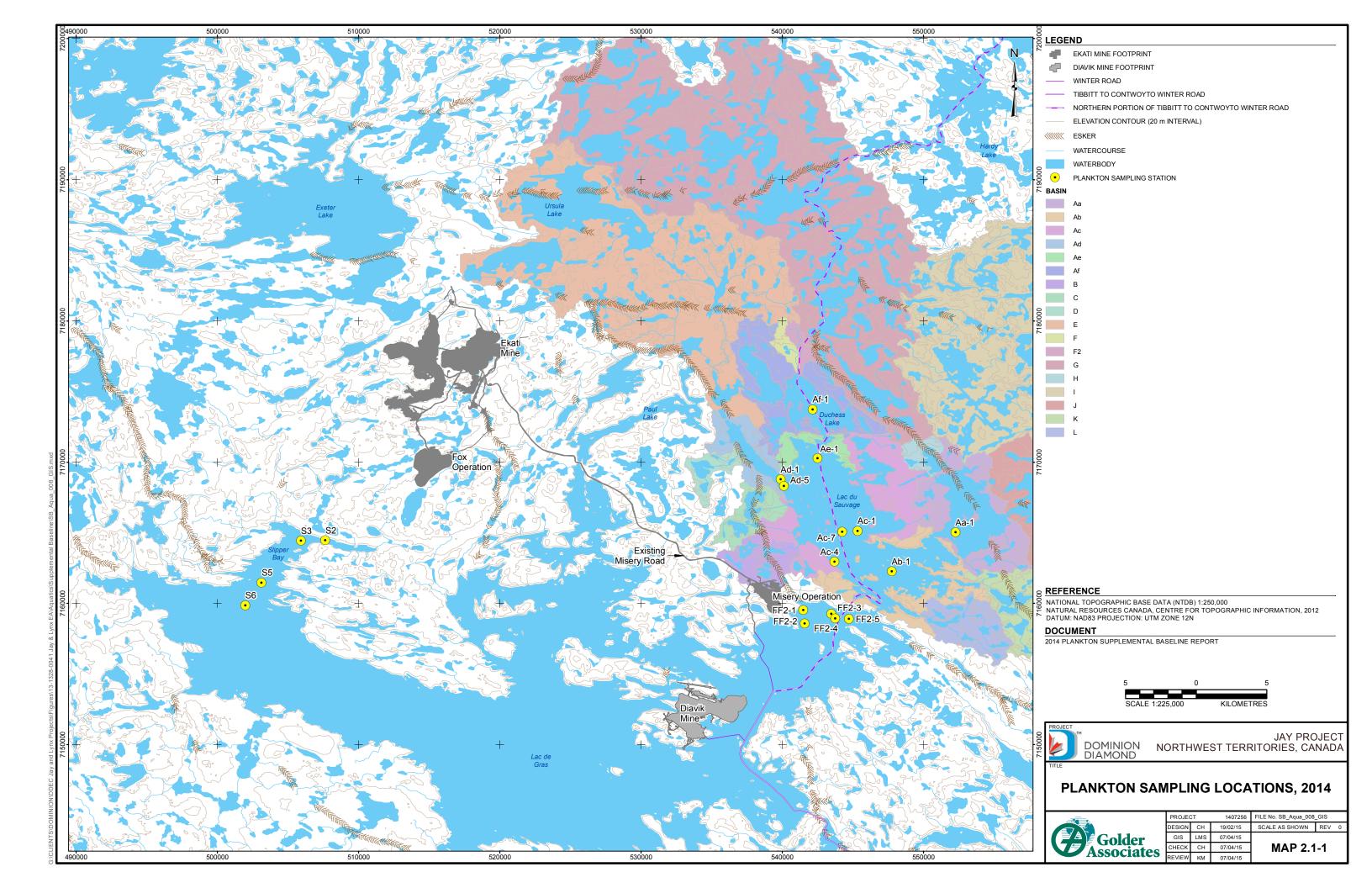
Note: UTM coordinates are in Zone 12V, North American Datum (NAD) 83.

a) UTM coordinates are from the first sampling event at each station.

b) Duplicate phytoplankton and zooplankton samples were collected.

c) During the late spring sampling period, the coordinates sampled for Station Ad-1 were far enough away from the actual Station Ad-1 that this station was assigned a unique station identifier (Station Ad-5). The original Station Ad-1 was sampled during the summer and fall sampling programs.

UTM = Universal Transverse Mercator coordinate system; m = metre; X = samples were collected; - = indicates that samples were not collected due to weather, environment, schedule limitations, or other reasons.





2.1.2 Field Methods

2.1.2.1 Supporting Variables, Nutrients, and Chlorophyll a

Depth-integrated and discrete nutrient samples, and chlorophyll *a* samples were collected as part of the water quality component, and as supporting data for the plankton component. Depth-integrated nutrient samples and chlorophyll *a* samples were collected from the euphotic zone, which is defined as the extent of the water column that is exposed to sufficient sunlight for photosynthesis to occur (typically to a depth where 1 percent [%] of the surface irradiance is measured). During the field program, the euphotic zone was calculated as two times the Secchi depth (Koenings and Edmundson 1991; AENV 2006). Discrete nutrient samples were collected from one of the top, mid, or bottom depths depending on the presence of thermal or oxic stratification (Dominion Diamond 2015).

Stratification was determined by meeting temperature and dissolved oxygen (DO) criteria as follows:

- temperature difference of 1 degree Celsius (°C) or greater over 1 metre (m) depth within the water column; and,
- DO difference of greater than, or equal to, 5 milligrams per litre (mg/L) between the top and the bottom of the water column; or
- DO in the top of the water column greater than the chronic guideline for the protection of aquatic life (6.5 mg/L for cold water species) and DO at the bottom of the water column less than the aquatic life guideline (CCME 2004).

If either of these conditions were met, the water column was considered to be stratified and discrete surface (top) and bottom (bottom) samples were collected. If neither of these conditions were met, the station was considered to be fully mixed or homogenous, and a single mid-column (mid) sample was collected. Lake stations were typically stratified during the under-ice period and fully mixed during the open-water period.

The depth-integrated samples were analyzed for nutrients (i.e., total nitrogen [TN], total dissolved nitrogen, total Kjeldahl nitrogen [TKN], dissolved Kjeldahl nitrogen, total ammonia, nitrate, nitrite, total phosphorus [TP], total dissolved phosphorus, dissolved orthophosphate, and soluble reactive silica [SRSi]) and chlorophyll *a*. Nutrient samples were analyzed by ALS Environmental, Edmonton, Alberta. Chlorophyll *a* samples were analyzed for total chlorophyll *a* by the Biogeochemical Analytical Service Laboratory at University of Alberta, Edmonton, Alberta.

In situ water quality profiles (i.e., pH, temperature, dissolved oxygen, and specific conductivity), light levels, and Secchi depths were measured in conjunction with the water quality baseline program. Detailed field methods for the collection and analysis of the nutrient and chlorophyll *a* samples, and field measurements are presented in the 2014 Water and Sediment Quality Supplemental Baseline Report (Dominion Diamond 2015).

2.1.2.2 Phytoplankton

Phytoplankton samples were collected from the euphotic zone, defined as two times the Secchi depth. At each station, discrete water samples were collected at 2 m intervals within the euphotic zone using a



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Kemmerer sampler. For example, if the water depth was 6 m, then samples were collected at the surface (0 m), 2 m, 4 m, and 6 m. If the water depth was less than 6 m, then samples were collected at surface (0 m), 2 m, and 4 m. If the water depth was less than the Secchi depth, then samples were collected every 2 m from the surface to a depth of 2 m above the lake bottom.

Equal volumes of water from each depth were combined in a large clean bucket and mixed thoroughly to form a composite sample. Samples of water were taken from this composite water sample and used to fill individual 250 millilitre (mL) amber Nalgene bottles for phytoplankton. Phytoplankton samples were preserved with six to eight drops (approximately 2.5 mL) of acid Lugol's solution. Samples were stored in the dark at 4°C before shipping to EcoAnalysts, Inc. (EcoAnalysts) in Moscow, Idaho, United States for taxonomic identification (to the lowest practical taxonomic level), and abundance and biomass estimates.

2.1.2.3 Zooplankton

Maximum water depth was measured before plankton sampling to determine zooplankton sampling depth. A 30 centimetre (cm) diameter, 80 micron (μ m) mesh Turtox plankton tow net was used to collect a single zooplankton sample at each station. A single vertical haul was taken for each zooplankton sample. The plankton net was lowered to a depth of 1 m above the bottom and then pulled vertically through the water column at a rate of approximately 0.5 metres per second (m/s).

Haul depths were recorded for each sample and were used to calculate the volume of water filtered through the net (Table 2.1-2). The plankton net was rinsed by splashing lake water on the outside to wash clinging zooplankton into the bottom of the plankton net. A 250 mL clear Nalgene bottle was placed below the tube of the plankton net. The stop-cock was then opened and the sample was transferred into the sample bottle below.

Before preservation, one half of an Alka-Seltzer tablet was added as a narcotizing agent to each sample bottle to prevent the zooplankton from being contorted by the preservative, thereby allowing for easier identification by the taxonomist. Each sample was preserved by doubling the sample volume with 10% buffered formalin solution. Samples were stored at 4°C and sent to EcoAnalysts for taxonomic identification to the lowest practical taxonomic level, and abundance and biomass estimates.

			Zoop	lankton Haul Depth	(m)
Basin	Waterbody Name	Station	Late Spring	Summer	Fall
		Aa-1	-	9.7	8.7
		Ab-1	8.7	10.7	11.7
	Lac du Sauvage	Ac-1	12.7	10.7	9.7
		Ac-4	8.7	9.7	10.7
Lac du Sauvage		Ac-7	11.7	10.7	10.7
		Ad-1	_(a)	10.7	8.7
		Ad-5	23.7	_(a)	_(a)
		Ae-1	13.7	9.7	8.7
	Duchess Lake	Af-1	-	13.7	9.7

Table 2.1-2 Zooplankton Haul Depths for Stations Sampled in the Jay Project Area, 2014



			Zooplankton Haul Depth (m)			
Basin	Waterbody Name	Station	Late Spring	Summer	Fall	
		S2	4.7	4.7	5.7	
	Clipper Dev	S3	10.7	11.7	9.7	
	Slipper Bay	S5	15.7	16.7	12.7	
		S6	23.7	25.7	-	
Lac de Gras		FF2-1	19.7	19.7	17.7	
		FF2-2	17.7	16.7	17.7	
	Far-field 2	FF2-3	17.7	15.7	18.7	
		FF2-4	17.7	18.7	18.7	
		FF2-5	17.7	19.7	18.7	

Table 2.1-2 Zooplankton Haul Depths for Stations Sampled in the Jay Project Area, 2014

a) During the late spring sampling period, the coordinates sampled for Station Ad-1 were far enough away from the actual Station Ad-1 that this station was assigned a unique station identifier (Station Ad-5). The original Station Ad-1 was sampled during the summer and fall sampling programs.

m = metre; - = indicates that samples were not collected due to weather, environment, schedule limitations, or other reasons.

2.1.3 Laboratory Methods

2.1.3.1 Phytoplankton

Phytoplankton samples were analyzed at the lowest possible taxonomic level (typically species), and abundance and biomass by EcoAnalysts (2009a). A 5 to 25 mL aliquot was extracted for analysis of softbodied algae and diatoms depending on cell and detritus density. Samples were homogenized and aliquots were placed into a Utermohl counting chamber (22 by 22 millimetres [mm]) to allow them to settle overnight. Samples were examined at 630 times (X) magnification using a Leica inverted microscope to evaluate whether the sample was too dense or dilute to achieve a desirable cell count (approximately 15 to 20 counting units per field of view). Samples were diluted or concentrated, as necessary, and the new volume and concentration ratios were noted.

Soft-bodied algae and diatom units were counted and identified at 630X to the lowest practical taxonomic level using the transect method. A minimum of 300 units were counted for each sample. Counting units were individual cells, filaments, or colonies, depending on the organization of the algae and diatoms. Transects totaling 44 mm (1 full horizontal and 1 full vertical transect) were also counted at 200X enumerating only soft-bodied and diatom taxa between 20 to 100 μ m in size. Lastly, a full chamber, high-level scan at 100X was completed to enumerate large, rare soft-bodied and diatom taxa over 100 μ m in size. Taxonomic identifications were based on standard taxonomic references (Dillard 1991a,b, 1993; Wehr and Sheath 2003; Siver et al. 2005; Pfeil 2010; John et al. 2011).

Biovolume (cubic micrometre $[\mu m^3]$) of each soft-bodied and diatom species was estimated from the average dimensions and related to geometric shapes (Hillerbrand et al. 1999). The number of measurements recorded for each taxon in each sample was based on the level of scan and the relative abundance of the taxa. At 630X, at least one biovolume measurement was made for each soft-bodied and diatom taxon, representing less than 5% relative abundance in the sample. At least 10 biovolume



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measurements were made for every taxon representing greater than 5% of the sample. For the high-level (i.e., 200X and 100X) scans, at least one biovolume measurement was made, but no more than five, for every soft-bodied and diatom taxon encountered. If the taxonomist observed strong discontinuities in the size distribution of a taxon, then a minimum of 20 measurements were recorded per taxon per sample. Biovolumes for each taxon were then averaged together for one final value. The biovolumes of colonial taxa were based on the number of individuals within each colony. If a taxon was identified in more than one level of scan, the biovolume measurements were combined to provide an average biovolume for that taxon.

Average biovolume (cubic micrometres per litre $[\mu m^3/L]$) was converted to biomass for all individual phytoplankton taxa by assuming a specific gravity of 1 (i.e., $1x10^9 \mu m^3 = 1 mm^3 = 1 milligram [mg]$). Total sample biomass (reported as milligrams per cubic metre $[mg/m^3]$ wet weight) for each taxon was calculated by multiplying average biomass by the total abundance (reported as cells per litre [cells/L]).

2.1.3.2 Zooplankton

Zooplankton samples were analyzed at the lowest possible taxonomic level (typically species), abundance, and biomass according to methods provided by EcoAnalysts (2009b). Zooplankton samples were rinsed into a 400 mL beaker with 70% ethanol and allowed to settle overnight. To attain a reasonable density for counting, the supernatant was decanted from the samples using a variable flow chemical pump. The samples were decanted to a safe level to avoid disturbing the settled portion of the sample. Once a reasonable dilution was reached, the sample volumes were measured and recorded.

The sample was mixed thoroughly and a subsample was extracted using a 1 mL Hensen-Stempel pipette. Care was taken to capture the subsample while the sample was thoroughly mixed, to avoid bias resulting from the sinking of heavier organisms.

The 1 mL subsample was rinsed with water (with a drop of soap added to reduce surface tension) into a gridded Corning counting chamber. To achieve the target count of 200 to 400 organisms, adjustments were made either by increasing or reducing the volume, or taking aliquots from the first dilution into a second beaker and further diluting the subsample. To facilitate even distribution of organisms in the counting chamber, no more than 3 mL volumes were counted at one time and each dish was counted in its entirety. Coarse (non-rotifers) and fine (rotifers and copepod nauplii) zooplankters were identified separately. A Leica S8A10 Stereoscope (80X maximum) and a Zeiss Axiolab Compound scope (100X maximum) were used at an average magnification of 40X to identify and enumerate the zooplankton.

After the target count was reached, the ratio of volume counted to original volume was used to calculate abundances for the entire sample. The uncounted portion of the sample was scanned to identify any large or rare taxa that were not encountered during the analysis. Large or rare taxa identified during the scan of the uncounted portion were only included in the taxonomic richness and presence/absence data, and were excluded from the abundance and biomass estimates.

Cyclopoid and calanoid copepod specimens (mature and immature) were identified to species, with the exception of nauplii, which were classified as "copepod nauplii". Organisms that could not be identified to species were identified to genus. Taxonomic identifications were based primarily on Alberti et al. (2007), Edmondson (1959), and Stemberger (1979).



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Dry weight biomass (mg/m³) for each zooplankton taxon was based on published length-weight regressions and mean length measurements (Dumont et al. 1975; US EPA 2012). For each sample, length measurements were made for up to 15 individuals of each taxon contributing to the target count. Biomass calculations were based on the average of the 15 (or fewer, if less than 15 individuals were present in the counted portion of the sample) measurements for each taxon. Zooplankton lengths were determined directly on the microscope fitted with an ocular micrometer.

2.1.4 Data Analysis

2.1.4.1 Trophic Status Classification

The trophic status of each major waterbody was evaluated by examining the nutrient concentrations, chlorophyll *a*, and water transparency (Secchi depth). The trophic status was determined using the Vollenweider (1970) trophic classification scheme for lakes (using TP, TN, chlorophyll *a*, and Secchi depth), the Canadian Council of Ministers of the Environment (CCME 2004) trophic classification scheme for Canadian lakes and streams (using TP), and the Trophic State Index (TSI) developed by Carlson (1977). The TSI is a numerical trophic state index for lakes that classifies lakes on a scale of 0 to 100 (Carlson 1977). The index number is calculated from Secchi depth, chlorophyll *a*, and TP using the following equations (Carlson 1977):

$TSI(TP) = 10\left(6 - \frac{\ln\frac{48}{TP}}{\ln 2}\right)$	[Equation 2.1-1]
$TSI(Chl) = 10\left(6 - \frac{2.04 - 0.68\ln Chl}{\ln 2}\right)$	[Equation 2.1-2]
$TSI(SD) = 10\left(6 - \frac{\ln SD}{\ln 2}\right)$	[Equation 2.1-3]

where:

TSI = trophic state index; TP = total phosphorus; In = the natural logarithm; ChI = chlorophyll *a*; and SD = Secchi depth.

Values calculated using these equations are multiplied by 10 to give the scale a range of 0 to 100. The numerical scales for each of the trophic status indices are presented in the 2013 Water and Sediment Quality Baseline Report, Annex XI, of the DAR (Dominion Diamond 2014).

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2.1.4.2 Plankton Data Analysis

Phytoplankton and zooplankton data were analyzed separately, but the same approach was used to analyze both the datasets. Abundance and biomass data were divided into major taxonomic groups. Phytoplankton groups were Cyanobacteria, Chlorophyceae (chlorophytes), Chrysophyceae (chrysophytes), Cryptophyceae (cryptophytes), Dinophyceae (dinoflagellates), Bacillariophyceae (diatoms), and Euglenophyceae (euglenoids). Zooplankton groups were Cladocera (cladocerans), Calanoida (calanoid copepods), Cyclopoida (cyclopoid copepods), Rotifera (rotifers), and copepod nauplii. Cyclopoid and calanoid copepods were considered separately because of taxonomic and ecological differences. Copepod nauplii were not identified as either cyclopoid or calanoid copepods, but occurred in high abundances in certain samples; therefore, they were treated as a unique taxonomic group for plotting purposes.

Total abundance and total biomass for phytoplankton and zooplankton were plotted as bar graphs by major taxonomic groups. For stations where duplicate quality control (QC) samples were collected, total abundance and total biomass were calculated as the average of the two duplicate samples. The relative proportion accounted for by each major taxonomic group, based on both abundance and biomass, was calculated separately for each station to evaluate variability in community structure among stations.

Total taxonomic richness at the genus level was summarized for each station and plotted as a bar graph for both phytoplankton and zooplankton. Taxonomic richness provides an indication of the diversity at a station; higher richness values typically indicate more healthy and balanced communities. For stations where duplicate QC samples were collected, total taxonomic richness was calculated as the average richness of the two duplicate samples.

2.1.5 Quality Assurance and Quality Control

Quality assurance and quality control (QA/QC) practices determine data integrity and are relevant to all aspects of this baseline sampling program. These practices are applied throughout the activities undertaken within the program, from sample collection to data analysis, and reporting. Quality assurance (QA) encompasses management and technical practices designed to make sure that the data generated are of consistent high quality. Quality control is an aspect of QA that includes the procedures used to measure and evaluate data quality, and the corrective actions to apply when data quality objectives are not met.

The QA/QC procedures were applied during all aspects of the plankton component to verify that the data collected were of acceptable quality. The QA/QC practices applied during this study are described in this section. An evaluation of the QC data and a description of the implications of QC results to the interpretation of study results are provided in Section 3.3.

2.1.5.1 Field Quality Control Procedures

During each sampling program (i.e., late spring, summer, and fall), two pairs of duplicate phytoplankton and zooplankton samples were collected and submitted to the taxonomist for QC purposes. Duplicate samples were used to check within-site variation, and the precision of field sampling methods and laboratory analysis.



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Detailed QC methods and results for the supporting variables (i.e., depth-integrated nutrients and chlorophyll *a*) are presented in Appendix A of the 2014 Water and Sediment Quality Supplemental Baseline Report (Dominion Diamond 2015).

2.1.5.2 Laboratory Quality Control

EcoAnalysts performed an internal QC of the phytoplankton and zooplankton data by having a separate taxonomist re-analyze 10% of the samples to verify taxonomic accuracy and reproducibility of the processing and analytical methods. The percent similarity index was calculated from the two independent plankton counts. The internal QC standards set by EcoAnalysts required that the dominant plankton taxa were aligned, percent similarity was greater than or equal to 50%, and the common plankton taxa accounting for more than 10% relative abundance were identified similarly by both taxonomists. If any one of these criteria were not met, the original sample and its corresponding QC sample were reanalyzed. Discrepancies between taxonomists were resolved by re-examining digital images and/or preserved specimens, and the final organism counts and identifications were adjusted according to the recommendations of both taxonomists.

2.1.5.3 Quality Control Data Evaluation

To examine the variability introduced by field sampling procedures, taxonomic accuracy, and reproducibility of the processing and analysis methods, duplicate phytoplankton and zooplankton samples were analyzed by two QC approaches: Bray-Curtis index, and relative percent difference (RPD).

The Bray-Curtis dissimilarity index is a measure of the ecological distance between two communities. The Bray-Curtis index was calculated in SYSTAT (2009), according to the formula below, to evaluate the overall dissimilarity between the original and duplicate plankton samples:

$$b = \frac{\sum_{k=1}^{n} |x_{ik} - x_{jk}|}{\sum_{k=1}^{n} (x_{ik} + x_{jk})}$$

[Equation 2.1-4]

where,

 x_{ik} and x_{jk} = the abundance from the original and duplicate samples, respectively.

Since the Bray-Curtis index only allows comparisons of entire samples, the RPD was also calculated to compare abundances of each major group between duplicate samples. The RPD was calculated using the following formula:

$$RPD = \left| \frac{(sample-duplicate)}{\frac{(sample+duplicate)}{2}} \right| \times 100$$
 [Equation 2.1-5]



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where,

//= the absolute value;

sample = the abundance in the original sample; and,

duplicate = the abundance in the duplicate sample.

The QC assessment criteria for duplicate samples required the following:

- dominant taxa were aligned;
- the RPD values comparing abundances of major taxa met the established criterion (i.e., less than 50%); and,
- the Bray-Curtis dissimilarity index comparing the original and duplicate samples was less than 0.5.

If any one of these criteria was not met, the sample was flagged. Flagged data were not automatically rejected; rather, they were evaluated on a case-by-case basis, as a certain level of within-station variability is expected in plankton data.



3 **RESULTS**

Raw phytoplankton data are presented in Appendix A, Tables A-3 to A-6 and summarized in Appendix A, Table A-7. Raw zooplankton data are presented in Appendix A, Tables A-8 to A-11 and summarized in Appendix A, Table A-12.

Supporting information, including depth-integrated nutrient concentrations, chlorophyll *a* results, and light attenuation profiles, are presented in the 2014 Water and Sediment Quality Supplemental Baseline Report (Dominion Diamond 2015).

3.1 Lac du Sauvage Basin

3.1.1 Lac du Sauvage and Duchess Lake

3.1.1.1 Trophic Status Classification

Trophic status was evaluated by examining the concentrations of TP, chlorophyll *a*, and water transparency (Secchi depth) (Vollenweider 1970). The discrete water sampling program for Lac du Sauvage yielded mean annual concentrations of 0.0078 milligrams per litre as phosphorus (mg-P/L) for TP and 5.6 m for Secchi depth. The corresponding TSI values were 33.8 using TP and 35.1 using Secchi depth, for a rounded average of 34.5. The depth-integrated sampling program yielded mean annual concentrations of 0.0083 mg-P/L for TP, 5.6 m for Secchi depth, and 2.35 micrograms per litre (µg/L) for chlorophyll *a* (Dominion Diamond 2015). The corresponding TSI values were 34.7 using TP, 38.9 using chlorophyll *a*, and 35.1 using Secchi depth, for a rounded average of 36.3. Based on these TSI values, and the classification systems of Vollenweider (1970), and Carlson (1977), Lac du Sauvage is classified as an oligotrophic lake. Lac du Sauvage can also be classified as oligotrophic (i.e., between 0.004 and 0.01 mg-P/L), based on CCME (2004) TP trigger ranges for Canadian lakes.

The discrete water sampling program for Duchess Lake yielded mean annual concentrations of 0.0125 mg-P/L for TP and 3.6 m for Secchi depth. The corresponding TSI values were 40.6 using TP and 41.5 using Secchi depth, for a rounded average of 41.1. The depth-integrated sampling program yielded mean annual concentrations of 0.015 mg-P/L for TP, 6.79 μ g/L for chlorophyll *a*, and 3.6 m for Secchi depth (Dominion Diamond 2015). The corresponding TSI values were 43.2 using TP, 49.4 using chlorophyll *a*, and 41.5 using Secchi depth, for a rounded average of 44.7. Based on these TSI values, Duchess Lake is classified as mesotrophic using Vollenweider (1970) and Carlson (1977). Duchess Lake can also be classified as mesotrophic (i.e., between 0.01 and 0.02 mg-P/L), based on CCME (2004) TP trigger ranges for Canadian lakes.

3.1.1.2 Phytoplankton

Abundance and Biomass

Seasonal and spatial variation in total phytoplankton abundance was observed in Lac du Sauvage and Duchess Lake during the open-water season (Figure 3.1-1). Total phytoplankton abundance peaked in the summer at the majority of stations in Lac du Sauvage. The highest phytoplankton abundance was observed at stations Aa-1 (1,565,282 cells/L) and Ae-1 (1,643,909 cells/L) in summer. The extent of spatial variation observed in phytoplankton abundance in Lac du Sauvage was greater in the summer (676,103 to 1,643,908 cells/L) compared to late spring (457,734 to 955,784 cells/L) and fall (502,497 to



899,589 cells/L). Phytoplankton abundance varied little among seasons at stations Ac-1, Ac-4, Ac-7, and Ad-1. Phytoplankton abundance in Duchess Lake in fall was similar to Lac du Sauvage stations; however, total abundance in Duchess Lake in summer (3,287,508 cells/L) was notably higher than in Lac du Sauvage (676,103 to 1,643,908 cells/L). The high phytoplankton abundance observed at Station Af-1 in summer was driven by the chrysophyte, *Ochromonas* spp., and to a lesser extent the chrysophyte, *Uroglenopsis americana* and the cryptophyte, *Komma caudata*.

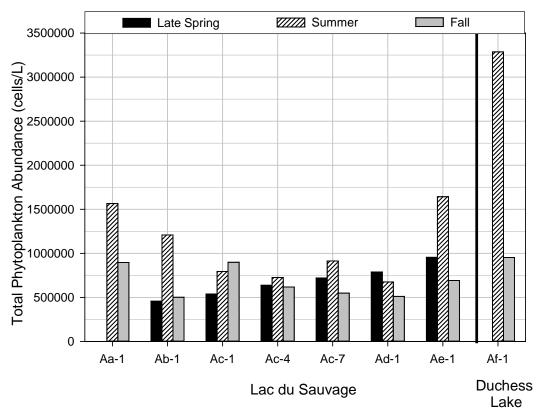


Figure 3.1-1 Total Phytoplankton Abundance in Lac du Sauvage and Duchess Lake in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; late spring samples were not collected at stations Aa-1 and Af-1. cells/L = cells per litre,

During the open-water season, phytoplankton biomass in Lac du Sauvage ranged from 197 to 747 mg/m³, with the exception of stations Aa-1 and Ac-1; however, no clear seasonal trend in total phytoplankton biomass was observed (Figure 3.1-2). Unusually high phytoplankton biomass was observed in fall at Station Aa-1 (1,515 mg/m³), and to a lesser extent Ac-1 (1,049 mg/m³). Total phytoplankton biomass was consistently lower in late spring compared to summer and fall, with values at the majority of stations below 747 mg/m³. Four out of seven stations (Aa-1, Ab-1, Ac-1, and Ae-1) exhibited increases in total phytoplankton biomass throughout the open-water season. The relatively high biomass observed at Station Aa-1 in fall was largely driven by the large dominant dinoflagellate,



Peridinium sp., the chrysophytes, *Ochromonas* spp. and *Mallomonas* sp., and the diatom, *Asterionella formosa*. The dominant taxa driving the high phytoplankton biomass at Station Ac-1 in fall was the chrysophyte, *Mallomonas* sp. (298 mg/m³) and the diatom, *Tabellaria fenestrate* (214 mg/m³). Unusually high phytoplankton biomass was also observed at Station Af-1 in Duchess Lake in summer (1,327 mg/m³); high biomass at Duchess Lake in summer was largely driven by the chrysophyte, *Dinobryon divergens*.

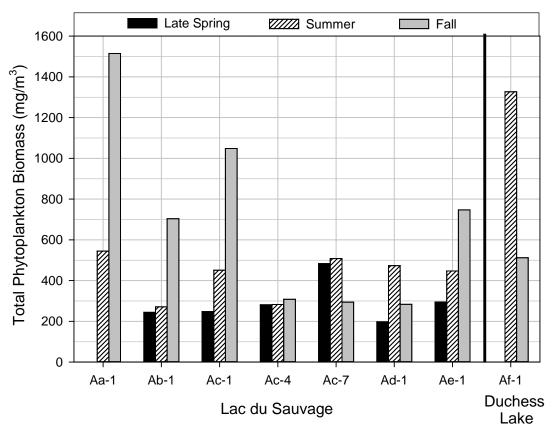


Figure 3.1-2 Total Phytoplankton Biomass in Lac du Sauvage and Duchess Lake in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; late spring samples were not collected at stations Aa-1 and Af-1. $ma/m^3 = milliarama par subjacements$

 $mg/m^3 = milligrams$ per cubic metre.

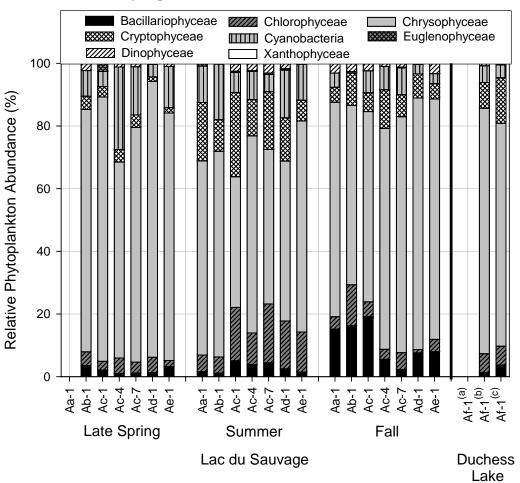
Community Composition

Chrysophytes consistently dominated the community composition by abundance in Lac du Sauvage and Duchess Lake throughout the open-water season, making up 63% to 88% of the phytoplankton assemblage in late spring, 42% to 78% in summer, and 57% to 80% in fall (Figure 3.1-3). Other major taxonomic groups such as cryptophytes (less than 27%), chlorophytes (less than 19%), cyanobacteria (less than 26%), and diatoms (less than 19%) were present in Lac du Sauvage at varying relative



abundances throughout the open-water season. Together, euglenoids, dinoflagellates, and xanthophytes made up less than 8% of the phytoplankton assemblage by abundance in Lac du Sauvage and Duchess Lake throughout the open-water season.

Figure 3.1-3 Relative Phytoplankton Abundance in Lac du Sauvage and Duchess Lake in Late Spring, Summer, and Fall, 2014



Note: Late spring samples were not collected at stations Aa-1 and Af-1; bars represent a single composite sample collected within the euphotic zone.

- a) late spring.
- b) summer.
- c) fall.

% = percent.

In terms of biomass, the phytoplankton community composition in Lac du Sauvage varied seasonally and spatially throughout the open-water season (Figure 3.1-4). In late spring, stations Ab-1 and Ac-7 were co-dominated by dinoflagellates (44% and 44%, respectively) and chrysophytes (27% and 30%, respectively). Stations Ac-4 and Ae-1 were co-dominated by cyanobacteria (32% and 29%, respectively)

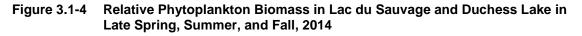


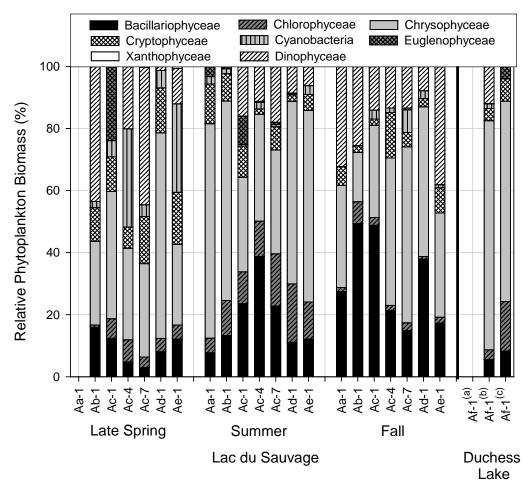
and chrysophytes (30% and 26%, respectively). Two major taxonomic groups, chrysophytes (41%) and euglenoids (24%), made up the majority of the phytoplankton biomass at Station Ac-1 in the late spring. Station Ac-1 was the only station in Lac du Sauvage with a notable (greater than 20%) biomass of euglenoids in late spring. Chrysophytes (66%) dominated the phytoplankton assemblage by biomass at Station Ad-1 in late spring; the composition closely resembled the composition of station Aa-1 in the summer. Cryptophytes (7% to 17%) and diatoms (37% to 16%) made up a relatively small proportion of the biomass in Lac du Sauvage in late spring.

In the summer, the phytoplankton community in Lac du Sauvage was co-dominated by chrysophytes (31% to 69%) and diatoms (8% to 39%); however, the percentages of these groups varied among stations. The remainder of the phytoplankton assemblage was made up of mainly chlorophytes (less than 19%), dinoflagellates (less than 18%), and cryptophytes (less than 13%). In fall, the dominant phytoplankton taxonomic groups by biomass were consistent among stations in Lac du Sauvage; however, the percentages of these groups varied among stations. A mixture of diatoms (15% to 49%), chrysophytes (16% to 57%), and dinoflagellates (8% to 38%) dominated the phytoplankton biomass in Lac du Sauvage in fall.

Chrysophytes dominated the phytoplankton assemblage by biomass in Duchess Lake in the summer and fall (no late spring sample was collected at Station Af-1); the composition closely resembled that at certain stations in Lac du Sauvage in the summer.







Note: Late spring samples were not collected at stations Aa-1 and Af-1; bars represent a single composite sample collected within the euphotic zone.

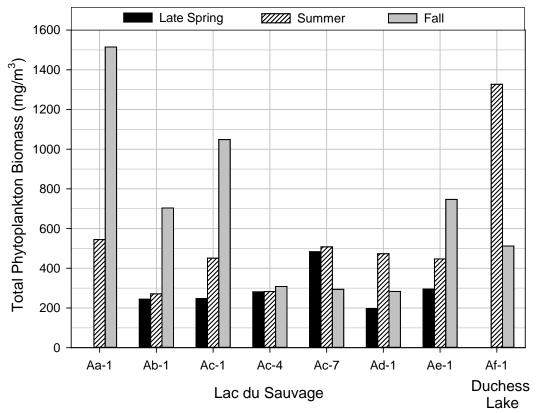
- a) late spring.b) summer.
- c) fall.
- % = percent.

In total, 118 phytoplankton taxa were identified in Lac du Sauvage in 2014: 52 chlorophytes, 18 diatoms, 16 chrysophytes, 15 cyanobacteria, 7 dinoflagellates, 7 cryptophytes, 2 euglenoids, and 1 xanthophyte (Appendix A, Table A-3). In total, 42 phytoplankton taxa were identified in Duchess Lake in 2014: 19 chlorophytes, 6 chrysophytes, 5 cyanobacteria, 4 diatoms, 4 cryptophytes, 3 dinoflagellates, and 1 euglenoid (Appendix A, Table A-4). The lower total phytoplankton richness observed in Duchess Lake reflects the lower sampling effort (i.e., a single sampling station) in this lake compared to Lac du Sauvage. Seasonal variation in phytoplankton taxonomic richness was observed in Lac du Sauvage, with richness values ranging from 18 taxa (stations Aa-1 and Ad-1) in fall to 48 taxa (Station Ac-1) in summer.



A similar level of richness was observed at Station Af-1 in Duchess Lake, with 34 taxa in summer and 16 taxa in fall. The highest phytoplankton richness was observed in the summer (30 to 48 taxa) at all stations in Lac du Sauvage and Duchess Lake (Figure 3.1-5).





Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; late spring samples were not collected at stations Aa-1 and Af-1. No. = number.

3.1.1.3 Zooplankton

Abundance and Biomass

Seasonal and spatial variation were observed in total zooplankton abundance in Lac du Sauvage and Duchess Lake; however, no consistent trends were observed (Figure 3.1-6). Overall, zooplankton abundance in Lac du Sauvage and Duchess Lake ranged from 32 to 66 organisms per litre (org/L) in late spring, 47 to 95 org/L in summer, and 46 to 80 org/L in fall. Seasonal peaks in total zooplankton abundance were observed in the summer at five stations (Aa-1, Ac-1, Ac-4, Ad-1, and Ae-1) in Lac du Sauvage, while abundance was lowest in the summer at Station Ab-1. Total zooplankton abundance at Station Ac-7 decreased throughout the open-water season, while abundance at Station Af-1 in Duchess Lake increased from summer to fall.



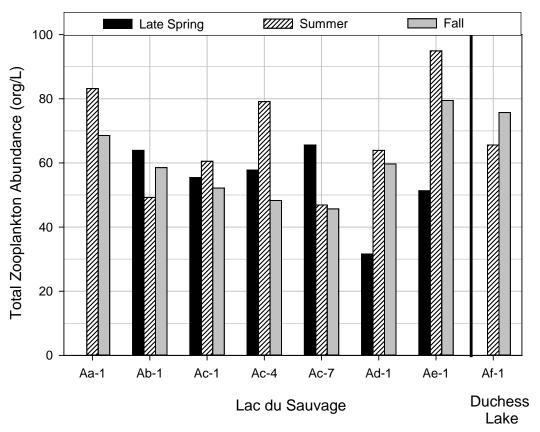
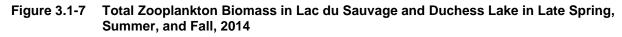


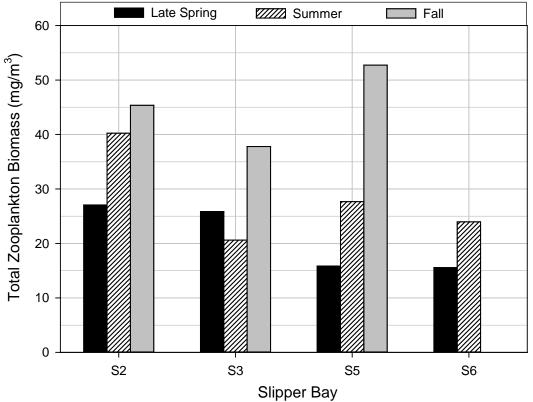
Figure 3.1-6 Total Zooplankton Abundance in Lac du Sauvage and Duchess Lake in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single a vertical haul taken throughout the water column; a mean value for field duplicates are shown; late spring samples were not collected at stations Aa-1 and Af-1. org/L = organisms per litre.

Seasonal and spatial variation in total zooplankton biomass were observed in Lac du Sauvage and Duchess Lake (Figure 3.1-7). Total zooplankton biomass peaked in summer at all stations in Lac du Sauvage and Duchess Lake, ranging from 41 to 152 mg/m³, compared to late spring (13 to 50 mg/m³) and fall (29 to 55 mg/m³). Spatial variability among stations in total zooplankton biomass was greatest in the summer, with lower variability observed in the late spring or fall. Two stations, Ac-4 and Ae-1, had notably higher zooplankton biomass in the summer (97 and 152 mg/m³, respectively) compared to other stations and sampling periods. The dominant taxa driving the high zooplankton biomass in summer at stations Ac-4 and Ae-1 were the cladocerans, *Holopedium gibberum, Daphnia thomasii,* and *Daphnia longiremis.* Total zooplankton biomass in Duchess Lake in the summer and fall (41 and 33 mg/m³, respectively) was similar to Lac du Sauvage, particularly stations Ac-7 and Ad-1.







Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; late spring samples were not collected at stations Aa-1 and Af-1. mg/m³ = milligrams per cubic metre.

Community Composition

Zooplankton community composition by abundance in Lac du Sauvage was similar among stations, but varied with sampling period (Figure 3.1-8). In late spring and summer, zooplankton abundance in Lac du Sauvage was dominated by rotifers (67% to 85% and 49% to 76%, respectively). The relative abundance of rotifers in Lac du Sauvage was lower in fall (46% to 70%), as the proportion of copepod nauplii (11% to 32%) and cyclopoid copepods (9% to 16%) increased. Calanoid copepods made up a very small fraction of the total zooplankton abundance in Lac du Sauvage (less than 2%) and Duchess Lake (less than 1%) throughout the open-water season. Cladocera made up 11% to 31% of the zooplankton abundance in summer in Lac du Sauvage, compared to the late spring (3% to 13%) and fall (3% to 8%).

In summer and fall, the zooplankton assemblage by abundance in Duchess Lake was co-dominated by Cladocera (25% and 48%, respectively) and rotifers (39% and 70%, respectively); however, the proportion of rotifers was higher in fall. Copepod nauplii made up a small proportion of the total abundance in Duchess Lake in summer (12%) and fall (4%).



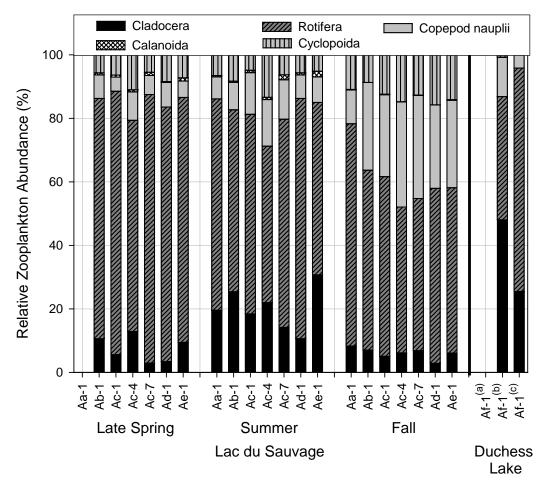


Figure 3.1-8 Relative Zooplankton Abundance in Lac du Sauvage and Duchess Lake in Late Spring, Summer, and Fall, 2014

Note: Late spring samples were not collected at stations Aa-1 and Af-1; bars represent a single composite sample collected within the euphotic zone.

a) late spring.

b) summer.

c) fall.

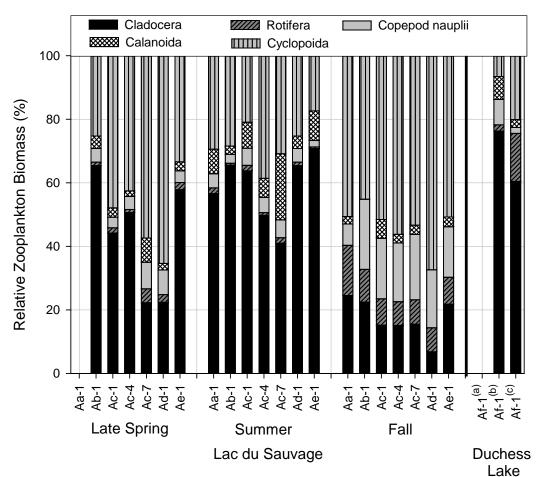
% = percent.

Zooplankton community composition by biomass in Lac du Sauvage varied among stations and throughout the open-water season (Figure 3.1-9). In general, stations in Lac du Sauvage were codominated by Cladocera and cyclopoid copepods in late spring and summer. In late spring, the zooplankton community by biomass in Lac du Sauvage was made up of 22% to 65% Cladocera and 25% to 65% cyclopoid copepods. In summer, Cladocera (41% to 71%) made up more of the zooplankton assemblage by biomass in Lac du Sauvage than cyclopoid copepods (17% to 39%). The relative proportions of these two major taxonomic groups varied among stations in late spring and summer. Stations Ac-7 and Ad-1 had higher proportions of cyclopoid copepods in late spring than the other stations in Lac du Sauvage.



In terms of biomass, Lac du Sauvage stations were dominated by cyclopoid copepods (45% to 67%) in fall. Cladocera (7% to 25%), copepod nauplii (7% to 22%), and rotifers (7% to 16%) made up the majority of the remaining zooplankton biomass in Lac du Sauvage in fall. Throughout the open-water season, the total zooplankton biomass in Lac du Sauvage and Duchess Lake consisted of up to 21% calanoid copepods. The zooplankton assemblage by biomass in Duchess Lake was dominated by Cladocera in summer and fall (76% and 60%, respectively). Despite making up a large proportion of the zooplankton abundance in Lac du Sauvage and Duchess Lake (Figure 3.1-8), rotifers made up a small proportion of total zooplankton biomass due to their small body size (Figure 3.1-9). Rotifers made up a larger fraction of the zooplankton biomass in Lac du Sauvage and Duchess Lake in fall, ranging from 7% to 15%, compared to late spring (less than 4%) and summer (less than 2%).





Note: Late spring samples were not collected at stations Aa-1 and Af-1; bars represent a single composite sample collected within the euphotic zone.

a) late spring.

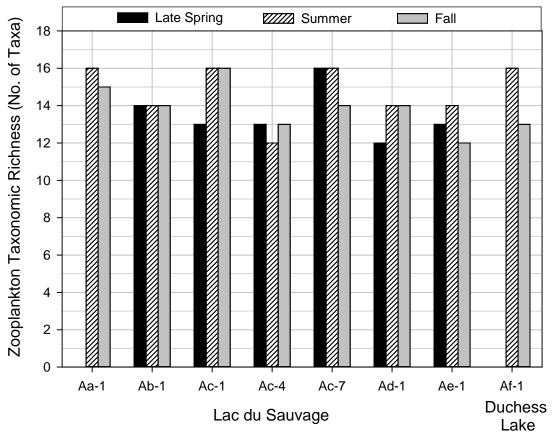
- b) summer.
- c) fall.

% = percent.



In total, 31 zooplankton taxa were identified in Lac du Sauvage in 2014: 18 rotifers, 6 cladocerans, 5 calanoid copepods, and 2 cyclopoid copepods (Appendix A, Table A-8). In total, 19 zooplankton taxa were identified in Duchess Lake in 2014: 13 rotifers, 3 cladocerans, 2 calanoid copepods, and 1 cyclopoid copepod (Appendix A, Table A-9). The lower total zooplankton richness observed in Duchess Lake reflects the lower sampling effort (i.e., a single sampling station) in this lake compared to Lac du Sauvage. Zooplankton taxa (Figure 3.1-10). Zooplankton richness in Duchess Lake in the summer and fall was similar to Lac du Sauvage (16 and 13 taxa, respectively). Some stations in Lac du Sauvage (Aa-1 [16 taxa], Ac-7 [16 taxa], and Ae-1 [14 taxa]) and Duchess Lake station Af-1 (16 taxa) showed a small peak in zooplankton richness in summer, while a slight drop in richness was observed in summer at other stations (Ab-1 and Ac-4).

Figure 3.1-10 Total Zooplankton Taxonomic Richness in Lac du Sauvage and Duchess Lake in Late Spring, Summer, and Fall, 2014



Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; late spring samples were not collected at stations Aa-1 and Af-1.

No. = number.



3.1.2 Comparison of 2013 and 2014 Plankton Data

Results from the 2014 supplemental baseline plankton sampling program were compared to baseline data collected in 2013 (from Lac du Sauvage and Duchess Lake) (Annex XII of the DAR) to describe the range in temporal variation in plankton communities present in these lakes.

3.1.2.1 Trophic Status Classification

The 2014 baseline study evaluated trophic status as the TSI, based on discrete and depth-integrated TP, Secchi depth, and chlorophyll *a* concentrations from the open-water season. The trophic status of the lakes sampled in 2014 ranged from oligotrophic (Lac du Sauvage) to mesotrophic (Duchess Lake). Results from the 2014 trophic status classifications are consistent with the 2013 classifications for Lac du Sauvage and Duchess Lake.

3.1.2.2 Phytoplankton

Based on comparing the 2013 and 2014 phytoplankton data for Lac du Sauvage and Duchess Lake, the following conclusions were made:

- Total phytoplankton abundance and biomass in Lac du Sauvage in 2014 were within the range observed in 2013. However, seasonal and spatial variability in the timing of peak total phytoplankton abundance and biomass in Lac du Sauvage was not consistent among stations or sampling years.
- In 2013, Duchess Lake had higher phytoplankton biomass than Lac du Sauvage, but abundance was comparable between lakes. In 2014, phytoplankton abundance was higher in Duchess Lake than in Lac du Sauvage, while biomass was comparable between the two lakes.
- The peak in total phytoplankton abundance observed in Duchess Lake in the summer of 2014 was two times higher than the maximum abundance observed in Duchess Lake in 2013. However, the phytoplankton abundance in Duchess Lake in fall of 2014 was within the 2013 range.
- The peak in total phytoplankton biomass observed in Duchess Lake in the summer of 2014 was less than half the maximum biomass observed in Duchess Lake in 2013. However, the phytoplankton biomass in Duchess Lake in fall of 2014 was within the 2013 range.
- Chrysophytes consistently dominated the community composition by abundance in Lac du Sauvage throughout the open-water season in 2013 and 2014, while the phytoplankton community by abundance in Duchess Lake differed between years.
- Seasonal and spatial differences in community composition by biomass in Lac du Sauvage and Duchess Lake were observed in 2013 and 2014, and no consistent dominant major taxonomic group was identified.

Overall, the variation observed in community metrics in the baseline phytoplankton dataset underscores the importance of having multiple years of data to characterize the phytoplankton communities present in Lac du Sauvage and Duchess Lake.



3.1.2.3 Zooplankton

Based on comparing the 2013 and 2014 zooplankton data for Lac du Sauvage and Duchess Lake, the following conclusions were made:

- Total zooplankton abundance in Lac du Sauvage in 2014 was within the range observed in Lac du Sauvage in 2013.
- Total zooplankton biomass in Lac du Sauvage in 2014 was notably higher than the range in biomass observed in Lac du Sauvage in 2013. The annual difference in zooplankton biomass in Lac du Sauvage was mainly attributed to the large peaks in biomass recorded in the summer of 2014.
- Total zooplankton abundance and biomass in Duchess Lake in 2014 were within the range observed in Duchess Lake in 2013.
- The timing of seasonal peaks in total zooplankton abundance and biomass in Lac du Sauvage and Duchess Lake were not consistent between 2013 and 2014.
- In 2014, zooplankton community composition by abundance in Lac du Sauvage was similar among stations, but varied with sampling period; the same was true for 2013.
- The zooplankton community by abundance in Lac du Sauvage in the late spring and summer 2014 was similar to the open-water season in 2013 (dominated by rotifers), but differed in fall (codominated by rotifers and copepod nauplii).
- The zooplankton assemblage by abundance in Duchess Lake was consistent between 2013 and 2014 (co-dominated by Cladocera and rotifers).
- Zooplankton community composition by biomass varied seasonally and spatially in Lac du Sauvage and Duchess Lake in 2013 and 2014.

Overall, the variation observed in community metrics in the baseline zooplankton dataset underscores the importance of having multiple years of data to characterize the zooplankton communities present in Lac du Sauvage and Duchess Lake.

3.2 Lac de Gras Basin

3.2.1 Slipper Bay Stations

3.2.1.1 Trophic Status Classification

The discrete water sampling program for Slipper Bay yielded mean annual concentrations of 0.0041 mg-P/L for TP and 8.0 m for Secchi depth. The corresponding TSI values were 24.5 using TP and 30.0 using Secchi depth, for a rounded average of 27.3. The depth-integrated sampling program yielded mean annual concentrations of 0.0033 mg-P/L for TP, 1.49 μ g/L for chlorophyll *a*, and 8.0 m for Secchi depth (Dominion Diamond 2015). The corresponding TSI values were 24.1 using TP, 34.5 using chlorophyll *a*, and 30.0 using Secchi depth, for a rounded average of 28.6. Based on these TSI values, and the classification systems of Vollenweider (1970) and Carlson (1977), the Slipper Bay area of Lac de Gras is classified as oligotrophic. The Slipper Bay area can also be classified as oligotrophic (i.e., between 0.004 and 0.01 mg-P/L), based on CCME (2004) TP trigger ranges for Canadian lakes.



3.2.1.2 Phytoplankton

Abundance and Biomass

Total phytoplankton abundance in Slipper Bay varied seasonally and spatially (Figure 3.2-1). Total phytoplankton abundance was lowest in late spring (209,568 to 808,527 cells/L) at all stations, with the exception of Station S5. Peaks in total phytoplankton abundance occurred in the summer at stations S3 (1,631,311 cells/L), S5 (1,032,607 cells/L), and S6 (948,606 cells/L). Station S2 exhibited a peak in total phytoplankton abundance in fall (1,626,511 cells/L). A spatial gradient in phytoplankton abundance was observed in the Slipper Bay area, whereby stations located closest to the inflow (i.e., stations S2 and S3) had higher abundances than those closer to the open-water basin of Lac de Gras (i.e., stations S5 and S6). This gradient may be related to mine discharge from the Ekati Mine that enters Slipper Bay through the inflow near Station S2.

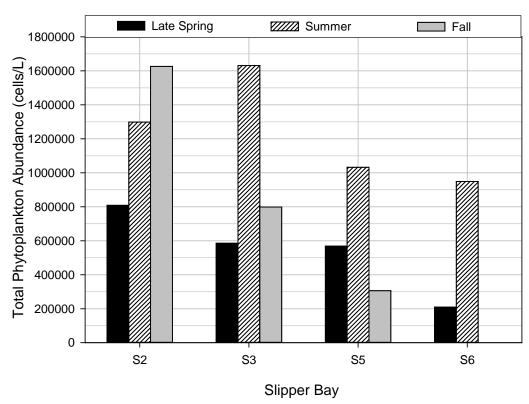


Figure 3.2-1 Total Phytoplankton Abundance in Slipper Bay Area, Lac de Gras in Late Spring, Summer, and Fall, 2014

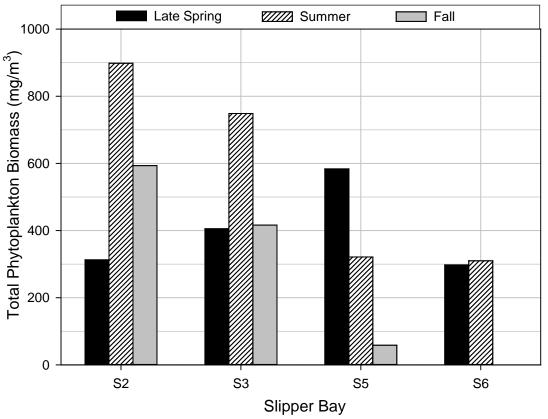
Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6. cells/L = cells per litre.

Seasonal and spatial variability were observed in total phytoplankton biomass in Slipper Bay; however, no clear seasonal trends were observed (Figure 3.2-2). Total phytoplankton biomass peaked in summer at



stations S2 (898 mg/m³) and S3 (749 mg/m³) and was 2.5 to 3 times higher than at stations S5 (322 mg/m³) and S6 (310 mg/m³) in summer. One chrysophyte, *Ochromonas* spp., and one diatom, *Cyclotella* sp., were the main drivers of the high phytoplankton biomass observed at Station S2 in the summer. The dominant taxa driving the high biomass at Station S3 in summer were the chrysophyte, *Ochromonas* sp., and the diatom, *Cyclotella*. In late spring, phytoplankton biomass was highest at Station S5 (584 mg/m³), while Station S2 had the highest biomass in fall (593 mg/m³). Total phytoplankton biomass at Station S6 exhibited little variation between late spring (298 mg/m³) and summer (310 mg/m³). A spatial gradient in phytoplankton biomass was observed in the Slipper Bay Area during the summer and fall. Stations located closest to the inflow (i.e., stations S5 and S6), possibly related to mine discharge from the Ekati Mine that enters Slipper Bay through the inflow near Station S2.

Figure 3.2-2 Total Phytoplankton Biomass in Slipper Bay Area, Lac de Gras in Late Spring, Summer, and Fall, 2014



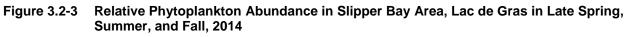
Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

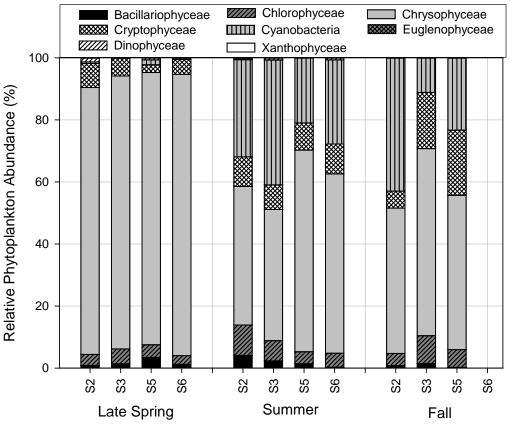
 $mg/m^3 = milligrams$ per cubic metre.



Community Composition

In general, the phytoplankton assemblage by abundance in Slipper Bay was relatively consistent among stations, but varied among sampling periods (Figure 3.2-3). Chrysophytes (86% to 91%) made up the majority of the phytoplankton assemblage by abundance in late spring. In general, two major taxonomic groups (i.e., chrysophytes and cyanobacteria) dominated the phytoplankton assemblage by abundance in the summer (42% to 65% and 21% to 40%, respectively) and fall (47% to 60% and 1% to 43%, respectively). Other taxonomic groups (cryptophytes, chlorophytes, and diatoms) made up a small fraction of the phytoplankton community composition by abundance in Slipper Bay and the percentages of these groups varied, particularly among sampling periods.





Slipper Bay

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

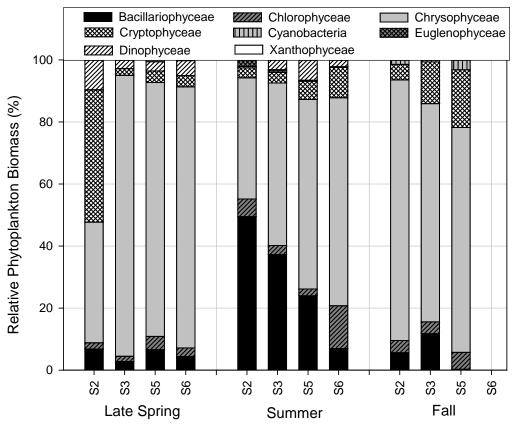
% = percent.

Chrysophytes dominated the phytoplankton composition by biomass at all stations in Slipper Bay in late spring (82% to 91%), with the exception of Station S2, which was co-dominated by chrysophytes (39%) and cryptophytes (43%) (Figure 3.2-4). The remainder of the community in late spring consisted of



variable amounts of diatoms (less than 49%), chlorophytes (less than 14%), and dinoflagellates (less than 10%). In the summer, stations S2 and S3 were co-dominated by chrysophytes (39% and 52%, respectively) and diatoms (49% and 37%, respectively). Chrysophytes made up the majority of the phytoplankton assemblage by biomass at stations S5 and S6 in the summer (61% and 67%, respectively). Diatoms were present at stations S5 and S6 (24% and 7%, respectively) in the summer, but represented a smaller percentage of the community compared to stations S2 and S3 (37% and 49%, respectively). In fall, chrysophytes dominated the phytoplankton community in Slipper Bay by biomass (70% to 84%). The remainder of the phytoplankton biomass in fall was made up mainly of varying proportions of cryptophytes (5% to 19%), diatoms (0% to 12%), and chlorophytes (4% to 5%).





Slipper Bay

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

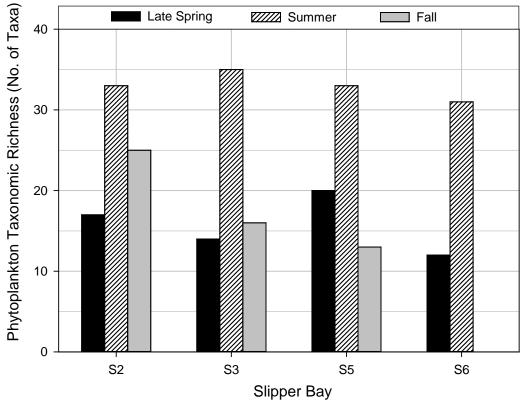
% = percent.

In total, 93 phytoplankton taxa were identified in Slipper Bay in 2014: 41 chlorophytes, 10 diatoms, 14 cyanobacteria, 14 chrysophytes, 6 dinoflagellates, 6 cryptophytes, 1 euglenoid, and 1 xanthophyte (Appendix A, Table A-5). Phytoplankton taxonomic richness was highest in the summer at all stations in



Slipper Bay (31 to 35 taxa) (Figure 3.2-5). Overall, taxonomic richness exhibited less spatial variation in the summer (31 to 35 taxa) compared to late spring (12 to 17 taxa) and fall (13 to 25 taxa).





Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

No. = number.

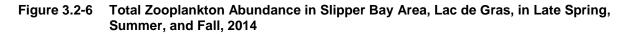
3.2.1.3 Zooplankton

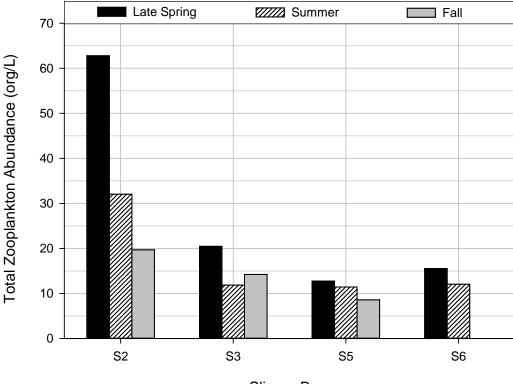
Abundance and Biomass

In general, total zooplankton abundance decreased throughout the open-water season in Slipper Bay; abundance ranged from 13 to 63 org/L in late spring, compared to summer (11 to 32 org/L) and fall (9 to 20 org/L) (Figure 3.2-6). Station S2 showed the largest decrease in total zooplankton abundance throughout the open-water season, ranging from 63 org/L in late spring to 20 org/L in fall. Station S2 also had notably higher zooplankton abundance compared to the other Slipper Bay stations, particularly in the late spring and summer. The high total zooplankton abundance observed at Station S2 in late spring was driven by the rotifers, *Kellicottia longispina* and *Conochilus unicornis*. A spatial gradient in zooplankton abundance was observed in the Slipper Bay Area. Station S2, located closest to the inflow, had higher



zooplankton abundance than stations closer to the open-water basin of Lac de Gras, possibly related to the inflow of discharge from the Ekati Mine.





Slipper Bay

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

org/L = organisms per litre.

Clear seasonal patterns in total zooplankton biomass were observed in Slipper Bay (Figure 3.2-7). In general, total zooplankton biomass increased throughout the open-water season, with the exception of Station S3 where a slight decrease was observed in summer. Total zooplankton biomass in Slipper Bay ranged from 38 to 53 mg/m³ in fall compared to late spring (16 to 27 mg/m³) and summer (21 to 40 mg/m³). Despite having the highest zooplankton abundance, zooplankton biomass at Station S2 was comparable to the other Slipper Bay stations throughout the open-water season.



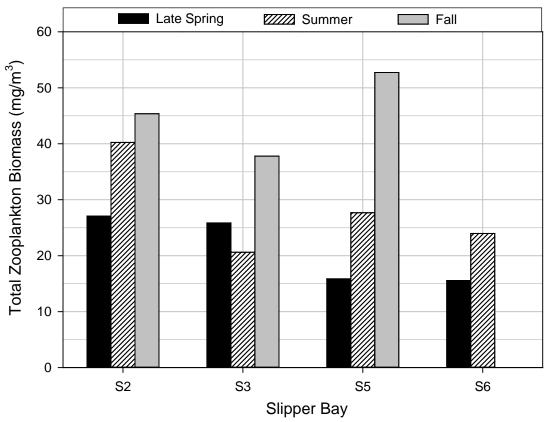


Figure 3.2-7 Total Zooplankton Biomass in Slipper Bay Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

 $mg/m^3 = milligrams$ per cubic metre.

Community Composition

Zooplankton community composition by abundance was generally similar among stations and seasons in Slipper Bay (Figure 3.2-8). Rotifers dominated the zooplankton community by abundance (48% to 91%) throughout the open-water season, with the exception of Station S5 in fall (24%). The zooplankton assemblage by abundance at Station S5 in fall was made up of calanoid copepods (34%) copepod nauplii (27%), and rotifers (24%). Station S2 was dominated by rotifers throughout the open water season, but unlike the other stations in Slipper Bay, copepod nauplii made up less than 2% of the abundance at Station S2 in late spring. Copepod nauplii (1% to 27%), calanoid copepods (1% to 34%), and cyclopoid copepods (6% to 18%) made up varying proportions of the total zooplankton abundance in Slipper Bay throughout the open-water season.



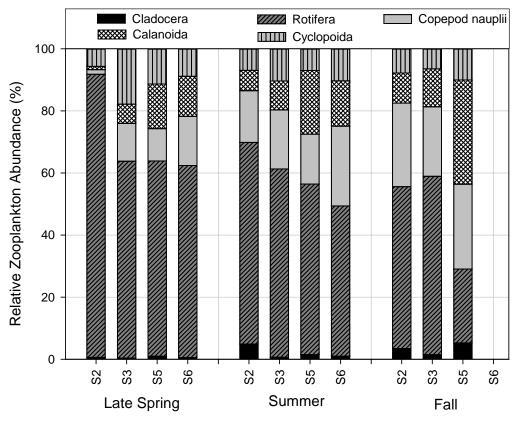


Figure 3.2-8 Relative Zooplankton Abundance in Slipper Bay Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

Slipper Bay

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

% = percent.

Zooplankton community by biomass was relatively consistent among stations in Slipper Bay, but the relative biomass of the major taxonomic groups varied with sampling period (Figure 3.2-9). Together, cyclopoid and calanoid copepods made up 58% to 92% of the zooplankton assemblage by biomass in Slipper Bay throughout the open-water season. The relative biomass of cyclopoid copepods decreased throughout the open-water season, from 44% to 73% in late spring to 7% to 15% in fall, as the relative proportion of calanoid copepods and Cladocera increased. Calanoid copepods made up 12% to 43% of the zooplankton biomass in late spring, compared to summer (45% to 67%) and fall (50% to 70%). The relative biomass of cyclopoid copepods decreased from late spring (44% to 73%) to fall (7% to 15%). Copepod nauplii were present in relatively low abundances in Slipper Bay throughout the open-water season (less than 6%). Overall, rotifers represented only a small fraction of the total zooplankton biomass in Slipper Bay, particularly in summer (less than 1%) and fall (less than 1%).



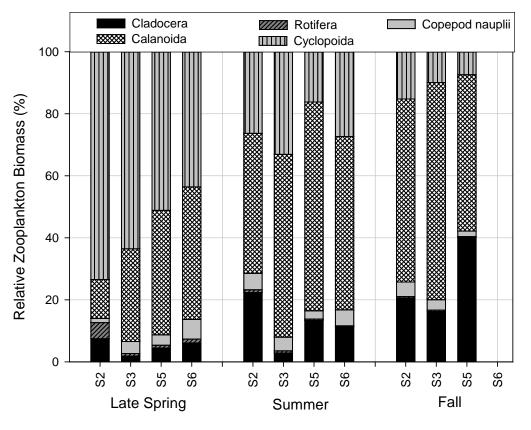


Figure 3.2-9 Relative Zooplankton Biomass in Slipper Bay Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

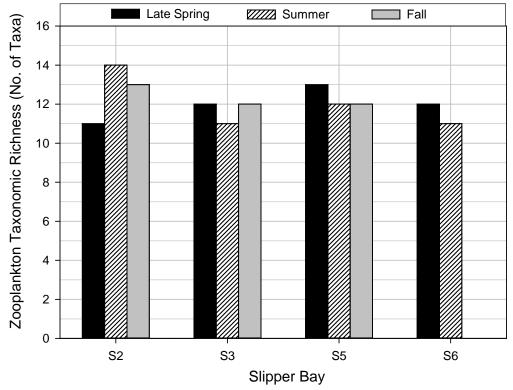
Slipper Bay

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6. % = percent.

In total, 31 zooplankton taxa were identified in Slipper Bay in 2014: 17 rotifers, 8 calanoid copepods, 4 cladocerans, and 2 cyclopoid copepods (Appendix A, Table A-10). There was little seasonal or spatial variability in total zooplankton richness in Slipper Bay throughout the open-water season (Figure 3.2-10). The highest and lowest zooplankton taxonomic richness in Slipper Bay was observed at Station S2 in summer (14 taxa) and late spring (11 taxa), respectively.







Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown; fall samples were not collected at Station S6.

No. = number.

3.2.2 Far-field 2 Stations

3.2.2.1 Trophic Status Classification

The discrete water sampling program in the FF2 area of Lac de Gras yielded mean annual concentrations of 0.0048 mg-P/L for TP and 5.8 m for Secchi depth. The corresponding TSI values were 26.8 using TP and 34.6 using Secchi depth, for a rounded average of 30.7. The depth-integrated sampling program yielded mean annual concentrations of 0.0047 mg-P/L for TP, 2.99 μ g/L for chlorophyll *a*, and 5.8 m for Secchi depth (Dominion Diamond 2015). The corresponding TSI values were 26.5 using TP, 41.3 using chlorophyll *a*, and 34.6 using Secchi depth, for a rounded average of 34.1.

Based on these TSI values, and the classification system of Vollenweider (1970), the FF2 area of Lac de Gras is classified as oligotrophic. According to the classification system of Carlson (1977), FF2 area is classified as oligotrophic based on TP and Secchi depth and mesotrophic based on chlorophyll *a*. The FF2 area can also be classified as oligotrophic (i.e., between 0.004 and 0.01 mg-P/L), based on CCME (2004) TP trigger ranges for Canadian lakes.



3.2.2.2 Phytoplankton

Abundance and Biomass

Total phytoplankton abundance in the FF2 area varied seasonally and spatially (Figure 3.2-11). Peaks in total phytoplankton abundance occurred in the summer at stations FF2-1, FF2-2, and FF2-3. Station FF2-1 had higher total phytoplankton abundance in the summer (1,764,661 cells/L), compared to the other stations (1,306,209 to 1,391,157 cells/L). The relatively high abundance observed at Station FF2-1 in late spring was largely driven by the chrysophyte, *Ochromonas* spp. and the cyanobacteria, *Leptolyngbya* sp. and *Aphanothece clathrata*. At stations FF2-1, FF2-1, and FF2-3, phytoplankton abundance was higher in the late spring (1,033,300 to 1,168,780 cells/L) compared to fall (605,545 to 815,498 cells/L). Total phytoplankton abundance in fall was highest at stations FF2-4 (1,514,140 cells/L) and FF2-5 (1,426,239 cells/L). Station FF2-4 exhibited an increasing trend throughout the open-water season, while Station FF2-5 exhibited little variation in total phytoplankton abundance between summer (1,375,229 cells/L) and fall (1,426,239 cells/L).

The timing of seasonal peaks in phytoplankton abundance appears to follow a spatial gradient, possibly related to the inflow of more productive waters from Lac du Sauvage. Phytoplankton abundance peaked during fall at the stations located closest to the outflow from Lac du Sauvage (i.e., FF2-4 and FF2-5), while abundance peaked in summer at the stations located furthest from the Lac du Sauvage outflow (i.e., FF2-1 and FF2-2).

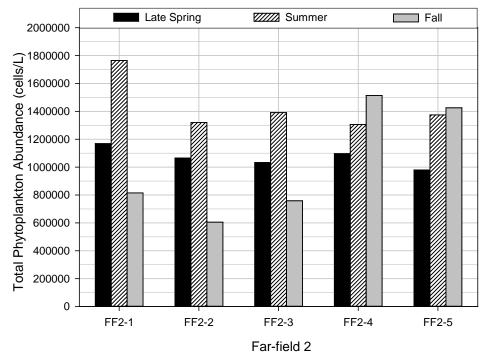


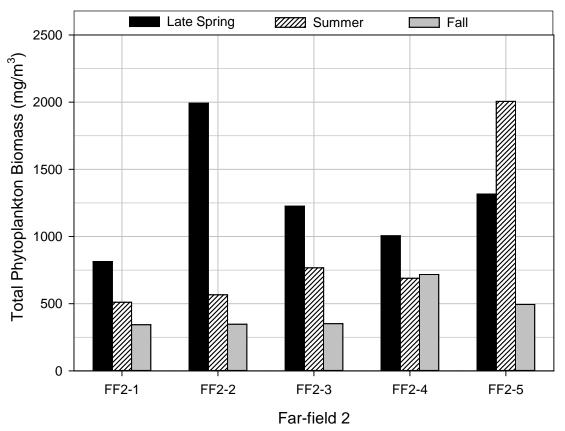
Figure 3.2-11 Total Phytoplankton Abundance in Far-Field 2 Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. cells/L = cells per litre.



Seasonal and spatial variability were observed in total phytoplankton biomass in the FF2 area (Figure 3.2-12). Total phytoplankton biomass was highest in late spring (1,006 to 1,993 mg/m³), with the exception of Station FF2-5 (814 mg/m³). Total phytoplankton biomass decreased throughout the openwater season at stations FF2-1, FF2-2, and FF2-3, from 814 to 1,993 mg/m³ in late spring to 344 to 351 mg/m³ in fall. Little variation in phytoplankton biomass was observed at Station FF2-4 between summer (690 mg/m³) and fall (718 mg/m³). The highest phytoplankton biomass in the late spring was observed at Station FF2-2 (1,993 mg/m³). Total phytoplankton biomass peaked in summer at Station FF2-5 (2,006 mg/m³); biomass values at this station were higher than the other FF2 stations in summer, but comparable to Station FF2-2 in the late spring. The high phytoplankton biomass observed at Station FF2-2 in late spring was largely driven by the chrysophyte, *Dinobryon divergens*, while the high biomass at Station FF2-5 in summer was largely driven by the diatom, *Cyclotella* sp. Phytoplankton biomass was generally low at all FF2 stations in fall (344 to 718 mg/m³).

Figure 3.2-12 Total Phytoplankton Biomass in Far-Field 2 Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

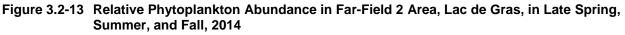


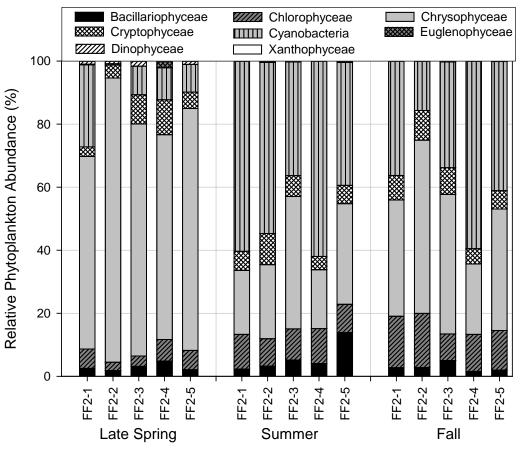
Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. $mg/m^3 = milligrams$ per cubic metre.



Community Composition

In general, two major taxonomic groups (i.e., chrysophytes and cyanobacteria) dominated the phytoplankton assemblage by abundance in the FF2 area throughout the open-water season, but the percentages of these groups varied among stations and sampling periods (Figure 3.2-13). The phytoplankton community by abundance was co-dominated by cyanobacteria and chrysophytes in summer (36% to 62% and 19% to 42%, respectively) and fall (16% to 60% and 22% to 55%, respectively). Chrysophytes made up the majority of the phytoplankton assemblages in late spring (61% to 90%); cyanobacteria made up a smaller percentage of the community composition in the FF2 area in late spring compared to summer and fall. The remainder of the phytoplankton abundance was made up of varying percentages of chlorophytes (3% to 17%), cryptophytes (3% to 11%), diatoms (2% to 14%), and dinoflagellates (less than 2%). Together, euglenoids and xanthophytes (less than 2%) made up a small fraction of the total phytoplankton abundance in the FF2 area throughout the open-water season.



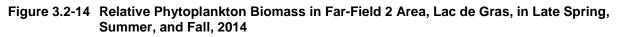


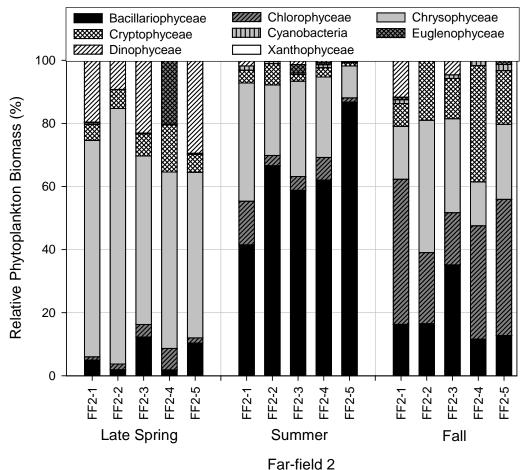
Far-field 2

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. % = percent.



Phytoplankton composition by biomass in the FF2 area varied seasonally and spatially (Figure 3.2-14). Chrysophytes (53% to 81%) dominated the phytoplankton community by biomass in late spring. Together, dinoflagellates and cryptophytes made up 15% to 35% of the phytoplankton assemblage by biomass in late spring. Throughout the open-water season, xanthophytes and euglenoids made up a small percentage of the total biomass in the FF2 area, with the exception of Station FF2-4 in the late spring (20% euglenoids). Diatoms (41% to 87%) and chrysophytes (22% to 37%) co-dominated the phytoplankton assemblage by biomass in summer. A mixture of chlorophytes (17% to 46%), chrysophytes (17% to 42%), diatoms (12% to 35%), and cryptophytes (7% to 37%) made up the majority of the phytoplankton biomass in the FF2 area in fall.





Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. % = percent.



In total, 108 phytoplankton taxa were identified in the FF2 area in 2014: 48 chlorophytes, 10 diatoms, 19 cyanobacteria, 15 chrysophytes, 6 dinoflagellates, 6 cryptophytes, 2 euglenoids, and 2 xanthophytes (Appendix A, Table A-6). Seasonal variation in phytoplankton taxonomic richness was observed in this area, but spatial variation was not apparent (Figure 3.2-15). Phytoplankton richness peaked in summer at all stations in the FF2 area (36 to 42 taxa); richness observed in the summer was higher compared to late spring (17 to 25 taxa) and fall (24 to 28 taxa).

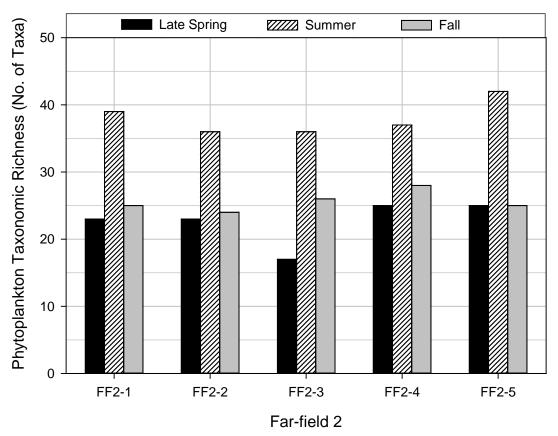


Figure 3.2-15 Total Phytoplankton Taxonomic Richness in Far-Field 2 Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. No. = number.

3.2.2.3 Zooplankton

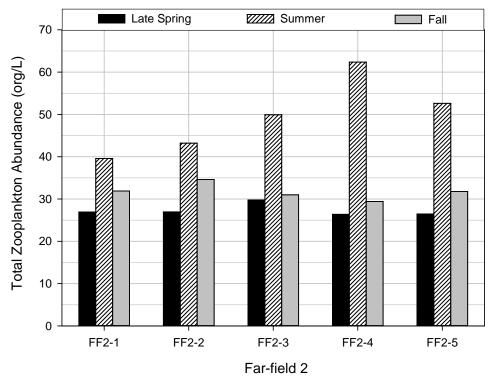
Abundance and Biomass

Stations in the FF2 area showed similar trends in total zooplankton abundance throughout the open-water season (Figure 3.2-16). Total zooplankton abundance peaked in summer at all stations in this area, ranging from 40 to 62 org/L, compared to late spring (26 to 30 org/L) and fall (29 to 35 org/L). The spatial variability in the range of zooplankton abundance in the FF2 area was also greatest in the summer.



During the summer, a spatial gradient in zooplankton abundance was observed in the FF2 area, possibly related to the inflow of more productive waters from Lac du Sauvage. Zooplankton abundance was higher at the stations located closest to the outflow from Lac du Sauvage (i.e., FF2-3, FF2-4, and FF2-5), compared to the stations located further from the Lac du Sauvage outflow (i.e., FF2-1 and FF2-2).

Figure 3.2-16 Total Zooplankton Abundance in Far-Field 2 Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014



Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. org/L = organisms per litre.

Seasonal and spatial patterns in total zooplankton biomass were observed in the FF2 area (Figure 3.2-17). Total zooplankton biomass at FF2 stations was generally lower in late spring (21 to 26 mg/m³) compared to summer (36 to 57 mg/m³) and fall (36 to 53 mg/m³). At stations FF2-1 and FF2-2, zooplankton biomass increased from late spring (19 mg/m³ and 26 mg/m³, respectively) to summer (38 mg/m³ and 53 mg/m³, respectively) and then remained steady in fall (40 mg/m³ and 53 mg/m³, respectively). A steady increase in zooplankton biomass throughout the open-water season was observed at Station FF2-3 (from 21 to 53 mg/m³), while biomass peaked in summer at stations FF2-4 (57 mg/m³) and FF2-5 (52 mg/m³).



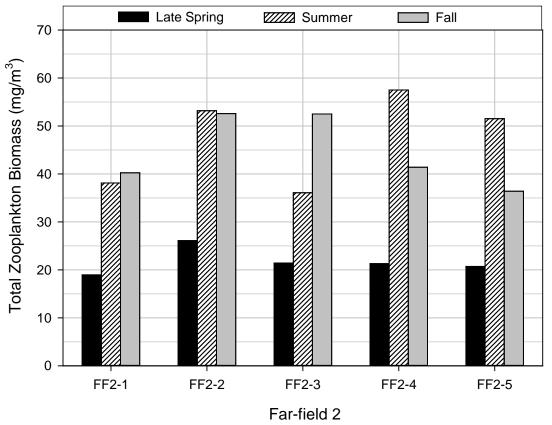


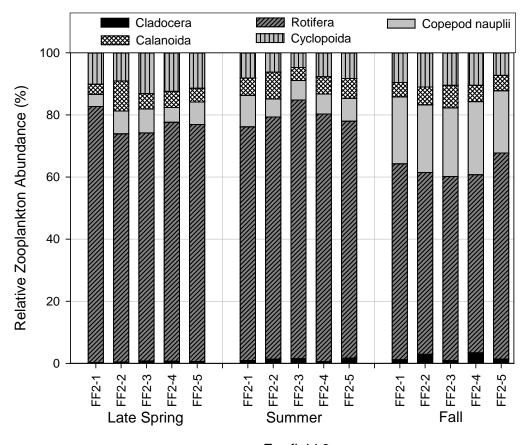
Figure 3.2-17 Total Zooplankton Biomass in Far-Field 2 Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

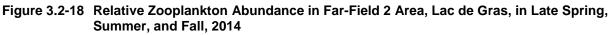
Note: Bars represent a single composite sample collected within the euphotic zone; a mean value for field duplicates are shown. $mg/m^3 = milligrams$ per cubic metre.

Community Composition

The zooplankton community by abundance in the FF2 area remained relatively consistent throughout the open-water season (Figure 3.2-18). Rotifers dominated the zooplankton abundance at all stations in this area during all three sampling periods (late spring: 73% to 82%; summer 75% to 83%; fall: 57% to 66%). Copepods (including copepod nauplii, cyclopoid copepods and calanoid copepods) made up the majority of the remaining zooplankton abundance throughout the open-water season. The relative proportion of cyclopoid and calanoid copepods at the FF2 stations varied little seasonally. The relative proportion of copepod nauplii at the FF2 stations was higher in fall (20% to 24%) compared to late spring and summer (4% to 7% and 6% to 10%, respectively). Cladocera made up less than 4% of the zooplankton abundance in the FF2 area throughout the open-water season.



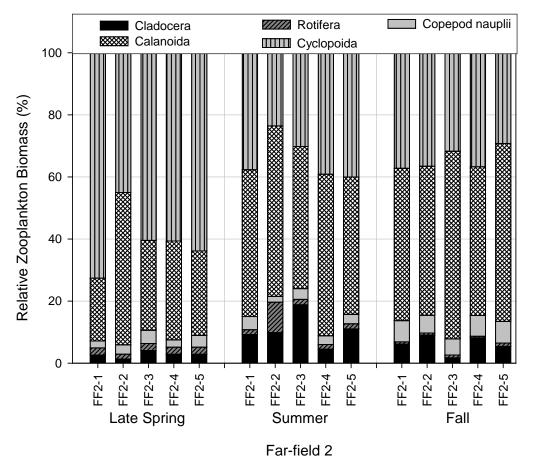




Far-field 2 Note: Bars represent a single a vertical haul taken throughout the water column. % = percent.

Two major taxonomic groups (i.e., calanoid and cyclopoid copepods) dominated the zooplankton assemblage by biomass at all stations in the FF2 area throughout the open-water season, but the percentages of these groups varied among sampling periods (Figure 3.2-19). Cyclopoid copepods made up a higher percentage of the relative biomass in late spring (45% to 73%) compared to summer and fall (24% to 40% and 29% to 37%, respectively). The proportion of calanoid copepods was higher in summer and fall (44% to 55% and 48% to 60%, respectively) compared to late spring (20% to 49%). Cladocera (1% to 19%), rotifers (1% to 10%), and copepod nauplii (2% to 7%) made up the remaining zooplankton biomass in the FF2 area throughout the open-water season.







Note: Bars represent a single a vertical haul taken throughout the water column. % = percent.

In total, 37 zooplankton taxa were identified in the FF2 area in 2014: 21 rotifers, 8 calanoid copepods, 6 cladocerans, and 2 cyclopoid copepods (Appendix A, Table A-11). With the exception of Station FF2-3, zooplankton taxonomic richness in this area was similar among stations and sampling periods, ranging from 13 to 16 taxa (Figure 3.2-20). At Station FF2-3, zooplankton richness decreased throughout the open-water season from 18 taxa in late spring to 11 taxa in fall.



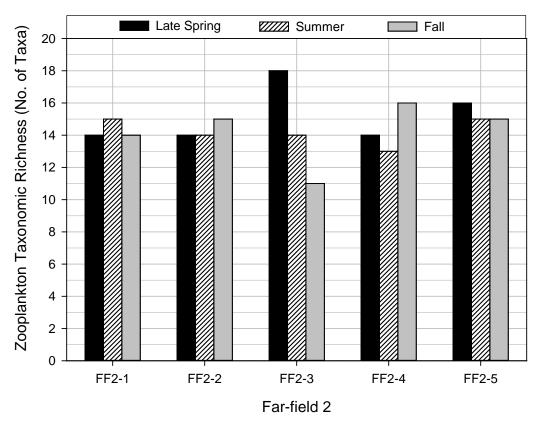


Figure 3.2-20 Total Zooplankton Taxonomic Richness in Far-Field 2 Area, Lac de Gras, in Late Spring, Summer, and Fall, 2014

Note: Bars represent a single a vertical haul taken throughout the water column; a mean value for field duplicates are shown. No. = number.

3.2.3 Comparison of 2014 Plankton Data to Previously Collected Data

Results from the 2014 supplemental baseline plankton sampling program were compared to data collected by previous studies in Lac de Gras (stations FF2-2, FF2-5, S2, and S3) to describe the range in temporal variation in plankton communities present in this lake. Historical data for Lac de Gras, previously summarized in Annex XII of the DAR (Dominion Diamond 2014), were obtained from the following sources:

- baseline and long-term AEMP data collected in the FF2 area from 1997 to 2013 for the Diavik Mine (Golder 2011; DDMI 2012, 2013; Golder 2014a,b); and,
- baseline and long-term AEMP data collected in Slipper Bay at stations S2 and S3 from 1997 to 2012 for the Ekati Mine (ERM Rescan 2013).



Phytoplankton biomass results for Slipper Bay could not be compared in this section, because historical phytoplankton biomass data are not available. In addition, previous estimates of zooplankton biomass were based on wet weight biomass derived from length-weight regressions (FF2 area) and dry weight AFDM estimates (Slipper Bay). Therefore, the historical and recent (i.e., 2012 and 2013) zooplankton biomass data for Lac de Gras were not comparable to the dry weight biomass results from 2014.

3.2.3.1 Trophic Status Classification

Based on the TSI calculations for discrete and depth-integrated TP, Secchi depths, and chlorophyll *a* concentrations from the open-water season in 2014, the trophic status of Slipper Bay was in the oligotrophic range. The FF2 area was also classified as oligotrophic based on Vollenweider (1970) and CCME (2004), and within the range of an oligotrophic to mesotrophic system based on the Carlson (1977) classification. Results are consistent with previous baseline and historical data for Lac de Gras; the FF2 area and Slipper Bay (and Lac de Gras in general) were classified as oligotrophic based on chlorophyll *a* data collected from 1997 to 2012 (Golder 2011; DDMI 2012, 2013; ERM Rescan 2013).

3.2.3.2 Phytoplankton

Based on comparing the previously collected and 2014 phytoplankton data for Lac de Gras, the following conclusions were made:

- Between 1997 and 2012, total phytoplankton abundance ranged from 285,133 to 1,877,100 cells/L at Station S2 and from 194,130 to 1,161,400 cells/L at Station S3. Total phytoplankton abundances observed in Slipper Bay in 2014 (209,568 to 1,631,311 cells/L) were within the historical range.
- In 2014, total phytoplankton abundance in the FF2 area ranged from 605,545 to 1,764,661 cells/L and was within the range observed in this area between 2003 and 2013 (1,228,280 cells/L and 6,254,450 cells/L).
- Between 2003 and 2013, phytoplankton biomass in the FF2 area ranged from 171 mg/m³ in 2004 to 701 mg/m³ in 2013. Total phytoplankton biomass in this area in 2014 ranged from 344 to 2,006 mg/m³, and were generally higher than historical values.
- Cyanobacteria were the dominant group at stations S2 and S3 in 1997 and 1998. Since 1998, chrysophytes, chlorophytes, and diatoms accounted for the majority of the phytoplankton community by abundance at stations S2 and S3. In 2014, seasonal differences were observed in phytoplankton composition by abundance in Slipper Bay; chrysophytes made up the majority of the phytoplankton assemblage in late spring, while chrysophytes and cyanobacteria dominated the phytoplankton assemblage in the summer and fall.
- Chrysophytes and cyanobacteria co-dominated the phytoplankton community by abundance in the FF2 area from 2003 to 2006, and 2008 to 2010, and 2013; in 2007, "others" dominated the community and in 2011 cyanobacteria were dominant. Chrysophytes and cyanobacteria also co-dominated the phytoplankton assemblage by abundance in the FF2 area throughout the open-water season in 2014.
- Between 2003 and 2013, cyanobacteria and chrysophytes generally co-dominated the phytoplankton community by biomass in the FF2 area, while the phytoplankton composition by biomass in 2014 varied seasonally and spatially. Chrysophytes dominated in late spring, diatoms and chrysophytes co-



dominated in summer, and a mixture of chlorophytes, chrysophytes, diatoms, and cryptophytes made up the majority of the phytoplankton biomass in the FF2 area in fall.

Overall, the variation observed in community metrics in the baseline phytoplankton dataset underscores the importance of having multiple years of data to characterize the phytoplankton communities present in the Slipper Bay and FF2 areas of Lac de Gras.

3.2.3.3 Zooplankton

Based on comparing the previously collected and 2014 zooplankton data for Lac de Gras, the following conclusions were made:

- Summer zooplankton abundances at stations S2 and S3 have varied since 1997, ranging between 9 and 41 org/L. Despite seasonal fluctuations, total zooplankton abundance in Slipper Bay in 2014 (9 to 63 org/L) were generally comparable to historical values.
- Historical (i.e., 2008 to 2013) zooplankton abundances in the FF2 area ranged from 5 to 62 org/L. Total zooplankton abundance in this area in 2014 ranged from 26 to 62 org/L and were generally comparable to historical values.
- In general, rotifers dominated the zooplankton community by abundance in Slipper Bay throughout the open-water season in 2014, consistent with historical (i.e., 1997 to 2012) results for stations S2 and S3.
- In 2014, rotifers dominated the zooplankton abundance at all stations in the FF2 area during all three sampling periods, consistent with historical (i.e., 2008 to 2013) results for this area.
- Between 2008 and 2012, the zooplankton community by biomass in the FF2 area was dominated by calanoid copepods; cyclopoid copepods were the sub-dominant group. The same two major taxonomic groups (i.e., calanoid and cyclopoid copepods) co-dominated the zooplankton assemblage by biomass at all stations in this area throughout the open-water season in 2014, while total zooplankton biomass in the FF2 area in 2013 was dominated by a mixture of cladocerans, calanoid copepods, and cyclopoid copepods.

Overall, the variation observed in community metrics in the baseline zooplankton dataset underscores the importance of having multiple years of data to characterize the zooplankton communities present in the Slipper Bay and FF2 areas of Lac de Gras.

3.3 Quality Assurance and Quality Control

Detailed QC results for phytoplankton and zooplankton are presented in Appendix A, Tables A-1 and A-2. Overall, differences between original and QC samples were within the range of variability expected for plankton samples and subsampling variance.

3.3.1.1 Laboratory Quality Control

For phytoplankton QC comparisons, EcoAnalysts re-counted two samples from late spring (stations Ac-1 and FF2-4), two samples from summer (stations Ad-1 and FF2-5), and two samples from fall (stations



Ac-1 and FF2-4). All six phytoplankton QC samples were above the 50% sample similarity criterion (ranging from 83% to 93%). QC results indicate that the dominant phytoplankton taxa (i.e., *Ochromonas* sp., *Komma caudata, and Cyclotella*) were the same in both the original and the QC samples.

For zooplankton QC comparisons, EcoAnalysts re-counted two samples from late spring (stations Ac-4 and S6), two samples from summer (stations FF2-5 and S6), and two samples from fall (stations Ae-1 and S2). All six zooplankton QC samples were above the 50% similarity criterion in both the fine and coarse fractions (ranging from 84% to 99%). QC results indicate that the dominant zooplankton taxa in the coarse fraction (i.e., *Daphnia longiremis, Cyclops strenuous*, cyclopoid copepodites, and calanoid copepodites) and fine fraction (i.e., *Kellicottia longispina, Keratella cochlearis, Conochilus unicornis*, and copepod nauplii) were the same in both the original and QC samples.

Internal QC results indicate that the overall occurrence of dominant taxa was consistent between the original and duplicate samples for phytoplankton and zooplankton. Sample percent similarities met the 50% criterion for all internal laboratory QC comparisons for both phytoplankton and zooplankton. The taxonomist concluded that differences observed between the original and QC samples for phytoplankton and zooplankton were due to sub-sampling variance. Overall, internal QC comparison results by EcoAnalysts suggest that the phytoplankton and zooplankton data are of acceptable quality.

3.3.1.2 Quality Control Data Evaluation

Duplicate phytoplankton and zooplankton samples were collected from stations FF2-5 and S5 in late spring, Stations FF2-4 and S3 in summer, and Station Ac-1 in fall. Only one set of duplicate samples was collected in fall, because of field sampling error.

The Bray-Curtis index values for the phytoplankton samples were acceptable for abundance, and ranged from 0.04 to 0.29 (Appendix A, Tables A-1). For zooplankton abundance, the Bray-Curtis index values ranged from 0.01 to 0.08 and no data required further investigation (Appendix A, Table A-2). For phytoplankton and zooplankton abundance, duplicate samples yielded RPD values lower than 50% in the major taxonomic group comparison (Appendix A, Tables A-1 and A-2).

Overall, differences between original and QC phytoplankton and zooplankton samples were deemed minor, and within the range of variability expected in plankton data. Therefore, QC results indicated that the 2014 phytoplankton and zooplankton data were considered to be of acceptable quality and no data were invalidated.



4 SUMMARY

The 2014 plankton sampling program was designed to supplement the existing plankton baseline dataset. The 2014 program was carried out over three sampling programs (late spring, summer, and fall) during open-water conditions. In total, 18 stations were sampled for plankton across three lakes (Lac du Sauvage, Duchess Lake, and Lac de Gras). A high-level summary of the findings from the 2014 sampling program is provided in this section.

4.1 Trophic Status Classification

Based on the TSI calculations for discrete and depth-integrated TP concentrations, Secchi depths, and chlorophyll *a* concentrations from the open-water season in 2014, the trophic status of Lac du Sauvage and Slipper Bay were oligotrophic, while Duchess Lake was mesotrophic (Table 4.1-1). The FF2 area of Lac de Gras was classified as oligotrophic based on Vollenweider (1970) and CCME (2004), and as oligotrophic to mesotrophic based on the Carlson (1977) classification.

			Lac de Gras		
Community Variable	Lac du Sauvage	Duchess Lake	Slipper Bay Area	Far-field 2 Area	
Trophic Status	oligotrophic	mesotrophic	oligotrophic	oligotrophic to mesotrophic ^(a)	
Phytoplankton					
Total Abundance (cells/L)	457,734 to 1,643,909 954,969 to 3,28		209,568 to 1,761,738	605,545 to 1,764,661	
Total Biomass (mg/m ³)	198 to 1,515	512 to 1,327	59 to 898	344 to 2,006	
Total Richness (No. of taxa)	118	42	93	108	
Zooplankton					
Total Abundance (org/L)	32 to 95	66 to 76	9 to 63	25 to 67	
Total Biomass (mg/m ³)	al Biomass (mg/m ³) 13 to 152		16 to 53	18 to 59	
otal Richness (No. of taxa) 31		19	31	37	

 Table 4.1-1
 Summary of Baseline Plankton Community, 2014

a) Oligotrophic system based on Vollenweider (1970) and CCME (2004); oligotrophic to mesotrophic system based on Carlson (1977).

cells/L = cells per litre; mg/m³ = milligrams per cubic metre; No. = number; org/L = organisms per litre.

4.2 Phytoplankton

Total phytoplankton abundance and biomass varied both within and among lakes in 2014 (Table 4.1-1). Seasonal and spatial variability was observed in total phytoplankton abundance and biomass in Lac du Sauvage, Duchess Lake, and Lac de Gras. Total phytoplankton abundance peaked in the summer at the majority of lake stations, while the timing of peak total phytoplankton biomass was not consistent among lakes.

The range in total phytoplankton abundance was comparable among lakes, with the exception of a peak in total phytoplankton abundance in Duchess Lake. Slipper Bay had the lowest total phytoplankton biomass, consistent with its classification as an oligotrophic system. Lac du Sauvage was also classified as an oligotrophic system; however, the range in total phytoplankton biomass was higher than in Slipper



Bay, because of the occurrence of seasonal peaks. High total phytoplankton biomass was observed in the FF2 area of Lac de Gras, including numerous seasonal peaks greater than 1,000 mg/m³, reinforcing the oligotrophic to mesotrophic classification. Total phytoplankton biomass in Duchess Lake was lower than in the FF2 area and comparable to Lac du Sauvage (an oligotrophic system), despite being classified as an oligotrophic to mesotrophic system.

The total number of phytoplankton taxa identified in Lac du Sauvage, Duchess Lake, and Lac de Gras ranged from 42 taxa (Duchess Lake) to 118 taxa (Lac du Sauvage). The lower richness in Duchess Lake reflects the lower sampling effort in this lake in 2014 compared to Lac du Sauvage and Slipper Bay. Chrysophytes consistently dominated the community by abundance in Lac du Sauvage and Duchess Lake throughout the open-water season. Chrysophytes also made up the majority of the phytoplankton assemblage by abundance in Slipper Bay and the FF2 area in late spring, while chrysophytes and cyanobacteria co-dominated the phytoplankton assemblage by abundance in these areas of Lac de Gras in summer and fall.

In terms of biomass, the phytoplankton community composition in Lac du Sauvage, Duchess Lake and the FF2 area varied seasonally and spatially throughout the open-water season. However, no consistent trends in dominant major taxonomic groups were identified. Chrysophytes dominated the phytoplankton composition by biomass at the majority of stations in Slipper Bay and Duchess Lake throughout the open-water season.

4.3 Zooplankton

Total zooplankton abundance and biomass varied both within and among lakes in 2014 (Table 4.1-1). Seasonal and spatial variability was observed in total zooplankton abundance and biomass in Lac du Sauvage, Duchess Lake, and Lac de Gras. The timing of seasonal peaks in total zooplankton abundance and biomass were not consistent among lakes. Total zooplankton abundance peaked in summer at Lac du Sauvage and the FF2 area of Lac de Gras, fall in Duchess Lake, and late spring in Slipper Bay. Summer peaks in total zooplankton biomass were observed in Lac du Sauvage and Duchess Lake, while biomass peaked in fall in Slipper Bay; the FF2 area exhibited no consistent seasonal peak.

In general, lakes in the Lac du Sauvage basin (Lac du Sauvage and Duchess Lake) had higher total zooplankton abundance throughout the open-water season compared to lakes in the Lac de Gras basin (Slipper Bay and FF2 area). With the exception of Station S2, Slipper Bay had notably lower zooplankton abundance compared to the FF2 area and Lac du Sauvage basin lakes. Total zooplankton biomass was similar among Slipper Bay, FF2 area, and Duchess Lake, and was generally lower than the biomass observed in Lac du Sauvage.

The total number of zooplankton taxa identified in lakes in the Lac du Sauvage and Lac de Gras basins ranged from 19 taxa (Duchess Lake) to 37 taxa (FF2). The lower richness in Duchess Lake reflects the lower sampling effort in this lake in 2014 compared to Lac du Sauvage and Slipper Bay. Rotifers dominated the zooplankton community by abundance at the majority of stations in Slipper Bay and the FF2 area throughout the open-water season. Rotifers also dominated the zooplankton abundance in Lac du Sauvage in late spring and summer, while the assemblage in fall was co-dominated by rotifers and copepod nauplii. Zooplankton abundance in Duchess Lake was co-dominated by Cladocera and rotifers.



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The same two major taxonomic groups (i.e., calanoid and cyclopoid copepods) made up the majority of the zooplankton assemblage by biomass in Slipper Bay and the FF2 area in Lac de Gras throughout the open-water season. The zooplankton community composition by biomass in the Lac du Sauvage basin lakes differed from Lac de Gras basin lakes. In late spring and summer, the zooplankton community by biomass in Lac du Sauvage was co-dominated by Cladocera and cyclopoid copepods, while the assemblage was dominated by cyclopoid copepods in fall. The zooplankton assemblage by biomass in Duchess Lake was dominated by Cladocera.



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6 GLOSSARY

Term	Description
Abundance	The number of individuals in a given area or sample.
Aquatic Effects Monitoring Program	A monitoring program designed to evaluate the effect of mining activities and mitigation on the aquatic environment. Also referred to as an AEMP.
Bacillariophyceae	Diatoms; a group of algae that are encased within a frustule (shell) made of silica; a component of phytoplankton.
Baseline	A base for measurement or comparison.
Biomass	The weight of living matter in a given area or sample.
Bray-Curtis Index	A distance measure used to quantify the compositional dissimilarity between two different communities, based on counts of organisms.
Calanoida	An order of copepods (crustaceans); small animals in the water column that are a component of zooplankton.
Chlorophyll a	The primary photosynthetic pigment contained in the phytoplankton (primary producers; small plants in the water column).
Chlorophyceae	Chlorophytes; Green algae; a component of phytoplankton.
Chrysophyceae	Chrysophytes; Golden-brown algae; a component of phytoplankton.
Cladocera	A group of small planktonic animals (crustaceans) also known as water fleas; a component of zooplankton.
Community composition	The assortment of different organisms that constitute an ecological community.
Composite sample	A sample taken by combining several volumes of water from different depths within the water column of a lake into a common vessel. A composite sample can also be obtained as a combination of samples taken from different parts of a waterbody laterally.
Colonial	Individuals of the same species clustered together to form a group.
Conductivity	A measure of the ability of a solution to conduct electrical current; an indirect measure of the salinity of the water.
Copepoda	An order of planktonic crustaceans; a component of zooplankton.
Cryptophyceae	Cryptophytes; flagellated algae also known as cryptomonads; a component of phytoplankton.
Crustaceans	A large group of primarily aquatic arthropods of the class Crustacea, which are free-living, have a segmented body and an exoskeleton.
Cyanobacteria	Blue-green algae; a component of phytoplankton.
Cyclopoida	An order of copepods (crustaceans); small animals in the water column that are a component of zooplankton.
Depth-integrated composite sample	A sample made up of subsamples of equal volume collected from discrete depths throughout the water column that represents the average composition of the water column.
Dinophyceae	Dinoflagellates; a group of unicellular flagellated algae, many of which are motile; 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.
Euglenophyceae	Euglenoids; one of the best-known groups of flagellated algae; a component of phytoplankton.
Euphotic zone	The upper layer of a waterbody as defined by light penetration, with the upper limit determined by the water surface and the lower limit determined as the depth to which sufficient light for photosynthesis can penetrate (nominally 1% of the surface ambient light, measured as photosynthetically active radiation, or PAR).
Eutrophic	Trophic state classification for lakes characterized by high level of productivity and nutrient inputs (particularly total phosphorus).
Headwater	The source of water at the top of a watershed, typically a lake or marsh.
Hydrology	The study of flowing water and effects of flowing water on the Earth's surface, in the soil and underlying



Term	Description					
	rocks, and in the atmosphere.					
In situ measurement	The on-site measurement of physical water quality constituents in a waterbody.					
Kimberlite	Igneous rocks that originate deep in the Earth's mantle and intrude the Earth's crust. These rocks typically form narrow pipe-like deposits that sometimes contain diamonds.					
Kimberlite pipe	A more or less vertical, cylindrical body of kimberlite that resulted from the forcing of the kimberlite material to the Earth's surface.					
Mesotrophic	Trophic state classification for lakes characterized by moderate productivity and nutrient inputs (particularly total phosphorus).					
Nutrients	Environmental substances (elements or compounds) such as nitrogen or phosphorus, which are necessary for the growth and development of plants and animals.					
Oligotrophic	Trophic state classification for lakes characterized by low productivity and nutrient inputs (particularly total phosphorus).					
Open-water conditions	The period of time when the surface of a waterbody is completely free of ice.					
рН	A measure of the acidity or alkalinity of water.					
Photosynthesis	A chemical reaction that occurs in the chloroplasts of algae and plants and involves the conversion of water and carbon dioxide into organic carbon.					
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.					
Quality Assurance	Management and technical practices designed so that the data generated are of consistent high quality. They include standardization and review by field and office personnel of procedures used in the collection, transport, and analyses of samples. Also referred to as QA.					
Quality Control	Internal techniques used to measure and assess data quality, including samples that are used to detect and reduce systematic and random errors that may occur during field sampling and laboratory procedures. Also referred to as QC.					
Relative abundance	The proportional representation of the abundance of each species in a sample or a community.					
Relative biomass	The proportional representation of the biomass of each species in a sample or a community.					
Richness	The number of different types of animals present in a sample or at a location.					
Rotifera	A phylum of microscopic and near-microscopic pseudocoelomate animals; 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.					
Specific conductivity	A measure of the capacity of water to conduct an electrical current. It is the reciprocal of resistance. This measurement provides an estimate of the total concentration of dissolved ions in the water (specific conductance is normalized to 25°C).					
Stratification	The separation of lakes into three layers: well mixed top layer, middle layer (see Thermocline), and a bottom layer. In freshwater lakes, stratification usually occurs as a result of temperature effects that cause changes in water density. Stratification may also result in vertical variation in water quality.					
Таха	A group of organisms of any taxonomic rank (e.g., family, genus, species).					
Taxon	A group of organisms at the same level of the standard biological classification system; the plural of taxon is taxa.					
Taxonomic group	Biological organisms that have shared characteristics and are therefore grouped under a common name at a higher taxonomic level.					
Total Kjeldahl nitrogen	The sum of organic nitrogen; ammonia (NH3) and ammonium (NH4+).					
Total nitrogen	A measurement of the sum of all forms of particulate and dissolved nitrogen in water. Also referred to as TN.					
Total phosphorus	A measurement of the sum of particulate and dissolved phosphorus and phosphate in water. Also referred to as TP.					
Total richness	The total number of different taxa occupying a given area.					
Trophic status	Eutrophication is the process by which lakes are enriched with nutrients, increasing the production of					

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Term	Description
	rooted aquatic plants and algae. The extent to which this process has occurred is reflected in a lake's trophic classification or status, which can be oligotrophic (nutrient poor), mesotrophic (moderately productive), or eutrophic (very productive).
Waterbody	A general term that refers to rivers, streams, and lakes.
Xanthophyceae	Xanthophytes; yellow-green algae; a component of phytoplankton.
Zooplankton	Small, sometimes microscopic, animals that live in the water column of lakes and mainly eat primary producers (phytoplankton).



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APPENDIX A

2014 PLANKTON TAXONOMY DATA



Tables

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Abbreviations

Abbreviation	Definition				
Dominion Diamond	Dominion Diamond Ekati Corporation				
FF	Far-field				
Golder	Golder Associates Ltd.				
QC	quality control				
RPD	relative percent difference				
sp.	species				
spp.	multiple species				

Units of Measure

Unit	Definition				
%	percent				
cells/L	cells per litre				
mg/m ³	milligrams per cubic metre				
org/L	organisms per litre				



	Station	Sampling Period	Major Taxonomic Group	Total Abundance (cells/L)			Bray Curtis
Basin				Replicate 1	Replicate 2	RPD (%)	Dissimilarity Index
			Bacillariophyceae	15,086	22,581	10	
			Chlorophyceae	71,852	55,021	7	
			Chrysophyceae	568,336	814,842	9	•
			Cryptophyceae	25,998	59,489	20	
Lac de Gras – Far- field 2	FF2-5	Late Spring	Cyanobacteria	97,528	82,018	4	0.18
			Dinophyceae	15,771	6,824	20	
			Euglenophyceae	0	0	0	
			Xanthophyceae	396	1,034	22	
			Total	794,966	1,041,809	7	
			Bacillariophyceae	20,865	19,193	2	
			Chlorophyceae	15,682	25,326	12	
			Chrysophyceae	612,537	461,872	7	
			Cryptophyceae	19,320	12,631	10	
Lac de Gras - Slipper Bay	S5	Late Spring	Cyanobacteria	2,816	10,800	29	0.15
Зпррег вау			Dinophyceae	968	3,831	30	
			Euglenophyceae	0	0	0	
			Xanthophyceae	0	594	50	
			Total	672,188	534,247	6	
			Bacillariophyceae	26,743	86,515	26	
			Chlorophyceae	182,993	83,263	19	
			Chrysophyceae	57,647	485,314	39	
			Cryptophyceae	17,416	102,291	35	
Lac de Gras – Far- field 2	FF2-4	Summer	Cyanobacteria	819,828	747,780	2	0.29
			Dinophyceae	220	2,409	42	
			Euglenophyceae	-	0	0	
			Xanthophyceae	-	0	0	
			Total	1,104,847	1,507,572	8	
			Bacillariophyceae	11,810	59,462	33	
			Chlorophyceae	161,183	58,643	23	
			Chrysophyceae	734,807	644,239	3	
			Cryptophyceae	114,553	139,958	5	
Lac de Gras - Slipper Bay	S3	Summer	Cyanobacteria	735,880	579,866	6	0.13
Chipper Day			Dinophyceae	2,955	18,717	36	
			Euglenophyceae	275	-	50	
			Xanthophyceae	275	0	50	
			Total	1,761,738	1,500,885	4	

Table A-1Quality Control Comparison of Phytoplankton Abundance Data in Duplicate
Samples for Lakes in the Jay Project Area, July to September, 2014



Table A-1Quality Control Comparison of Phytoplankton Abundance Data in Duplicate
Samples for Lakes in the Jay Project Area, July to September, 2014

					undance Is/L)		Bray Curtis
Basin	Station	Sampling Period	Major Taxonomic Group	Replicate 1	Replicate 2	RPD (%)	Dissimilarity Index
	Ac-1	Fall	Bacillariophyceae	178,134	174,018	1	0.04
Lac du Sauvage			Chlorophyceae	44,088	43,192	1	
			Chrysophyceae	530,442	567,188	2	
			Cryptophyceae	46,396	58,023	6	
			Cyanobacteria	57,071	67,732	4	
			Dinophyceae	17,690	22,935	6	
			Euglenophyceae	-	-	0	
			Xanthophyceae	-	-	0	
			Total	873,821	933,088	2	

Notes: RPD (%) and total abundance values are rounded to the nearest whole number.

cells/L = number of cells per litre; % = percent; RPD = relative percent difference; Late Spring = July; Summer = August; Fall = September; - = not applicable or zero abundance.



Table A-2 Quality Control Comparison of Zooplankton Abundance Data in Duplicate Samples for Lakes in the Jay Project Area, July to September, 2014

				. Total Abundance (org/L)			Bray Curtis
Basin	Station	Sampling Period	Major Taxonomic Group	Replicate 1	Replicate 2	RPD (%)	Dissimilarity Index
			Cladocera	1	0	26	
			Rotifera	21	19	2	
Lac de Gras – Far-field	FF2-5	Lata Carina	Copepoda - nauplii	2	2	9	0.07
2	гг2-э	Late Spring	Calanoida	1	1	1	0.07
			Cyclopoida	4	3	9	
			Total	28	25	2	
			Cladocera	0	0	21	
			Rotifera	8	8	1	
Lac de Gras - Slipper	05	Lete Carian	Copepoda - nauplii	1	2	7	0.07
Bay	S5	Late Spring	Calanoida	2	2	9	0.07
			Cyclopoida	1	2	3	
			Total	12	14	3	
	fF2-4	Summer	Cladocera	0	0	0	0.08
			Rotifera	46	53	4	
Lac de Gras – Far-field			Copepoda - nauplii	3	5	10	
2			Calanoida	3	3	0	
			Cyclopoida	4	5	3	
			Total	57	67	4	
			Cladocera	0	0	2	
			Rotifera	7	7	0	
Lac de Gras - Slipper	S3	C	Copepoda - nauplii	2	2	1	0.01
Bay	53	Summer	Calanoida	1	1	2	0.01
			Cyclopoida	1	1	1	
			Total	12	12	0	
			Cladocera	3	3	1	
			Rotifera	29	30	1	1
Loo du Couvers	A a 1	Fall	Copepoda - nauplii	14	13	1	0.02
Lac du Sauvage	Ac-1	Fall	Calanoida	0	0	6	0.03
			Cyclopoida	7	6	4	
			Total	53	52	0	

Notes: RPD (%) and total abundance values are rounded to the nearest whole number.

org/L = number of organisms per litre; % = percent; RPD = relative percent difference; Late Spring = July; Summer = August; Fall = September; - = not applicable or zero abundance.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Achnanthidium sp.	6,731	1.11
		Desillerienkusses	Cyclotella sp.	9,424	37.41
		Bacillariophyceae	Tabellaria fenestrata	74	0.17
			Tabellaria flocculosa	110	0.31
			Lagerheimia genevensis	17,502	0.49
		Chlorenhueses	Monoraphidium irregulare	1,417	0.10
		Chlorophyceae	Monoraphidium tortile	1,012	0.12
			Mougeotia sp.	37	1.04
			Dinobryon bavaricum	957	1.85
			Dinobryon bavaricum var. vanhoeffenii	129	0.40
L.L.		Chrysophyceae	Dinobryon divergens	129	0.33
July	Ab-1		Mallomonas sp.	202	0.95
			Ochromonas sp.	352,724	62.37
			Cryptomonas ovata	1,822	15.34
		Cryptophyceae	Cryptomonas sp.	4,048	9.14
			Komma caudata	13,463	1.94
		Cyanobacteria Dinophyceae	Anabaena sp.	19,835	4.64
			Aphanocapsa delicatissima	12,144	0.02
			Chroococcus sp.	5,385	0.18
			Leptolyngbya sp.	37	0.00
			Gyrodinium helveticum	1,012	18.22
			Peridinium sp.	9,542	88.06
			Asterionella formosa	2,846	10.15
			Cyclotella sp.	6,351	3.59
			Fragilaria sp.	205	0.19
		Bacillariophyceae	Nitzschia sp.	19	0.81
			<i>Surirella</i> sp.	205	0.59
			Tabellaria fenestrata	1,637	15.36
			Ankistrodesmus falcatus	1,228	0.35
			Dictyosphaerium pulchellum	3,478	0.43
			Elakatothrix gelatinosa	2,455	0.30
July	Ac-1		Lagerheimia genevensis	1,588	0.11
		Chlorophyceae	Monoraphidium tortile	205	0.05
			Roya sp.	818	1.41
			Teilingia granulata	3,832	11.08
			Xanthidium johnsonii	1,588	1.97
			Chrysocapsa planktonica	4,334	0.94
			Chrysococcus rufescens	1,588	2.81
		Chrysophyceae	Dinobryon bavaricum	2,920	18.41
			Dinobryon bavaricum var. vanhoeffenii	800	2.15



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Dinobryon borgei	1,432	7.35
			Dinobryon divergens	1,693	10.95
			Mallomonas sp.	1,432	5.02
			Ochromonas sp.	439,800	50.70
			Synuropsis janei	1,637	3.48
			Cryptomonas erosa	5,115	23.22
		On intershuiseee	Cryptomonas marssonii	1,588	1.50
		Cryptophyceae	Cryptomonas sp.	3,175	1.81
			Komma caudata	7,939	1.13
		Querra ha staria	Anabaena sp.	21,176	12.81
		Cyanobacteria	Chroococcus sp.	4,763	0.16
			Amphidinium sp.	7,939	5.40
			Gyrodinium helveticum	614	10.03
		Discul	Peridinium inconspicuum	1,588	3.18
		Dinophyceae	Peridinium sp.	2,455	17.50
			Peridinium umbonatum	818	5.75
			Peridinium willei	614	17.40
			Amphora sp.	220	0.31
		Bacillariophyceae	Asterionella formosa	3,300	2.83
			Cyclotella sp.	440	2.71
			Fragilaria sp.	440	0.62
			Navicula sp.	440	0.61
			Nitzschia sp.	220	0.23
			Stauroneis sp.	20	0.86
			Tabellaria fenestrata	1,060	3.57
			Tabellaria flocculosa	520	1.74
			Botryococcus braunii	6,160	0.52
			Dictyosphaerium pulchellum	5,280	0.43
July	Ac-4		Elakatothrix gelatinosa	880	0.05
-			Gonatozygon brebissonii	660	2.55
			Lagerheimia genevensis	11,175	1.10
		Chlorophyceae	Monoraphidium arcuatum	220	0.05
			Monoraphidium komarkovae	3,300	0.70
			Mougeotia sp.	160	2.54
			Roya sp.	1,320	1.05
			Staurodesmus triangularis	220	5.24
			Teilingia granulata	2,040	5.80
			Chrysocapsa planktonica	16,762	2.21
		Chrysophyceae	Dinobryon bavaricum	8,773	17.30
		Chrysophyceae		· · ·	



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Dinobryon divergens	3,640	13.54
			Mallomonas sp.	1,320	8.78
			Ochromonas sp.	365,968	38.97
			Synuropsis janei	440	0.81
			Uroglenopsis americana	1,200	0.11
			Cryptomonas erosa	220	0.60
		Crumtanhuaaaa	Cryptomonas ovata	1,540	13.28
		Cryptophyceae	Cryptomonas sp.	1,397	2.22
			Komma caudata	22,349	3.15
		Querra ha staria	Anabaena sp.	128,553	89.10
		Cyanobacteria	Aphanocapsa delicatissima	39,600	0.13
			Amphidinium sp.	1,397	1.98
		Dinophyceae	Peridinium sp.	5,500	25.98
			Peridinium willei	440	28.53
			Asterionella formosa	2,260	5.97
		Bacillariophyceae	Cyclotella sp.	3,841	0.98
			Fragilaria sp.	880	1.01
			Nitzschia sp.	440	0.48
			Tabellaria fenestrata	980	5.62
			Tabellaria flocculosa	220	0.36
		Chlorophyceae	Euastrum elegans	440	3.28
			Lagerheimia genevensis	17,286	0.32
			Monoraphidium tortile	4,400	0.44
			Roya sp.	2,200	1.28
			Staurastrum sp.	220	1.44
			Staurastrum spp.	220	9.63
			Dinobryon bavaricum	1,500	3.98
July	Ac-7		Dinobryon bavaricum var. vanhoeffenii	1,800	8.50
		Chrysophyceae	Dinobryon divergens	6,300	16.63
			Ochromonas sp.	522,413	85.32
			Synuropsis janei	8,402	30.98
			Cryptomonas ovata	3,300	19.90
		Cryptophyceae	Cryptomonas sp.	19,706	52.53
			Komma caudata	5,762	1.15
			Anabaena sp.	73,040	17.69
		Cyanobacteria	Aphanocapsa delicatissima	26,400	0.07
			Chroococcus sp.	11,524	0.39
			Gyrodinium helveticum	2,640	78.72
		Dinophyceae	Peridinium sp.	3,080	21.56
			Peridinium willei	1,760	113.89



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m ³)
		Xanthophyceae	Ophiocytium parvulum	220	1.21
			Cyclotella sp.	7,529	13.41
		Bacillariophyceae	Navicula sp.	2,510	1.51
			Tabellaria fenestrata	255	0.97
			Euastrum elegans	647	2.56
			Lagerheimia genevensis	35,135	1.38
		Chlorophycoco	Monoraphidium komarkovae	647	0.20
		Chlorophyceae	<i>Mougeotia</i> sp.	78	0.34
			Roya sp.	1,725	1.80
			Staurastrum sp.	216	2.28
			Chrysocapsa planktonica	5,174	1.13
le de c	A -1 4		Chrysocapsella planctonica	5,019	2.35
July	Ad-1		Dinobryon bavaricum	1,098	2.72
		Chrysophyceae	Dinobryon bavaricum var. vanhoeffenii	314	1.32
			Dinobryon divergens	39	0.15
			Ochromonas sp.	680,110	114.78
			Synuropsis janei	3,156	8.50
		Cryptophyceae Cyanobacteria	Cryptomonas ovata	4,096	27.45
			Komma caudata	7,529	1.21
			Anabaena circinalis	1,960	0.38
			Anabaena sp.	19,404	10.58
			Chroococcus sp.	10,039	0.14
		Dinophyceae	Amphidinium sp.	2,510	2.54
			Achnanthidium sp.	18,773	2.12
			Asterionella formosa	1,152	1.76
			Cyclotella sp.	8,046	7.11
		Decilleringhouses	Eunotia sp.	211	3.20
		Bacillariophyceae	Fragilaria sp.	1,056	0.76
			Nitzschia sp.	230	13.98
			Tabellaria fenestrata	845	6.81
			Tabellaria flocculosa	19	0.05
July	Ae-1		Euastrum elegans	211	1.33
			Gloeocystis sp.	2,682	6.90
			Gonatozygon monotaenium	19	0.97
		Oblasarburga	Lagerheimia genevensis	13,410	0.53
		Chlorophyceae	Monoraphidium komarkovae	1,267	0.41
			Roya sp.	1,056	1.04
			Staurastrum sp.	211	0.29
			Staurastrum spp.	211	1.94
		Chrysophyceae	Dinobryon bavaricum	154	0.30



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Dinobryon bavaricum var. vanhoeffenii	58	0.13
			Dinobryon sp.	8,046	2.68
			Mallomonas sp.	422	0.89
			Ochromonas sp.	737,524	71.81
			Uroglenopsis americana	8,640	0.87
			Cryptomonas erosa	1,901	20.74
		Cryptophyceae	Cryptomonas ovata	2,746	25.12
		Стурюрпусеае	Cryptomonas sp.	1,056	1.90
			Komma caudata	10,728	1.82
		Cyanobacteria	Anabaena sp.	125,523	84.19
			Amphidinium sp.	5,364	4.42
		Dinophyceae	Gyrodinium helveticum	211	6.50
			Peridinium sp.	3,802	22.76
		Xanthophyceae	Ophiocytium parvulum	211	1.87
			Achnanthidium sp.	3,361	0.24
			Asterionella formosa	8,104	15.77
			Cocconeis sp.	193	2.99
		Bacillariophyceae	Cyclotella sp.	10,083	5.05
		Dacinanophyceae	Navicula sp.	385	0.50
			Navicymbula sp.	193	3.43
			Synedra sp.	18	0.05
			Tabellaria fenestrata	3,010	14.15
			Botryococcus braunii	11,550	0.60
			Closteriopsis acicularis	3,658	0.50
			Crucigenia tetrapedia	10,083	6.53
			Crucigeniella irregularis	5,390	3.93
August	A a 1		Crucigeniella rectangularis	3,080	2.25
August	Aa-1	Chlorophyceae	Dictyosphaerium pulchellum	20,983	2.24
			Elakatothrix sp.	11,238	0.25
			Gloeocystis sp.	3,361	1.92
			Quadrigula closterioides	4,620	0.19
			Scenedesmus sp.	6,722	0.56
			Spondylosium sp.	1,925	6.48
			Chrysocapsa planktonica	26,889	7.21
			Chrysocapsella planctonica	46,197	35.46
			Dinobryon bavaricum	473	1.04
		Chrysophyceae	Dinobryon divergens	35,230	130.93
			Ochromonas spp.	648,694	177.55
			Synuropsis janei	7,019	8.57
			Uroglenopsis americana	206,027	16.09



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Chroomonas sp.	3,746	6.24
		Cryptophyceae	Cryptomonas ovata	11,816	41.45
			Komma caudata	275,611	22.02
			Anabaena affinis	525	0.09
			Anabaena circinalis	29,663	9.17
			Anabaena sp.	5,075	1.24
		Cyanobacteria	Anabaena spp.	7,508	1.40
		Cyanobacteria	Chroococcus limneticus	3,080	0.55
			Leptolyngbya sp.	893	0.00
			Merismopedia sp.	121,000	0.29
			Woronichinia naegeliana	14,438	0.06
		Euglenophyceae	Trachelomonas sp.	13,444	17.90
			Asterionella formosa	6,110	15.04
			Cyclotella sp.	2,881	1.01
		Bacillariophyceae	Synedra sp.	431	1.49
			Tabellaria fenestrata	3,919	17.58
			Tabellaria flocculosa	263	0.95
			Closteriopsis acicularis	206	0.02
			Elakatothrix gelatinosa	975	0.38
			Elakatothrix sp.	17,698	0.40
			Microspora sp.	656	0.62
			Monoraphidium komarkovae	21,640	0.82
			Monoraphidium minutum	2,881	0.01
		Chlorophyceae	<i>Oocystis</i> sp.	2,881	0.54
			Quadrigula closterioides	3,300	0.13
A			Roya sp.	7,396	20.70
August	Ab-1		Schroederia sp.	1,444	0.42
			Spondylosium sp.	581	2.25
			Staurastrum sp.	206	3.82
			Tetraselmis sp.	2,881	0.45
			Bitrichia chodatii	206	0.05
			Chrysocapsella planctonica	63,381	15.02
			Chrysococcus rufescens	17,286	2.66
			Dinobryon bavaricum	1,519	4.08
		Chrysophyceae	Dinobryon divergens	5,644	11.24
			Mallomonas insignis	413	0.27
			Mallomonas sp.	1,650	2.73
			Ochromonas spp.	645,333	132.83
			Uroglenopsis americana	58,875	5.33
		Cryptophyceae	Cryptomonas ovata	6,587	14.31



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Komma caudata	115,238	9.59
			Anabaena sp.	12,638	3.87
			Aphanocapsa sp.	158,452	0.23
		Cyanobacteria	Leptolyngbya sp.	8,081	0.06
			Merismopedia sp.	34,571	0.14
			Pseudanabaena catenata	750	0.02
		Euglenophyceae	Trachelomonas sp.	2,881	2.01
			Achnanthidium sp.	5,762	0.93
			Asterionella formosa	12,921	31.82
			Cocconeis sp.	1,921	4.98
			Cyclotella sp.	3,841	0.87
		Bacillariophyceae	Cymbella sp.	1,921	1.01
			Navicula sp.	3,841	2.06
			Synedra sp.	240	0.33
			Tabellaria fenestrata	9,618	62.25
			Tabellaria flocculosa	160	1.84
			Botryococcus braunii	36,080	0.51
			Chlamydomonas sp.	5,762	3.07
			Chlorella ellipsoidea	13,444	4.28
			Crucigenia tetrapedia	3,841	3.73
			Dictyosphaerium pulchellum	9,020	1.36
			Elakatothrix sp.	21,846	1.03
			Gloeocystis sp.	1,921	0.73
August	Ac-1		Kirchneriella obesa	600	0.01
-			Monoraphidium arcuatum	220	0.02
			Monoraphidium komarkovae	3,300	0.23
		Chlorophyceae	Mougeotia sp.	160	0.40
			Oocystis sp.	880	1.79
			Quadrigula closterioides	3,520	0.18
			Roya sp.	9,561	8.99
			Schroederia sp.	440	0.02
			Sphaerocystis sp.	12,320	0.62
			Spondylosium sp.	2,101	9.92
			Staurodesmus triangularis var. inflatus	440	7.71
			Teilingia excavata	440	0.13
			Tetraedron triangulare	1,921	1.38
			Tetrastrum triangulare	7,683	0.26
			Chrysocapsella planctonica	59,379	24.76
		1			
		Chrysophyceae	Chrysococcus rufescens	21,127	4.15



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Dinobryon divergens	2,980	6.66
			Mallomonas insignis	3,241	3.53
			Mallomonas sp.	13,664	39.59
			Ochromonas spp.	205,508	50.25
			Uroglenopsis americana	23,800	2.75
			Chroomonas sp.	3,841	4.07
		Cruptophycoco	Cryptomonas ovata	10,162	24.05
		Cryptophyceae	Komma caudata	197,825	13.45
			Rhodomonas sp.	1,921	3.62
			Anabaena sp.	9,620	2.11
		Cyanobacteria	Aphanocapsa delicatissima	12,540	0.04
			Leptolyngbya sp.	28,786	0.38
			Amphidinium sp.	7,683	7.55
			<i>Gymnodinium</i> sp.	2,361	53.62
		Dinophyceae	Gyrodinium helveticum	220	4.05
			Peridinium inconspicuum	1,921	1.06
			Peridinium umbonatum	5,762	5.15
		Evolution	<i>Euglena</i> sp.	1,100	38.74
		Euglenophyceae	Trachelomonas sp.	1,921	2.38
		Xanthophyceae	Ophiocytium parvulum	1,921	0.68
			Achnanthidium sp.	1,646	0.21
			Asterionella formosa	9,850	18.94
			Pleurosigma sp.	23	14.24
		Bacillariophyceae	Synedra sp.	293	0.65
			Tabellaria fenestrata	16,286	74.23
			Tabellaria flocculosa	180	1.15
			Botryococcus braunii	4,950	0.03
			Chlorella ellipsoidea	23,048	1.31
			Cosmarium sp.	1,894	4.50
A	0 - 1		Dictyosphaerium pulchellum	4,208	0.59
August	Ac-4		Elakatothrix sp.	8,070	0.33
			Monoraphidium komarkovae	8,168	0.38
		Chlorophyceae	Oocystis submarina	495	0.08
			Quadrigula closterioides	2,475	0.48
			<i>Roya</i> sp.	5,220	19.42
			Schroederia sp.	1,238	0.26
			Sphaerocystis sp.	13,170	0.19
			Staurastrum sp.	248	4.80
		Ohmenshiring	Bitrichia chodatii	495	0.11
		Chrysophyceae	Chrysocapsa planktonica	8,415	0.12



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Chrysocapsella planctonica	69,143	12.58
			Dinobryon bavaricum	3,150	11.89
			Dinobryon divergens	5,400	10.83
			Mallomonas insignis	990	0.78
			Mallomonas sp.	14,816	7.95
			Ochromonas spp.	304,558	43.89
			Uroglenopsis americana	50,400	9.36
		Cryptophyceae	Komma caudata	83,959	4.83
			Anabaena circinalis	13,500	2.42
			Anabaena sp.	12,353	3.40
		Cyanobacteria	Aphanocapsa delicatissima	12,375	0.05
			Aphanocapsa sp.	24,750	0.10
			Leptolyngbya sp.	2,363	0.00
			Amphidinium sp.	3,293	4.32
		Dinophyceae	<i>Gymnodinium</i> sp.	495	5.51
			Peridinium umbonatum	13,170	21.89
		Euglenophyceae	Trachelomonas sp.	1,646	0.95
			Asterionella formosa	24,090	56.13
		Bacillariophyceae	Cyclotella sp.	10,109	9.49
			Cymbella sp.	25	0.20
			Eunotia sp.	300	9.69
			Fragilaria sp.	275	0.17
			Navicula sp.	275	1.08
			Synedra sp.	25	0.03
			Tabellaria fenestrata	5,325	38.82
			Chlamydomonas sp.	18,195	23.92
			Chlorella ellipsoidea	24,261	4.66
			Elakatothrix gelatinosa	3,300	0.05
August	Ac-7		Elakatothrix sp.	16,174	0.43
-			Lagerheimia genevensis	4,043	0.03
			Monoraphidium komarkovae	5,500	0.27
		Chlorophyceae	Mougeotia sp.	75	0.39
			Quadrigula closterioides	5,597	1.03
			Roya sp.	11,092	8.27
			Schroederia sp.	275	0.06
			Sphaerocystis sp.	62,271	3.23
			Spondylosium sp.	5,222	41.94
			Teilingia excavata	525	1.00
		Chrysophyceae	Chrysocapsa planktonica	6,793	1.79
		Cryptophyceae	Chroomonas sp.	2,022	1.37



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Cryptomonas ovata	8,540	25.47
			Komma caudata	157,694	11.23
			Anabaena affinis	1,975	0.07
			Anabaena sp.	20,025	3.17
		Cyanobacteria	Aphanocapsa delicatissima	25,625	0.09
			Leptolyngbya sp.	475	0.02
			Phormidium sp.	2,375	0.27
			Amphidinium sp.	4,043	5.81
		Dinanhyaaaa	Gyrodinium helveticum	275	4.93
		Dinophyceae	Peridinium umbonatum	22,239	55.58
			Peridinium willei	275	7.41
		Euglenophyceae	Trachelomonas sp.	2,022	1.08
			Asterionella formosa	7,155	14.71
			Aulacoseira sp.	1,841	1.60
		Bacillariophyceae	Cyclotella sp.	1,819	10.43
			Cymbella sp.	248	0.10
			Tabellaria fenestrata	4,725	24.25
		Chlorophyceae	Chlamydomonas sp.	15,714	4.29
			Chlorella ellipsoidea	25,143	4.28
			Cosmarium sp.	1,571	0.07
			Dictyosphaerium pulchellum	5,123	1.42
			Elakatothrix gelatinosa	1,868	0.23
			<i>Roya</i> sp.	11,298	44.90
			Sphaerocystis sp.	11,656	2.84
			Spondylosium sp.	1,980	17.36
			Bitrichia chodatii	495	0.12
August	Ad-1		Chrysocapsa planktonica	9,429	0.39
			Chrysocapsella planctonica	53,837	25.87
			Dinobryon bavaricum	5,869	71.43
			Dinobryon divergens	4,253	19.70
		Chrysophyceae	Mallomonas insignis	4,628	10.71
			Mallomonas multiunca	248	0.26
			Mallomonas sp.	28,533	77.55
			Ochromonas spp.	234,143	71.87
			Uroglenopsis americana	3,600	0.39
			Cryptomonas ovata	743	2.04
		Cryptophyceae	Cryptomonas sp.	1,571	0.44
			Komma caudata	91,143	7.37
		Cyanobacteria	Komma caudata Anabaena sp.	91,143 19,103	7.37



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Leptolyngbya sp.	25,245	0.09
			<i>Merismopedia</i> sp.	18,857	0.08
			Woronichinia naegeliana	29,700	0.17
		Dinophyceae	<i>Gymnodinium</i> sp.	2,066	28.17
		Diriophyceae	Peridinium umbonatum	9,429	12.36
		Euglenophyceae	Trachelomonas sp.	3,143	1.41
			Asterionella formosa	8,255	6.48
			<i>Cyclotella</i> sp.	7,683	6.64
		Bacillariophyceae	<i>Eunotia</i> sp.	20	0.47
			Synedra sp.	300	0.33
			Tabellaria fenestrata	8,551	40.48
			Botryococcus braunii	21,800	1.11
			Botryococcus protuberans	17,600	0.61
			Chlamydomonas sp.	30,730	11.82
			Chlorella ellipsoidea	23,048	4.08
			Cosmarium margaritatum	220	0.54
		Oblasshussa	Cosmarium sp.	880	1.00
			Dictyosphaerium pulchellum	20,020	4.50
			Elakatothrix gelatinosa	760	0.13
			Elakatothrix sp.	15,383	0.49
			Gloeocystis sp.	2,640	3.26
			Lagerheimia genevensis	3,841	0.15
August	A o 1	Chlorophyceae	Monoraphidium irregulare	4,061	0.05
August	Ae-1		Monoraphidium komarkovae	5,280	0.28
			Quadrigula closterioides	3,040	0.66
			<i>Quadrigula</i> sp.	1,840	0.72
			<i>Roya</i> sp.	6,160	4.86
			Sphaerocystis sp.	49,193	1.39
			Spondylosium sp.	880	2.41
			Staurastrum sp.	220	5.38
			Staurodesmus triangularis	220	0.82
			Staurodesmus triangularis var. inflatus	660	8.43
		Teilingia excavata	800	0.62	
			Chrysocapsella planctonica	80,024	35.67
			Dinobryon bavaricum	5,395	12.88
			Dinobryon divergens	5,020	17.71
		Chrysophyceae	Mallomonas insignis	5,381	8.21
			Mallomonas multiunca	3,841	2.44
			Mallomonas sp.	4,281	6.86
			Ochromonas spp.	902,698	180.55



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Uroglenopsis americana	100,806	11.68
		Cruptophycoco	Cryptomonas ovata	5,601	15.28
		Cryptophyceae	Komma caudata	103,714	7.75
			Anabaena circinalis	18,400	3.29
			Anabaena sp.	35,812	9.29
		Cyanobacteria	Aphanocapsa delicatissima	44,000	0.10
			Aphanocapsa elachista	2,000	0.01
			Aphanocapsa sp.	88,349	0.08
		Dipanhyaaaa	<i>Gymnodinium</i> sp.	220	2.52
		Dinophyceae	Peridinium umbonatum	4,281	25.15
			Asterionella formosa	91,637	232.32
		Decillerienhycese	<i>Cyclotella</i> sp.	2,235	0.57
		Bacillariophyceae	Nitzschia sp.	32	0.28
			Tabellaria fenestrata	41,947	182.70
		Chlorophyceae	Dictyosphaerium pulchellum	25,344	4.10
	Aa-1		Gonium pectorale	768	0.34
			Monoraphidium tortile	1,232	0.15
			<i>Roya</i> sp.	5,808	12.14
Contombor			Spondylosium planum	2,656	2.76
September		Chrysophyceae	Dinobryon divergens	400	1.48
			Mallomonas sp.	25,816	254.68
			Ochromonas sp.	587,784	242.59
		Ormitechusees	Cryptomonas ovata	11,616	84.82
		Cryptophyceae	Komma caudata	31,289	6.23
		Cyanobacteria	Anabaena sp.	288	0.06
			Chroococcus sp.	40,229	0.57
		Dinanhyanan	Amphidinium sp.	4,470	3.28
		Dinophyceae	Peridinium sp.	23,232	485.69
			Asterionella formosa	57,987	159.41
			Eunotia sp.	16	0.99
		Bacillariophyceae	Nitzschia sp.	16	0.13
			Tabellaria fenestrata	23,936	186.63
			Tabellaria flocculosa	128	0.27
Contombor	Ah 1		Botryococcus braunii	9,152	0.93
September	Ab-1		Dictyosphaerium pulchellum	39,424	5.89
			Elakatothrix gelatinosa	2,112	0.13
		Chlorophyceae	Monoraphidium tortile	1,232	0.16
			<i>Oocystis</i> sp.	11,175	3.73
			Roya sp.	880	0.75
			Spondylosium planum	832	0.73



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Staurodesmus triangularis	528	37.13
			Chrysocapsella planctonica	4,470	0.76
			Dinobryon bavaricum	720	1.76
		Ohmanaharan	Dinobryon divergens	80	0.34
		Chrysophyceae	Mallomonas sp.	1,117	0.94
			Ochromonas sp.	271,543	106.54
			Uroglenopsis americana	9,600	1.45
			Cryptomonas ovata	528	3.79
		Cryptophyceae	Komma caudata	50,286	8.36
			Rhodomonas sp.	1,117	2.50
		Cyanobacteria	Anabaena sp.	2,320	0.78
		Disastas	Amphidinium sp.	2,235	1.87
		Dinophyceae	Peridinium sp.	11,063	177.64
			Asterionella formosa	52,125	142.72
		Bacillariophyceae	Cyclotella sp.	13,829	36.19
			Fragilaria sp.	990	1.14
			Tabellaria fenestrata	115,306	214.01
		Chlorophyceae	Crucigeniella crucifera	6,914	28.32
			Dictyosphaerium pulchellum	35,640	4.43
			<i>Mougeotia</i> sp.	198	5.04
			Roya sp.	1,386	1.38
			Spondylosium planum	648	0.51
			Staurastrum sp.	198	3.49
0	A = 4(a)		Dinobryon bavaricum	342	0.72
September	Ac-1 ^(a)		Dinobryon divergens	8,640	8.68
		Chrysophyceae	Mallomonas sp.	14,027	298.24
			Ochromonas sp.	444,243	123.80
			Uroglenopsis americana	16,200	1.83
			Cryptomonas sp.	1,927	16.88
		Cryptophyceae	Komma caudata	32,843	5.07
			Anabaena circinalis	5,220	1.76
		Cyanobacteria	Anabaena sp.	15,228	9.79
			Chroococcus sp.	20,743	0.47
			Amphidinium sp.	5,186	6.26
		Dinophyceae	Peridinium sp.	7,260	187.63
			Achnanthidium sp.	10,243	1.16
			Asterionella formosa	59,180	198.89
September	Ac-1 ^(b)	Bacillariophyceae	Cyclotella sp.	10,243	3.22
-			Fragilaria sp.	220	0.45
			Nitzschia sp.	20	0.07



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Tabellaria fenestrata	94,111	319.70
			Cosmarium sp.	2,049	2.08
			Dictyosphaerium pulchellum	29,480	3.36
		Oblemation	Lagerheimia genevensis	6,146	0.13
		Chlorophyceae	Mougeotia sp.	100	0.41
			<i>Oocystis</i> sp.	4,317	16.13
			<i>Roya</i> sp.	1,100	1.57
			Chrysocapsella planctonica	10,243	1.93
			Dinobryon bavaricum	23,100	39.36
			Dinobryon divergens	25,474	30.77
		Chrysophyceae	Mallomonas sp.	4,689	85.67
			Ochromonas sp.	491,683	130.31
			Uroglenopsis americana	12,000	3.86
			Cryptomonas sp.	660	1.29
		Cryptophyceae	Komma caudata	51,217	8.09
			Rhodomonas sp.	6,146	11.35
		Querration text	Anabaena sp.	55,440	32.38
		Cyanobacteria	Chroococcus sp.	12,292	0.17
			Gyrodinium helveticum	220	4.61
		Dinophyceae	Peridinium sp.	14,520	118.67
			Peridinium umbonatum	8,195	16.41
			Achnanthidium sp.	6,470	0.85
		Bacillariophyceae	Asterionella formosa	10,531	25.49
			Cyclotella sp.	4,852	3.22
			Tabellaria fenestrata	12,270	36.21
			Gonatozygon brebissonii	20	0.29
			Lagerheimia genevensis	1,617	0.05
			Monoraphidium tortile	3,235	0.26
			<i>Oocystis</i> sp.	3,235	0.85
		Chlorophyceae	<i>Roya</i> sp.	1,100	1.08
September	Ac-4		Staurastrum arachne	20	0.31
			Teilingia granulata	1,020	1.62
			Tetraedron incus	3,235	0.16
			Tetrastrum triangulare	6,470	0.42
			Dinobryon divergens	40,333	84.51
		Ohman h	Mallomonas sp.	8,087	5.14
		Chrysophyceae	Ochromonas sp.	380,084	55.21
			Uroglenopsis americana	8,000	1.54
		Ormitant	Cryptomonas ovata	3,300	33.67
		Cryptophyceae	Cryptomonas sp.	1,617	0.66



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Komma caudata	71,165	10.93
		Quesekesterie	Anabaena sp.	6,520	3.79
		Cyanobacteria	Leptolyngbya sp.	39,600	0.58
		Disartura	Peridinium sp.	4,400	37.85
		Dinophyceae	Peridinium umbonatum	1,617	3.43
			Achnanthidium sp.	1,383	0.24
		Destillenten have an	Asterionella formosa	2,970	2.80
		Bacillariophyceae	Cyclotella sp.	8,102	39.70
			Nitzschia sp.	18	0.99
			Cosmarium sp.	198	1.37
			Dictyosphaerium pulchellum	23,958	1.84
		Chlorophyceae	Monoraphidium komarkovae	198	0.04
			Monoraphidium tortile	1,782	0.24
			<i>Roya</i> sp.	3,564	3.94
			Dinobryon bavaricum	18,036	18.15
September	Ac-7	Chrysophyceae	Dinobryon divergens	24,750	30.47
			Mallomonas sp.	3,759	44.35
			Ochromonas sp.	356,777	71.22
			Uroglenopsis americana	10,800	2.68
		Cryptophyceae	Cryptomonas sp.	2,769	8.68
			Komma caudata	35,954	4.93
		Cyanobacteria	Anabaena sp.	30,576	21.08
			Chroococcus sp.	16,594	0.37
			Amphidinium sp.	1,383	2.27
		Dinophyceae	Peridinium sp.	4,950	37.14
		Euglenophyceae	Trachelomonas sp.	1,383	1.77
			Asterionella formosa	27,874	78.89
			Cyclotella sp.	4,780	2.27
		Bacillariophyceae	Tabellaria fenestrata	6,300	26.05
			Tabellaria flocculosa	140	0.26
			Monoraphidium komarkovae	1,848	0.46
			Monoraphidium tortile	770	0.10
			<i>Mougeotia</i> sp.	350	0.59
September	Ad-1	Chlorophyceae	Roya sp.	462	0.31
			Spondylosium planum	910	0.78
			Teilingia excavata	910	0.27
			Dinobryon bavaricum	294	0.78
			Dinobryon divergens	52,360	55.88
		Chrysophyceae	Mallomonas sp.	3,585	8.65
			Ochromonas sp.	310,716	66.43



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Uroglenopsis americana	44,800	5.21
		Crimterburges	Komma caudata	38,242	5.63
		Cryptophyceae	Rhodomonas sp.	1,195	1.84
			Anabaena circinalis	5,600	1.98
		Cuanabastaria	Anabaena sp.	7,868	5.07
		Cyanobacteria	Leptolyngbya sp.	1,260	0.02
			Oscillatoria sp.	644	0.15
		Disarburgas	Peridinium sp.	1,694	11.91
		Dinophyceae	Peridinium willei	154	10.28
			Asterionella formosa	28,512	75.18
			Cyclotella sp.	11,645	7.73
		Bacillariophyceae	Fragilaria sp.	198	0.24
			Tabellaria fenestrata	14,520	45.98
			Tabellaria flocculosa	162	0.19
		Chlorophyceae	Dictyosphaerium pulchellum	21,384	1.76
			Monoraphidium tortile	1,456	0.12
			<i>Roya</i> sp.	990	1.57
			Spondylosium planum	198	0.17
			Staurastrum sp.	396	5.45
			Tetraedron triangulare	2,911	4.45
			Zygnema sp.	108	0.57
0 / /			Dinobryon bavaricum	1,296	2.61
September	Ae-1		Dinobryon divergens	105,461	155.95
		Chrysophyceae	Mallomonas sp.	792	8.84
			Ochromonas sp.	391,567	80.36
			Uroglenopsis americana	32,400	3.38
			Cryptomonas ovata	2,970	27.12
			Cryptomonas sp.	9,060	27.51
		Cryptophyceae	Komma caudata	20,379	2.13
			Rhodomonas sp.	1,456	3.95
		Querral i i	Anabaena circinalis	18,612	5.81
		Cyanobacteria	Anabaena sp.	3,600	0.99
			Peridinium sp.	21,780	241.87
		Dinophyceae	Peridinium willei	594	42.01
		Euglenophyceae	<i>Euglena</i> sp.	198	1.11

Table A-3	Phytoplankton Abundance and Biomass in Lac du Sauvage, 2014
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Notes: Samples were analyzed by EcoAnalysts, Inc. cells/L = number of cells per litre; mg/m³ = milligrams per cubic metre; Late Spring = July; Summer = August; Fall = September; sp. a single species; spp. = multiple species.
a) Replicate 1.
b) Replicate 2.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Achnanthidium sp.	6,914	2.0
		Desillerienhysees	Asterionella formosa	19,048	23.0
		Bacillariophyceae	Cyclotella sp.	13,829	23.7
			Tabellaria fenestrata	4,410	24.4
			Botryococcus braunii	52,470	2.8
			Chlorella ellipsoidea	27,657	10.2
			Coelastrum sp.	55,314	3.6
			Cosmarium sp.	743	2.0
			Crucigenia tetrapedia	13,829	10.5
			Crucigeniella irregularis	990	4.9
			Dictyosphaerium ehrenbergianum	4,703	0.8
			Dictyosphaerium pulchellum	9,158	0.6
		Chlorophyceae	Elakatothrix gelatinosa	3,353	0.4
			Monoraphidium arcuatum	248	0.0
			<i>Mougeotia</i> sp.	68	0.8
			Quadrigula closterioides	3,960	0.2
			<i>Roya</i> sp.	743	0.5
			Sphaerocystis sp.	15,593	0.8
			Spondylosium sp.	990	0.9
August	Af-1		Staurastrum sp.	248	3.3
			Xanthidium sp.	6,914	0.2
			Dinobryon bavaricum	2,520	7.1
			Dinobryon divergens	163,350	372.9
			Mallomonas insignis	990	0.7
		Chrysophyceae	Ochromonas spp.	1,770,057	544.2
			Synuropsis janei	1,238	1.4
			Uroglenopsis americana	637,843	53.3
			Chroomonas sp.	6,914	1.8
			Cryptomonas erosa	743	0.3
		Cryptophyceae	Cryptomonas ovata	5,445	28.3
			Komma caudata	255,829	22.0
			Anabaena circinalis	51,705	15.6
			Anabaena sp.	21,668	3.9
		Cyanobacteria	Merismopedia sp.	82,971	0.2
			Woronichinia naegeliana	19,800	0.1
			Gymnodinium sp.	495	5.7
		Dinophyceae	Peridinium sp.	4,253	33.1
			Peridinium umbonatum	20,264	119.4
		Euglenophyceae	<i>Euglena</i> sp.	248	1.7

Table A-4 Phytoplankton Abundance and Biomass in Duchess Lake, 2014



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Asterionella formosa	8,604	13.1
		Bacillariophyceae	Cyclotella sp.	27,043	28.7
			Tabellaria fenestrata	128	0.5
			Cosmarium sp.	176	2.7
			Dictyosphaerium pulchellum	12,672	1.4
		Chlorophyceae	Micrasterias fimbriata	16	51.3
			<i>Oocystis</i> sp.	31,959	15.0
			<i>Roya</i> sp.	12,227	11.3
		Chrysophyceae	Dinobryon bavaricum	11,616	20.6
September	Af-1		Dinobryon divergens	119,680	211.1
			Mallomonas sp.	4,917	9.6
			Ochromonas sp.	543,309	89.5
			Uroglenopsis americana	160	0.0
			Cryptomonas ovata	1,232	11.3
		Cryptophyceae	Cryptomonas sp.	2,288	5.7
			Komma caudata	135,213	19.9
		Cyanobacteria	Chroococcus sp.	39,335	0.6
		Disashyasaa	Peridinium sp.	1,936	14.3
		Dinophyceae	Peridinium umbonatum	2,458	5.5

 Table A-4
 Phytoplankton Abundance and Biomass in Duchess Lake, 2014

Notes: Samples were analyzed by EcoAnalysts, Inc.

cells/L = number of cells per litre; mg/m³ = milligrams per cubic metre; Late Spring = July; Summer = August; Fall = September; sp. = a single species; spp. = multiple species.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Asterionella formosa	6,434	19.3
		Decillerianhyasaa	Fragilaria sp.	197	1.1
		Bacillariophyceae	Tabellaria fenestrata	98	0.3
			Tabellaria flocculosa	328	0.6
			Ankistrodesmus falcatus	2,165	0.5
			Elakatothrix gelatinosa	361	0.0
		Chlorophysoco	Lagerheimia genevensis	5,040	0.1
		Chlorophyceae	Monoraphidium irregulare	722	0.1
			Monoraphidium komarkovae	18,318	3.0
			<i>Roya</i> sp.	1,984	2.6
July	S2		Dinobryon bavaricum	2,624	4.6
		Chrysophyceae	Dinobryon divergens	2,460	7.2
			Ochromonas sp.	690,445	110.1
	-		Cryptomonas ovata	5,051	50.9
			Cryptomonas sp.	19,629	70.4
		Cryptophyceae	Komma caudata	35,278	5.7
			Rhodomonas sp.	2,520	6.4
		Cyanobacteria	Anabaena sp.	459	0.2
			Leptolyngbya sp.	3,969	0.1
		Dinophyceae	Amphidinium sp.	7,560	5.5
			Peridinium sp.	2,886	24.5
			Achnanthidium sp.	5,106	1.0
			Asterionella formosa	792	1.1
		Bacillariophyceae	Cyclotella sp.	2,177	9.1
			Fragilaria sp.	29	0.1
			Ankistrodesmus falcatus	5,702	1.6
			Cosmarium sp.	5,106	1.2
		Chlorophyceae	Monoraphidium komarkovae	10,454	3.4
July	S3		Spirogyra sp.	43	0.3
			Tetraedron incus	6,808	0.4
			Dinobryon bavaricum	3,312	8.5
		Chrysophyceae	Dinobryon divergens	52,272	275.3
			Ochromonas sp.	459,534	83.7
			Cryptomonas sp.	1,109	3.4
		Cryptophyceae	Komma caudata	32,338	5.3
		Dinophyceae	Peridinium sp.	792	11.3
			Achnanthidium sp.	13,521	2.5
July	S5 ^(a)	Bacillariophyceae	Cyclotella sp.	6,763	21.3
-			Nitzschia sp.	581	0.6



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
	Station		Ankistrodesmus falcatus	1,162	0.2
			Monoraphidium arcuatum	194	0.0
		Chlorophysoco	Monoraphidium komarkovae	12,778	2.7
		Chlorophyceae	Monoraphidium tortile	581	0.1
			Roya sp.	194	0.6
			Staurodesmus sp.	774	32.7
			Chrysocapsella planctonica	11,590	1.7
			Dinobryon bavaricum	2,640	8.4
		Chrysophyceae	Dinobryon divergens	67,115	400.1
			Ochromonas sp.	531,193	113.7
			Cryptomonas ovata	968	7.8
		Cryptophyceae	Cryptomonas sp.	968	2.8
			Komma caudata	17,384	3.3
		Cyanobacteria	Leptolyngbya sp.	2,816	0.1
		Dinophyceae	Peridinium sp.	968	9.6
		Bacillariophyceae	Achnanthidium sp.	10,057	1.1
			Asterionella formosa	1,386	1.4
			Cyclotella sp.	6,939	39.3
			Fragilaria sp.	792	1.1
			Nitzschia sp.	18	0.1
	-		Ankistrodesmus falcatus	1,584	0.5
			Monoraphidium irregulare	990	0.1
			Monoraphidium komarkovae	21,780	4.9
		Chlorophyceae	Monoraphidium tortile	792	0.1
			Mougeotia sp.	126	2.4
	o r (b)		Spirogyra sp.	54	13.3
July	S5 ^(b)		Dinobryon bavaricum	576	1.2
			Dinobryon divergens	108,900	375.7
		Chrysophyceae	Mallomonas sp.	396	2.8
			Ochromonas sp.	352,000	83.1
			Cryptomonas erosa	2,574	18.8
		Cryptophyceae	Cryptomonas sp.	3,771	4.3
			Komma caudata	6,286	1.2
		Cyanobacteria	Leptolyngbya sp.	10,800	0.2
			Amphidinium sp.	1,257	0.6
		Dinophyceae	Peridinium sp.	2,574	19.2
		Xanthophyceae	Ophiocytium cochleare	594	4.3



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
		Bacillariophyceae	Asterionella formosa	189	0.2
		Bacillanophyceae	Cyclotella sp.	2,370	13.0
			Monoraphidium komarkovae	5,108	1.2
		Chlorophyceae	Monoraphidium tortile	568	0.1
			Staurodesmus sp.	189	6.8
			Chrysocapsella planctonica	4,204	1.8
			Dinobryon bavaricum	52	0.2
lub.	S6	Chrysophyceae	Dinobryon borgei	189	0.4
July	30		Dinobryon divergens	21,001	218.8
			Ochromonas sp.	164,574	29.5
			Cryptomonas ovata	378	3.3
		Cryptophyceae	Cryptomonas sp.	1,357	5.8
			Komma caudata	8,409	1.6
		Diasakusasa	Amphidinium sp.	601	0.6
		Dinophyceae	Gyrodinium helveticum	189	13.8
		Xanthophyceae	Ophiocytium cochleare	189	0.6
		Bacillariophyceae	Achnanthidium sp.	2,469	0.2
			Asterionella formosa	7,202	7.6
			Cyclotella sp.	42,970	428.5
			Fragilaria sp.	124	0.1
			Tabellaria flocculosa	608	8.2
			Ankistrodesmus falcatus	5,693	0.8
			Chlorella ellipsoidea	2,469	0.4
			Cladophora sp.	225	1.5
			Coelastrum sp.	22,224	0.7
			Crucigenia quadrata	7,408	2.5
			Elakatothrix gelatinosa	1,733	0.0
August	S2		Elakatothrix sp.	12,347	0.1
			Monoraphidium irregulare	2,469	0.1
		Chlorophysess	Monoraphidium komarkovae	4,331	0.3
		Chlorophyceae	Monoraphidium minutum	2,469	0.0
			Mougeotia sp.	225	0.9
			Oocystis submarina	9,878	0.7
			Quadrigula sp.	5,310	1.2
			Roya sp.	1,114	1.2
			Scenedesmus sp.	9,878	0.5
			Sphaerocystis sp.	36,608	39.4
			Spondylosium sp.	371	0.7
			Tetraedron incus	2,469	0.1



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
	otation		Chrysocapsella planctonica	16,425	7.0
			Dinobryon bavaricum	12,925	32.7
		Chrysophysopp	Dinobryon divergens	4,208	18.1
		Chrysophyceae	Mallomonas insignis	619	0.5
			Mallomonas sp.	2,469	2.1
			Ochromonas spp.	543,265	290.7
			Cryptomonas ovata	3,336	6.1
		Cryptophyceae	Cryptomonas sp.	6,795	15.3
			Komma caudata	113,592	7.0
			Aphanocapsa delicatissima	31,725	0.1
			Aphanocapsa holsatica	13,725	0.1
		Cuanabastaria	Aphanocapsa sp.	84,263	0.4
		Cyanobacteria	Aphanothece clathrata	59,175	0.3
			Leptolyngbya sp.	214,376	3.2
			Woronichinia naegeliana	3,094	0.0
			<i>Gymnodinium</i> sp.	4,939	6.9
		Dinophyceae	Gyrodinium helveticum	248	5.6
			Peridinium umbonatum	2,469	6.3
		Bacillariophyceae	Asterionella formosa	2,325	2.2
			Cyclotella sp.	7,560	77.1
			Synedra sp.	1,375	1.3
			Tabellaria flocculosa	550	4.4
			Ankistrodesmus falcatus	10,175	0.9
			Botryococcus protuberans	17,200	1.4
			Chlamydomonas sp.	2,955	0.8
			Cosmarium margaritatum	275	0.6
			Crucigenia quadrata	17,729	2.0
			Dictyosphaerium pulchellum	5,910	0.4
August	S3 ^(a)		Elakatothrix gelatinosa	550	0.0
			Elakatothrix sp.	11,819	0.1
		Chlorophyceae	Gloeocystis sp.	2,955	2.7
			Gonium sp.	4,400	0.2
			Monoraphidium arcuatum	2,955	0.1
			Monoraphidium irregulare	3,230	0.5
			Monoraphidium komarkovae	14,300	0.9
			Oocystis solitaria	2,955	1.9
			Oocystis sp.	17,729	7.8
			Quadrigula sp.	5,910	0.9
			<i>Roya</i> sp.	1,100	0.6



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Sphaerocystis sp.	32,503	0.8
			Spondylosium sp.	625	1.0
			Tetraedron incus	5,910	0.2
			Bitrichia chodatii	550	0.3
			Dinobryon bavaricum	4,625	19.7
			Dinobryon divergens	3,500	12.8
		Chrysophyceae	Mallomonas insignis	2,200	1.5
			Ochromonas spp.	718,022	441.3
			Pseudokephyrion sp.	2,955	0.8
			Synuropsis janei	2,955	2.6
			Cryptomonas ovata	1,375	2.2
		Cryptophyceae	Cryptomonas sp.	9,760	19.0
			Komma caudata	103,419	6.6
	-		Aphanocapsa delicatissima	101,750	0.2
			Aphanocapsa holsatica	18,750	0.1
			Aphanocapsa sp.	428,127	2.0
		Cyanobacteria	Aphanothece clathrata	33,000	0.1
			Leptolyngbya sp.	119,053	1.3
			Merismopedia warmingiana	35,200	0.0
	-	Dinophyceae	Gymnodinium sp.	2,955	5.8
	-	Euglenophyceae	Trachelomonas sp.	275	0.6
		Xanthophyceae	Ophiocytium cochleare	275	0.5
			Asterionella formosa	2,283	1.2
			Cyclotella sp.	55,639	527.4
		Bacillariophyceae	Fragilaria sp.	303	0.1
			Nitzschia sp.	(cells/L) 32,503 625 5,910 550 4,625 3,500 2,200 718,022 2,955 1,375 9,760 103,419 101,750 18,750 428,127 33,000 119,053 35,200 2,955 275 275 2,75 2,283 55,639	1.3
			Synedra sp.	330	0.6
			Ankistrodesmus falcatus	17,545	1.1
			Chlamydomonas sp.	2,817	1.5
			Elakatothrix gelatinosa	5,237	0.9
August	S3 ^(b)		Elakatothrix sp.	303	0.0
			Gloeocystis sp.	3,229	4.5
		Oblama a burnera	Monoraphidium irregulare	908	0.0
		Chlorophyceae	Monoraphidium komarkovae	8,773	0.5
			<i>Mougeotia</i> sp.	605	2.6
			Oocystis submarina	11,268	0.7
			Quadrigula closterioides	2,420	1.0
			Roya sp.		1.1
			Schroederia sp.		0.9



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Staurodesmus sp.	605	2.9
			Tetraedron incus	2,817	0.3
			Chrysocapsella planctonica	16,127	6.8
			Chrysococcus rufescens	19,719	2.7
			Dinobryon bavaricum	2,255	4.2
		Chrysophyceae	Dinobryon divergens	2,805	9.3
			Mallomonas insignis	3,025	2.1
			Ochromonas spp.	600,006	221.0
			Synuropsis janei	303	0.3
			Chroomonas sp.	5,634	0.9
		Crumtanhuasas	Cryptomonas ovata	7,563	8.3
		Cryptophyceae	Cryptomonas sp.	0	0.0
			Komma caudata	126,762	11.2
			Aphanocapsa delicatissima	79,750	0.4
			Aphanocapsa sp.	98,689	0.4
			Aphanothece clathrata	59,675	0.2
		Cyanobacteria	Aphanothece sp.	126,858	0.6
			Leptolyngbya sp.	122,029	0.8
			Merismopedia sp.	47,491	0.2
			Woronichinia naegeliana	45,375	0.1
			Amphidinium sp.	14,085	5.3
		Discrete	Gyrodinium helveticum	1,210	24.2
		Dinophyceae	Peridinium umbonatum	3,119	20.2
			Peridinium willei	303	3.2
			Asterionella formosa	225	0.1
			Aulacoseira sp.	(cells/L) 605 2,817 16,127 19,719 2,255 2,805 3,025 600,006 303 5,634 7,563 0 126,762 79,750 98,689 59,675 126,858 122,029 47,491 45,375 14,085 1,210 3,03	4.9
		Bacillariophyceae	Cyclotella sp.		60.6
			Fragilaria sp.	275	0.2
			Pinnularia sp.	(cells/L) 605 2,817 16,127 19,719 2,255 2,805 3,025 600,006 303 5,634 7,563 0 126,762 79,750 98,689 59,675 126,858 122,029 47,491 45,375 14,085 1,210 3,119 303 225 3,875 9,883 275 300 3,575 1,375 1,650 7,683 1,650 550 550 550 550 550	11.3
			Ankistrodesmus falcatus	3,575	0.4
			Dictyosphaerium pulchellum	1,375	0.3
August	S5		Elakatothrix gelatinosa	1,650	0.1
			Elakatothrix sp.	7,683	0.1
		Oblassi	Gloeocystis sp.	1,650	0.5
		Chlorophyceae	Monoraphidium komarkovae	6,325	0.3
			Mougeotia sp.	550	0.8
			Roya sp.	550	0.4
			Scenedesmus sp.	10,243	0.2
			Spondylosium sp.	550	1.6



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Staurodesmus sp.	275	1.0
			Staurodesmus triangularis var. inflatus	275	1.1
			Tetraedron incus	2,561	0.1
			Xanthidium sp.	2,561	0.4
			Chrysocapsella planctonica	14,643	4.7
			Chrysococcus rufescens	2,561	0.4
			Dinobryon bavaricum	125	0.3
		Chrysophyceae	Dinobryon divergens	1,950	5.7
			Mallomonas insignis	1,650	0.7
			Mallomonas sp.	2,561	1.4
			Ochromonas spp.	647,894	183.3
		Comptone	Cryptomonas ovata	3,300	7.2
		Cryptophyceae	Komma caudata	87,069	11.7
			Anabaena sp.	1,600	0.3
			Aphanocapsa delicatissima	60,500	0.2
			Aphanocapsa holsatica	22,000	0.1
			Aphanothece sp.	16,500	0.0
		Cyanobacteria	Gloeothece linearis	1,925	0.0
			Leptolyngbya sp.	22,650	0.1
			Merismopedia sp.	40,974	0.1
			Woronichinia naegeliana	49,500	0.2
		5.	Gymnodinium sp.	275	3.1
		Dinophyceae	Gyrodinium helveticum	550	17.9
			Asterionella formosa	90	0.1
		Bacillariophyceae	Cyclotella sp.	2,281	21.1
			Synedra sp.	275 2,561 2,561 14,643 2,561 125 1,950 1,650 2,561 647,894 3,300 87,069 1,600 60,500 22,000 16,500 22,000 16,500 1,925 22,650 40,974 49,500 275 550 90	0.4
			Ankistrodesmus falcatus		0.3
			Cosmarium sp.	2,034	1.3
			Elakatothrix gelatinosa	4,455	0.5
			Gloeocystis sp.	4,509	2.4
. .	0.5		Monoraphidium komarkovae	6,683	0.5
August	S6		Oedogonium sp.		1.4
		Chlorophyceae	<i>Oocystis</i> sp.	8,629	19.6
			Quadrigula closterioides		2.8
			Schroederia sp.		0.0
			Sphaerocystis sp.	1,733	1.2
			Spondylosium sp.		5.9
			Staurodesmus sp.		1.7
			Staurodesmus triangularis var. inflatus		2.9



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Tetraedron sp.	2,034	2.2
			Bitrichia chodatii	495	0.2
			Chrysocapsella planctonica	36,605	1.1
		Chrysophysopo	Dinobryon bavaricum	1,688	3.4
		Chrysophyceae	Dinobryon divergens	3,105	17.3
			Mallomonas insignis	2,529	2.7
			Ochromonas spp.	504,336	183.4
			Chroomonas sp.	4,067	0.2
			Cryptomonas erosa	2,281	6.2
		On mtanku saaa	Cryptomonas ovata	2,529	3.1
		Cryptophyceae	Cryptomonas sp.	2,776	9.2
			Komma caudata	77,277	10.2
			Rhodomonas sp.	2,034	1.4
			Aphanocapsa delicatissima	64,350	0.2
			Aphanocapsa holsatica	23,400	0.1
			Aphanocapsa spp.	34,650	0.1
		Cyanobacteria	Aphanothece clathrata	28,800	0.1
			Aphanothece sp.	23,513	0.1
			Leptolyngbya sp.	47,115	0.2
			Merismopedia tenuissima	35,640	0.0
		D . 1	Amphidinium sp.	4,067	1.8
		Dinophyceae	Gymnodinium sp.	2,034	4.9
			Asterionella formosa	6,555	20.1
		Bacillariophyceae	Cyclotella sp.	6,553	13.3
			Ankistrodesmus falcatus	5,632	1.2
			Botryococcus braunii	9,680	0.7
			Crucigenia tetrapedia	15,129	7.7
			Elakatothrix gelatinosa	3,872	0.2
			Hyalotheca dissiliens	640	3.2
		Chlorophyceae	Monoraphidium komarkovae	15,488	3.3
September	S2		Monoraphidium tortile	1,408	0.2
			Mougeotia sp.	192	1.2
			Oocystis sp.	11,347	2.6
			Spondylosium planum	352	0.3
			Zygnema sp.	144	2.6
			Chrysocapsa planktonica	81,317	13.3
		. .	Chrysocapsella planctonica	11,968	3.5
		Chrysophyceae	Dinobryon bavaricum	45,760	72.4
			Dinobryon divergens	193,600	299.7



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Mallomonas sp.	1,760	15.7
			Ochromonas sp.	368,762	61.3
			Synuropsis janei	7,564	24.3
			Uroglenopsis americana	51,200	8.7
		Cruptophysooo	Cryptomonas ovata	2,112	17.4
		Cryptophyceae	Komma caudata	86,990	12.2
			Anabaena sp.	1,408	0.4
		Oursehesterie	Aphanocapsa sp.	453,861	3.7
		Cyanobacteria	Leptolyngbya sp.	212,960	3.4
			Merismopedia sp.	30,257	0.8
			Asterionella formosa	2,544	7.6
		Bacillariophyceae	Cyclotella sp.	7,904	39.3
			Fragilaria sp.	1,584	2.1
	-		Ankistrodesmus falcatus	12,320	3.1
			Botryococcus braunii	16,320	3.0
		Chlorophyceae	Monoraphidium komarkovae	38,720	7.5
			Monoraphidium tortile	3,512	0.5
			Mougeotia sp.	80	1.1
September	mber S3		Spondylosium planum	432	0.5
	-		Chrysocapsella planctonica	50,924	14.4
			Dinobryon divergens	66,880	200.8
		Chrysophyceae	Mallomonas insignis	3,512	2.1
			Ochromonas sp.	359,982	75.8
			Uroglenopsis americana	453,861 212,960 30,257 2,544 7,904 1,584 12,320 16,320 38,720 3,512 80 432 50,924 66,880 3,512	0.0
			Cryptomonas ovata	4,048	31.1
		Cryptophyceae	Komma caudata	140,481	24.9
	-	Cyanobacteria	Leptolyngbya sp.	88,880	2.7
		Bacillariophyceae	Achnanthidium sp.	747	0.2
			Monoraphidium komarkovae	7,920	1.8
			Monoraphidium tortile	2,990	0.3
		Chlorophyceae	Oocystis sp.	2,990	0.4
			Roya sp.		0.5
September			Tetraedron incus	2,990	0.2
	S5		Chrysocapsella planctonica		0.5
		Chrysophyceae	Mallomonas insignis	3,737	3.0
			Ochromonas sp.	142,771	39.1
		Cryptophyceae	Komma caudata	64,284	10.9
			Aphanocapsa sp.	7,200	0.1
		Cyanobacteria	Chroococcus sp.	7,475	0.1



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Leptolyngbya sp.	15,480	0.3
			<i>Merismopedia</i> sp.	41,184	1.4

Notes: Samples were analyzed by EcoAnalysts, Inc.

cells/L = number of cells per litre; mg/m³ = milligrams per cubic metre; Late Spring = July; Summer = August; Fall = September; sp. = a single species; spp. = multiple species.

a) Replicate 1.

b) Replicate 2.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
Tenou			Asterionella formosa	3,564	5.0
			Cyclotella sp.	21,275	22.6
		Bacillariophyceae	<i>Fragilaria</i> sp.	3,168	3.9
			Tabellaria fenestrata	1,062	7.0
			Tabellaria flocculosa	360	1.8
			Ankistrodesmus fusiformis	792	0.3
			Botryococcus braunii	1,188	0.1
			Dictyosphaerium pulchellum	4,752	0.7
			Lagerheimia genevensis	6,382	0.2
		Oblassalassa	Monoraphidium arcuatum	396	0.1
		Chlorophyceae	Monoraphidium irregulare	14,520	1.0
			Monoraphidium komarkovae	16,335	3.2
			Monoraphidium tortile	13,756	1.1
			Roya sp.	594	1.0
July	FF2-1		Tetraedron incus	12,765	1.0
			Chrysocapsella planctonica	12,765	3.3
		Chrysophyceae Cryptophyceae Cyanobacteria	Dinobryon bavaricum	35,937	109.2
			Dinobryon divergens	116,679	355.4
			Ochromonas sp.	548,888	91.1
			Cryptomonas sp.	10,052	37.2
			Komma caudata	25,530	4.3
			Anabaena sp.	540	0.2
			Chroococcus sp.	42,549	0.3
			Leptolyngbya sp.	261,360	3.5
			Gyrodinium helveticum	792	26.6
		Dinophyceae	Peridinium sp.	10,454	131.2
		Euglenophyceae	Trachelomonas sp.	2,127	1.7
		Xanthophyceae	Ophiocytium parvulum	198	1.5
			Asterionella formosa	8,272	12.5
			Cyclotella sp.	6,666	8.0
		Bacillariophyceae	Fragilaria sp.	2,688	4.5
			Tabellaria fenestrata	1,636	12.4
			Tabellaria flocculosa	132	0.3
July	FF2-2		Ankistrodesmus falcatus	827	11.7
-			Elakatothrix gelatinosa	414	0.1
			Lagerheimia genevensis	8,888	0.2
		Chlorophyceae	Monoraphidium arcuatum	207	0.0
			Monoraphidium irregulare	3,102	0.2
			Monoraphidium komarkovae	12,408	4.8



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Monoraphidium tortile	1,034	0.1
			<i>Roya</i> sp.	620	1.0
			Staurastrum sp.	414	2.9
			Staurodesmus sp.	207	15.6
			Dinobryon bavaricum	282	0.9
			Dinobryon divergens	339,152	1444.5
		Chrysophyceae	Mallomonas sp.	2,429	38.2
			Ochromonas sp.	611,057	91.0
			Synuropsis janei	7,700	40.3
			Cryptomonas erosa	3,102	17.6
		Origitantia	Cryptomonas ovata	2,482	29.1
		Cryptophyceae	Cryptomonas sp.	17,935	69.8
			Komma caudata	19,998	3.0
			Anabaena circinalis	902	0.3
			Anabaena sp.	1,015	0.3
		Cyanobacteria	Leptolyngbya sp.	1,241	0.0
			Microcystis sp.	1,316	0.1
			Gyrodinium helveticum	620	20.6
		Dinophyceae	Peridinium sp.	7,378	91.8
			Peridinium willei	1,034	70.6
		Xanthophyceae	Ophiocytium cochleare	19	0.2
			Asterionella formosa	11,858	18.2
			Cyclotella sp.	15,397	119.7
		Bacillariophyceae	Fragilaria sp.	4,312	6.2
			Tabellaria fenestrata	(cells/L) 1,034 620 414 207 282 339,152 2,429 611,057 7,700 3,102 2,482 17,935 19,998 902 1,015 1,241 1,316 620 7,378 1,034 19 11,858 15,397	6.8
			Monoraphidium arcuatum		0.2
			Monoraphidium irregulare		0.5
			Monoraphidium komarkovae	23,716	5.9
		Chlorophyceae	Monoraphidium tortile	2,372	0.6
	FF0 0		Staurastrum sp.	216	2.6
July	FF2-3		Staurodesmus sp.	1,078	38.8
			Dinobryon bavaricum	211	0.6
			Dinobryon borgei	216	2.6
		Ohmensterreit	Dinobryon divergens	120,677	491.7
		Chrysophyceae	Mallomonas sp.	3,234	27.5
			Ochromonas sp.	629,354	101.8
			Synuropsis janei	6,642	30.7
		Ommetershires	Cryptomonas erosa	862	5.0
		Cryptophyceae	Cryptomonas ovata	6,468	63.5



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Cryptomonas sp.	2,587	5.2
			Komma caudata	86,044	12.7
		Cuenchesteria	Anabaena sp.	1,725	0.5
		Cyanobacteria	Leptolyngbya sp.	91,908	3.0
		Dinanhyanaa	Amphidinium sp.	2,458	2.6
		Dinophyceae	Peridinium sp.	14,230	279.1
			Achnanthidium sp.	23,560	3.9
		Desillerienhusses	Asterionella formosa	4,858	5.3
		Bacillariophyceae	Cyclotella sp.	23,560	6.0
			Tabellaria fenestrata	1,104	3.7
			Ankistrodesmus falcatus	(cells/L) 2,587 86,044 1,725 91,908 2,458 14,230 23,560 4,858 23,560	0.5
			Ankistrodesmus fusiformis	405	0.2
			Cosmarium sp.	202	0.6
			Gloeocystis sp.	9,424	1.1
			Lagerheimia genevensis	4,712	0.2
		Chlorophyceae	Monoraphidium irregulare	19,083	2.0
			Monoraphidium komarkovae	33,396	6.1
			Monoraphidium tortile	1,012	0.2
			Roya sp.	1,214	1.5
			Staurodesmus sp.	2,226	56.0
			Tetraedron incus	2,356	0.1
			Chrysococcus rufescens	16,492	11.5
July	FF2-4		Dinobryon bavaricum	239	0.5
·			Dinobryon bavaricum var. vanhoeffenii	129	0.3
		Chrysophyceae	Dinobryon divergens	186,208	414.4
July FF2-4		Mallomonas sp.	1,012	10.2	
			Ochromonas sp.	(cells/L) 2,587 86,044 1,725 91,908 2,458 14,230 23,560 4,858 23,560 4,858 23,560 1,104 810 405 202 9,424 4,712 19,083 33,396 1,012 1,214 2,226 2,356 16,492 239 129 186,208 1,012 508,891 1,417 14,061 11,780 91,883 2,356 2,576 108,944 11,780 2,024 8,618 202	126.2
			Cryptomonas erosa		15.3
	uly FF2-4		Cryptomonas ovata		103.3
		Cryptophyceae	Cryptomonas sp.	11,780	11.7
			Komma caudata	91,883	14.9
			Rhodomonas sp.	(cells/L) 2,587 86,044 1,725 91,908 2,458 14,230 23,560 4,858 23,560 1,104 810 405 202 9,424 4,712 19,083 33,396 1,012 1,214 2,226 2,356 16,492 239 129 186,208 1,012 508,891 1,417 14,061 11,780 91,883 2,356 2,576 108,944 11,780 2,024 8,618 202	5.3
			Anabaena sp.		0.7
		Cyanobacteria	Leptolyngbya sp.		1.9
			Amphidinium sp.		17.7
			Gyrodinium helveticum		89.1
		Dinophyceae	Peridinium sp.		82.0
			Peridinium willei		13.0
		Xanthophyceae	Ophiocytium parvulum		1.1



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Asterionella formosa	4,356	5.1
			Cyclotella sp.	8,894	52.3
		Bacillariophyceae	Stauroneis sp.	18	0.4
			Tabellaria fenestrata	1,620	9.2
			Tabellaria flocculosa	198	0.4
			Ankistrodesmus falcatus	990	0.3
			Ankistrodesmus fusiformis	396	0.1
			Elakatothrix gelatinosa	1,188	0.1
			Gloeocystis sp.	3,457	0.9
		Monoraphidium irregulare	14,520	1.0	
	July FF2-5 ^(a)		Monoraphidium komarkovae	43,560	8.4
		Chlorophyceae	Monoraphidium tortile	1,188	0.1
			<i>Mougeotia</i> sp.	54	1.8
			Spondylosium planum	594	0.4
			Staurastrum sp.	198	1.8
			Staurodesmus triangularis	792	7.6
			Teilingia granulata	3,186	9.8
July	ly FF2-5 ^(a)		Xanthidium johnsonii	1,729	2.1
			Dinobryon bavaricum	1,890	4.5
		Chrysophyceae	Dinobryon divergens	94,295	167.4
			Mallomonas sp.	11,163	55.2
			Ochromonas sp.	449,429	70.5
			Synuropsis janei	11,559	33.8
			Cryptomonas ovata	8,712	64.4
		Cryptophyceae	Komma caudata	17,286	2.8
			Anabaena sp.	5,922	3.8
			Chroococcus sp.	13,829	0.5
			Leptolyngbya sp.	19,440	0.2
		Cyanobacteria	Merismopedia sp.	41,486	5.2
			Microcystis sp.	6,480	0.8
			Pseudanabaena sp.	10,371	0.2
			Amphidinium sp.	8,643	20.9
		Dinophyceae	Gyrodinium helveticum	0	0.0
			Peridinium sp.	6,534	56.3
		Xanthophyceae	Ophiocytium parvulum	396	2.4
			Cyclotella sp.	16,583	147.7
	(b)	Bacillariophyceae	Fragilaria sp.	4,963	5.9
July	FF2-5 ^(b)		Tabellaria fenestrata	1,034	6.0
		Chlorophyceae	Chlorotetraedron incus	2,626	0.3



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Euastrum elegans	207	4.9
			Lagerheimia genevensis	21,008	1.1
			Monoraphidium irregulare	13,649	1.0
			Monoraphidium komarkovae	17,061	5.8
			Spondylosium planum	432	0.4
			<i>Zygnema</i> sp.	38	1.5
			Dinobryon bavaricum	4,512	10.5
			Dinobryon bavaricum var. vanhoeffenii	545	1.7
			Dinobryon borgei	1,448	3.4
		Chrysophyceae	Dinobryon divergens	119,841	637.1
			Mallomonas sp.	1,241	6.5
			Ochromonas sp.	685,394	144.5
			Synuropsis janei	1,861	8.0
			Cryptomonas ovata	2,482	30.3
		Cryptophyceae	Cryptomonas sp.	9,739	37.1
			Komma caudata	47,269	7.9
			Anabaena sp.	978	0.4
		Cyanobacteria	Chroococcus sp.	7,878	0.1
			Leptolyngbya sp.	73,162	1.3
			Gyrodinium helveticum	1,654	64.0
		Dinophyceae	Peridinium sp.	620	12.6
			Peridinium willei	4,550	404.7
		Xanthophyceae	Ophiocytium parvulum	1,034	2.5
			Asterionella formosa	2,633	3.1
			Cyclotella sp.	31,857	166.4
		Desillerianhusses	Fragilaria sp.	495	0.6
		Bacillariophyceae	Synedra sp.	(cells/L) 207 21,008 13,649 17,061 432 38 4,512 545 1,448 119,841 1,241 685,394 1,861 2,482 9,739 47,269 978 7,878 73,162 1,654 620 4,550 1,034 2,633 31,857	0.7
			Tabellaria fenestrata		39.1
			Tabellaria flocculosa	1,193	2.3
			Ankistrodesmus falcatus	5,198	0.3
August			Chlamydomonas sp.	6,914	2.6
August	FF2-1		Chlorella ellipsoidea	12,100	1.4
			Closteriopsis acicularis	743	0.2
		Chlorophyses	Crucigenia quadrata	990	0.1
		Chlorophyceae	Crucigeniella rectangularis	6,914	1.3
			Dictyosphaerium pulchellum	4,703	0.3
			Elakatothrix gelatinosa	8,174	5.0
			Elakatothrix sp.	180	0.0
			Gloeocystis sp.	2,471	3.8



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Monoraphidium irregulare	24,235	0.7
			Monoraphidium komarkovae	20,499	1.0
			Nephrocytium agardhiunum	2,475	0.5
			<i>Oocystis</i> sp.	2,719	19.6
			Quadrigula closterioides	1,238	0.1
			<i>Roya</i> sp.	5,689	5.4
			Sphaerocystis sp.	53,837	13.1
			Spirogyra sp.	90	1.9
			Spondylosium sp.	2,089	5.0
			Staurodesmus incus	1,238	6.1
			Tetraedron incus	31,114	2.5
			Chrysocapsella planctonica	8,643	1.0
			Dinobryon bavaricum	13,241	20.1
		0	Dinobryon divergens	6,395	30.5
		Chrysophyceae	Mallomonas insignis	1,980	2.1
			Ochromonas spp.	326,700	137.9
			Uroglenopsis americana	1,350	0.1
			Cryptomonas ovata	9,900	14.8
		Cryptophyceae	Komma caudata	95,071	4.0
			Rhodomonas sp.	1,729	1.9
			Anabaena sp.	1,418	0.0
			Aphanocapsa delicatissima	168,188	1.3
			Aphanocapsa elachista	80,438	0.3
			Aphanocapsa sp.	152,350	0.4
		Cyanobacteria	Aphanothece clathrata	298,350	1.5
			Leptolyngbya sp.	294,340	3.1
			Merismopedia punctata	17,820	0.0
			Merismopedia sp.	41,486	0.1
			Woronichinia naegeliana	9,900	0.1
		Dinophyceae	Gyrodinium helveticum	743	8.9
			Achnanthidium sp.	1,729	1.0
			Asterionella formosa	2,318	3.0
		Desillerississes	Cyclotella sp.	34,819	356.7
		Bacillariophyceae	Synedra sp.	158	0.1
August	FF2-2		Tabellaria fenestrata	3,128	11.9
			Tabellaria flocculosa	833	5.0
			Ankistrodesmus falcatus	5,198	0.5
		Chlorophyceae	Botryococcus protuberans	14,850	0.7
			Chlamydomonas sp.	1,729	0.2



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Chlorella ellipsoidea	8,643	1.5
			Closteriopsis acicularis	270	0.3
			Cosmarium sp.	248	0.4
			Dictyosphaerium pulchellum	3,214	1.0
			Elakatothrix gelatinosa	495	0.0
			Elakatothrix sp.	8,643	0.1
			Gloeocystis sp.	1,729	0.5
			Lagerheimia genevensis	1,729	0.2
			Monoraphidium arcuatum	2,471	0.4
			Monoraphidium irregulare	7,779	0.3
			Monoraphidium komarkovae	8,765	0.7
			<i>Oocystis</i> sp.	10,371	0.9
			Roya sp.	743	0.6
			Sphaerocystis sp.	9,900	1.8
			Spondylosium sp.	1,665	5.1
			Staurodesmus triangularis var. inflatus	248	0.3
			Tetraedron incus	25,929	3.1
			Chrysocapsella planctonica	9,389	3.2
		Chrysophyceae	Dinobryon bavaricum	4,915	13.8
			Dinobryon divergens	4,523	22.6
			Mallomonas insignis	3,461	5.7
			Mallomonas multiunca	248	0.4
			Ochromonas spp.	278,300	70.8
			Synuropsis janei	5,433	9.8
			Uroglenopsis americana	3,600	0.5
			Chroomonas sp.	3,457	1.6
		Cryptophyceae	Cryptomonas ovata	9,397	27.8
			Komma caudata	117,543	9.1
		Cyanobacteria	Aphanocapsa delicatissima	112,613	0.2
			Aphanocapsa holsatica	66,623	0.3
			Aphanocapsa sp.	255,416	1.0
			Aphanothece clathrata	76,950	0.5
			Aphanothece sp.	40,543	0.2
			Leptolyngbya sp.	164,462	1.8
		Dinophyceae	Gymnodinium sp.	5,186	1.4
		Euglenophyceae	Monomorphina sp.	248	0.4
August	FF2-3	Bacillariophyceae	Asterionella formosa	13,002	19.0
			Cyclotella sp.	54,467	403.7
			Synedra sp.	825	1.6



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Tabellaria fenestrata	3,100	23.7
			Tabellaria flocculosa	400	3.7
			Ankistrodesmus falcatus	4,950	0.7
			Botryococcus braunii	11,000	0.9
			Botryococcus protuberans	16,500	1.6
			Chlamydomonas sp.	9,603	3.8
			Chlorella ellipsoidea	9,603	1.7
			Cosmarium margaritatum	275	1.0
			Elakatothrix gelatinosa	300	0.0
			Elakatothrix sp.	7,202	0.1
			<i>Gloeocystis</i> sp.	4,802	1.1
			Monoraphidium arcuatum	2,676	0.1
		Chlorophysess	Monoraphidium irregulare	7,975	0.4
		Chlorophyceae	Monoraphidium komarkovae	9,900	0.4
			Monoraphidium minutum	2,401	0.1
			Quadrigula closterioides	1,100	0.3
			Roya sp.	5,976	5.1
			Schroederia sp.	550	0.2
			Sphaerocystis sp.	20,603	2.5
			Spondylosium sp.	1,300	5.1
			Staurodesmus incus	275	0.8
			Staurodesmus incus var. indentatus	1,100	4.2
Period			Staurodesmus triangularis	275	2.3
			Tetraedron incus	19,206	1.4
			Bitrichia chodatii	550	0.2
			Chrysocapsella planctonica	30,259	13.0
			Dinobryon bavaricum	11,399	48.6
		Chrysophyceae	Dinobryon divergens	8,250	37.0
			Mallomonas insignis	6,050	5.1
			Ochromonas spp.	516,171	123.8
			Uroglenopsis americana	11,900	4.3
			Cryptomonas ovata	7,975	11.3
		Cryptophyceae	Komma caudata	84,028	5.5
		<u> </u>	Anabaena sp.	500	0.1
			Aphanocapsa delicatissima	30,000	0.1
			Aphanocapsa elachista	49,500	0.1
		Cyanobacteria	Aphanocapsa holsatica	20,750	0.0
			Aphanocapsa sp.	132,917	0.8
			Aphanothece clathrata	57,000	0.2



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)	
			Aphanothece sp.	5,500	0.0	
			Leptolyngbya sp.	176,943	2.1	
			Pseudanabaena sp.	12,375	0.2	
			Woronichinia naegeliana	16,500	0.0	
		Dipaphyaaaa	Gyrodinium helveticum	275	8.0	
		Dinophyceae	Peridinium sp.	275	1.8	
		Euglenophyceae	Trachelomonas sp.	2,676	20.4	
			Achnanthidium sp.	4,852	0.9	
			Asterionella formosa	6,747	6.4	
		De sille sie sterre e s	Cyclotella sp.	10,584	139.1	
		Bacillariophyceae	Fragilaria sp.	1,320	0.9	
			Tabellaria fenestrata	2,800	46.4	
			Tabellaria flocculosa	440	5.0	
			Ankistrodesmus falcatus	7,920	1.3	
			Botryococcus braunii	2,200	0.7	
			Chlamydomonas sp.	1,617	0.6	
			Chlorella ellipsoidea	8,087	1.1	
			Crucigenia quadrata	8,087	(mg/m ³) 0.0 2.1 0.2 0.0 8.0 1.8 20.4 0.9 6.4 139.1 0.9 46.4 5.0 1.3 0.7 0.6	
			Dictyosphaerium pulchellum	2,200	(mg/m ³) 0.0 2.1 0.2 0.0 8.0 1.8 20.4 0.9 6.4 139.1 0.9 46.4 5.0 1.3 0.7 0.6 1.1 1.4 0.1 0.1 0.1 0.1 0.1 0.1 0.1 0.1	
			Elakatothrix gelatinosa	1,980	0.1	
			Elakatothrix sp.	4,852	0.1	
			Gloeocystis sp.	1,617	0.4	
A			Lagerheimia genevensis	1,617	0.1	
August	FF2-4 ^(a)	Ohlananharaaaa	Monoraphidium arcuatum	3,675	0.2	
		Chlorophyceae	Monoraphidium irregulare	6,820	0.3	
			Monoraphidium komarkovae	15,400	0.9	
			Nephrocytium sp.	880	0.1	
			<i>Oocystis</i> sp.	6,470	0.5	
			Quadrigula closterioides	4,180	1.7	
			<i>Roya</i> sp.	8,295	9.9	
			Sphaerocystis sp.	87,029	7.8	
			Staurodesmus incus	880	5.8	
			Staurodesmus incus var. indentatus	440	2.8	
			Staurodesmus triangularis	660	4.3	
			Tetraedron incus	8,087	0.4	
			Bitrichia chodatii	440	0.4	
		Ohmana	Chrysocapsella planctonica	16,019	2.0	
		Chrysophyceae	Dinobryon bavaricum	13,501	22.8	
			Dinobryon divergens	8,800	51.4	



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)		
			Mallomonas insignis	3,597	5.5		
			Mallomonas sp.	4,852	3.6		
			Ochromonas spp.	1,617	0.7		
Period			Synuropsis janei	220	0.3		
			Uroglenopsis americana	8,600	1.0		
			Cryptomonas ovata	3,597	8.7		
		Cryptophyceae	Cryptomonas sp.	880	1.5		
			Komma caudata	12,939	1.3		
			Anabaena sp.	3,480	0.2		
			Aphanocapsa delicatissima	138,900	0.5		
			Aphanocapsa holsatica	121,300	0.3		
			Aphanocapsa sp.	65,843	0.2		
		Cyanobacteria	Aphanothece clathrata	capsa sp. 65,843 hece clathrata 63,820 hece sp. 69,300 ece linearis 6,470 gbya sp. 292,490 pedia sp. 58,226 um helveticum 220 hidium sp. 2,134 ella formosa 7,025 la sp. 60,578			
Period			Aphanothece sp.	69,300	0.3		
			Gloeothece linearis	6,470	0.0		
			Leptolyngbya sp.	292,490	2.1		
			Merismopedia sp.	58,226	1.2		
		Dinophyceae	Gyrodinium helveticum	220	4.6		
			Achnanthidium sp.	2,134	0.5		
			Asterionella formosa	2,134 0.5 7,025 6.8			
			Cyclotella sp.	60,578	558.9		
		Bacillariophyceae	Synedra sp.	1,375	1.2		
			Tabellaria fenestrata	12,928	97.8		
			Tabellaria flocculosa	2,475	26.1		
Period			Ankistrodesmus falcatus	9,625	1.3		
			Chlamydomonas sp.	2,134	1.1		
			Crucigenia quadrata	2,134	0.6		
			Dictyosphaerium pulchellum	3,025	0.4		
Period	FF2-4 ^(b)		Elakatothrix gelatinosa	1,650	0.0		
			Elakatothrix sp.	4,268	0.1		
			Lagerheimia genevensis	2,134	0.0		
		Chlorophyceae	Monoraphidium arcuatum	7,227	0.3		
			Monoraphidium irregulare	5,225	0.2		
			Monoraphidium komarkovae	12,467	0.6		
August			Mougeotia sp.	275	1.0		
			Nephrocytium sp.	2,200	0.2		
			Oocystis sp.	550	0.1		
			Oocystis submarina	6,402	0.2		
			Quadrigula closterioides	450	0.1		



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			<i>Quadrigula</i> sp.	925	1.0
			<i>Roya</i> sp.	2,475	1.7
			Spondylosium sp.	4,059	11.7
			Staurastrum sp.	275	1.3
			Staurodesmus triangularis	825	5.1
			Tetraedron incus	14,938	0.9
			Chrysocapsella planctonica	19,206	2.0
			Dinobryon bavaricum	9,412	84.7
			Dinobryon divergens	4,725	18.8
			Mallomonas insignis	5,643	9.0
		Chrysophyceae	Mallomonas multiunca	825	1.7
			Mallomonas sp.	2,134	1.1
			Ochromonas spp.	435,344	148.1
			Synuropsis janei	275	0.5
			Uroglenopsis americana	7,750	0.9
			Chroomonas sp.	2,134	0.1
		Cryptophyceae	Cryptomonas ovata	8,393	21.7
			Komma caudata	91,764	5.1
			Anabaena sp.	5,600	0.3
			Aphanocapsa delicatissima	133,375	0.1
			Aphanocapsa holsatica	12,500	0.0
		Cyanobacteria	Aphanocapsa sp.	221,666	1.1
			Aphanothece clathrata	144,750	0.5
			Aphanothece sp.	13,750	0.0
			Leptolyngbya sp.	216,139	2.0
		Disasteras	Gymnodinium sp.	2,134	3.5
		Dinophyceae	Gyrodinium helveticum	275	11.0
			Asterionella formosa	6,563	6.9
			Cyclotella sp.	180,753	1689.5
		De elle de la	Fragilaria sp.	220	0.1
		Bacillariophyceae	Nitzschia sp.	220	0.1
			Tabellaria fenestrata	2,100	13.7
A			Tabellaria flocculosa	1,140	31.3
August	FF2-5		Ankistrodesmus falcatus	2,200	0.3
			Botryococcus braunii	8,800	0.3
		Ohlanasi	Chlamydomonas sp.	880	0.2
		Chlorophyceae	Chlorella ellipsoidea	2,364	0.8
			Cosmarium margaritatum	220	0.9
			Cosmarium sp.	440	0.6



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)		
			Crucigenia fenestrata	7,092	1.3		
			Dictyosphaerium pulchellum	5,940	0.4		
			Elakatothrix sp.	2,364	0.0		
			Gloeocystis sp.	2,364	0.6		
			Monoraphidium arcuatum	220	0.0		
			Monoraphidium irregulare	3,300	0.1		
			Monoraphidium komarkovae	10,371	0.8		
			Nephrocytium agardhiunum	880	0.2		
			Oocystis submarina	10,172	1.3		
			Quadrigula closterioides	2,640	0.2		
			Roya sp.	6,708	6.1		
			Sphaerocystis sp.	37,822	0.7		
			Spondylosium sp.	860	2.5		
			Staurastrum sp.	20	0.8		
			Staurodesmus incus	660	3.3		
			Staurodesmus sp.	220	3.4		
			Tetraedron incus	16,547	0.9		
			Chrysocapsella planctonica	4,400	0.7		
			Dinobryon bavaricum	7,402	24.8		
			Dinobryon divergens	6,893	40.7		
			Mallomonas insignis	1,980	1.5		
		Chrysophyceae	Mallomonas multiunca	660	1.7		
			Ochromonas spp.	408,947	129.3		
			Synuropsis janei	2,804	3.1		
			Synuropsis sp.	1,100	0.4		
			Uroglenopsis americana	5,000	0.7		
			Cryptomonas ovata	6,764	12.1		
		Cryptophyceae	Komma caudata	73,280	7.1		
			Anabaena sp.	15,280	1.4		
			Anabaena spp.	580	0.1		
			Aphanocapsa delicatissima	45,980	0.1		
			Aphanocapsa holsatica	19,020	0.1		
		Cyanobacteria	Aphanocapsa sp.	129,460	0.3		
			Aphanothece clathrata	107,800	0.5		
			Gloeothece linearis	1,980	0.1		
			Leptolyngbya sp.	206,756	1.6		
			Pseudanabaena sp.	9,455	0.1		
			Gyrodinium helveticum	220	1.4		
		Dinophyceae	Peridinium sp.	2,364	5.7		



September	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)	
			Peridinium umbonatum	2,364	0.9	
		Euglenophyceae	Monomorphina sp.	220	2.5	
		Xanthophyceae	Ophiocytium cochleare	440	1.5	
			Asterionella formosa	2,250	3.3	
		Pagillarianhyagaa	Cyclotella sp.	1,257	1.3	
		Bacillariophyceae	<i>Fragilaria</i> sp.	2,970	4.9	
			Tabellaria fenestrata	15,840	46.5	
			Ankistrodesmus falcatus	55,176	13.1	
			Elakatothrix gelatinosa	3,960	0.4	
			Monoraphidium arcuatum	198	0.0	
			Monoraphidium irregulare	2,376	0.2	
			Monoraphidium komarkovae	21,780	4.3	
			Monoraphidium tortile	3,504	0.9	
		Chlorophyceae	<i>Mougeotia</i> sp.	216	7.6	
			<i>Roya</i> sp.	4,752	4.1	
			Spondylosium planum	990	0.8	
September			Staurastrum sp.	198	1.1	
	FF2-1		Staurodesmus sp.	2,178	(mg/m ³) 0.9 2.5 1.5 3.3 1.3 4.9 46.5 13.1 0.4 0.0 0.2 4.3 0.9 7.6 4.1 0.8	
			Tetraedron incus	2,514	0.3	
			Tetrastrum triangulare	35,200	13.1 0.4 0.0 0.2 4.3 0.9 7.6 4.1 0.8 1.1 112.1 0.3 13.4 10.4 3.1 44.2 0.0 16.0 8.8 0.0 0.1 0.1	
			Dinobryon bavaricum	5,940	10.4	
			Dinobryon divergens	1,890	3.1	
		Chrysophyceae	Ochromonas sp.	292,914	44.2	
September			Uroglenopsis americana	36	0.0	
			Cryptomonas erosa	1,980	16.0	
		Cryptophyceae	Komma caudata	61,600	8.8	
			Anabaena circinalis	54	0.0	
			Chroococcus sp.	12,571	0.1	
		Cyanobacteria	Coelosphaerium sp.	30,171	0.1	
			Leptolyngbya sp.	252,189	4.0	
		Dinophyceae	Peridinium willei	594	40.0	
		Euglenophyceae	Trachelomonas sp.	198	2.8	
			Asterionella formosa	1,600	4.2	
			Cyclotella sp.	5,797	7.1	
		Bacillariophyceae	Fragilaria sp.	220	0.3	
September	FF2-2		Tabellaria fenestrata	9,429	45.9	
			Ankistrodesmus falcatus	14,080		
		Chlorophyceae	Monoraphidium arcuatum	1,100		
			Monoraphidium irregulare	4,840		



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Monoraphidium komarkovae	29,040	7.1
			Monoraphidium tortile	1,100	0.2
			<i>Oocystis</i> sp.	8,381	5.7
			<i>Roya</i> sp.	2,200	2.5
			Spondylosium planum	540	0.4
			Staurodesmus sp.	880	45.3
			Tetraedron incus	16,762	0.6
			Tetrastrum triangulare	24,947	10.3
			Dinobryon bavaricum	2,700	4.9
			Dinobryon divergens	740	2.1
			Mallomonas sp.	2,057	28.6
		Chrysophyceae	Ochromonas sp.	321,270	76.4
			Synuropsis janei	6,027	34.0
			Uroglenopsis americana	100	0.0
Period			Cryptomonas ovata	7,260	57.4
		Cryptophyceae	Cryptomonas sp.	660	1.5
Period			Komma caudata	48,889	6.4
			Anabaena sp.	40	0.0
			Aphanocapsa delicatissima	8,060	0.0
		Cyanobacteria	Aphanothece paralleliformis	28,000	0.2
			Chroococcus sp.	50,286	0.4
			Leptolyngbya sp.	8,541	0.1
			Achnanthidium sp.	2,794	0.4
			Asterionella formosa	800	1.0
			Cyclotella sp.	4,190	3.2
		Bacillariophyceae	Fragilaria sp.	660	1.0
			Tabellaria fenestrata	27,955	115.6
September			Tabellaria flocculosa	1,500	2.3
			Ankistrodesmus falcatus	11,440	2.6
			Dictyosphaerium pulchellum	2,640	0.4
September	FF2-3		Elakatothrix gelatinosa	1,760	0.3
			Monoraphidium irregulare	3,080	0.3
eptember F			Monoraphidium komarkovae	29,040	4.0
		Chlorophyceae	<i>Oocystis</i> sp.	5,587	1.7
			Roya sp.	5,500	7.2
September			Spondylosium planum	520	0.5
			Staurodesmus sp.	660	21.8
			Staurodesmus triangularis	440	18.7
			Tetrastrum triangulare	3,520	0.9



September	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)
			Chrysocapsella planctonica	4,190	0.9
			Dinobryon divergens	2,020	6.3
		Chrysophyceae	Ochromonas sp.	314,286	89.4
			Synuropsis janei	1,540	6.6
Period			Uroglenopsis americana	14,000	1.4
			Cryptomonas ovata	3,520	35.0
		Cryptophyceae	Cryptomonas sp.	660	1.4
			Komma caudata	60,063	8.6
			Anabaena sp.	80	0.0
		Quere la staria	Aphanothece paralleliformis	240	0.0
		Cyanobacteria	Chroococcus sp.	78,222	0.7
			Leptolyngbya sp.	176,000	3.3
Period		Dinophyceae	Peridinium sp.	2,200	16.0
			Asterionella formosa	7,740	24.4
			Cyclotella sp.	5,139	17.8
		Bacillariophyceae	Fragilaria sp.	2,376	2.7
September			Tabellaria fenestrata	8,473	38.4
			Ankistrodesmus falcatus	27,720	7.1
			Ankistrodesmus fusiformis	396	0.1
			Elakatothrix gelatinosa	792	0.1
			Euastrum elegans	396	6.2
			Monoraphidium irregulare	2,970	0.3
Period			Monoraphidium komarkovae	32,670	6.0
		Chlorophyceae	Roya sp.	7,920	6.7
			Spondylosium planum	2,268	3.4
			Staurodesmus sp.	3,762	210.8
	FF2-4		Tetraedron incus	11,853	0.6
			Tetrastrum triangulare	86,922	15.6
			Zygnema sp.	108	1.4
			Chrysocapsella planctonica	9,900	3.3
			Chrysococcus rufescens	9,219	3.5
September			Dinobryon bavaricum	2,052	4.2
		Chrysophyceae	Dinobryon divergens	1,512	2.2
			Mallomonas sp.	2,178	12.3
			Ochromonas sp.	280,522	70.0
			Uroglenopsis americana	32,400	4.4
			Cryptomonas ovata	14,385	248.8
		Cryptophyceae	Cryptomonas sp.	5,268	7.1
			Komma caudata	53,997	8.8



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (cells/L)	Biomass (mg/m³)					
			Aphanocapsa sp.	210,721	1.7					
			Aphanothece paralleliformis	23,400	0.1					
		Cyanobacteria	Chroococcus sp.	79,020	0.8					
			Leptolyngbya sp.	544,500	8.3					
			Merismopedia sp.	43,560	0.7					
			Asterionella formosa	1,440	2.0					
		Desillerienhysees	Cyclotella sp.	10,637	7.1					
		Bacillariophyceae	Fragilaria sp.	2,178	5.3					
			Tabellaria fenestrata	13,860	48.8					
			Ankistrodesmus falcatus	11,880	2.5					
			Ankistrodesmus fusiformis	11,880	2.8					
			Crucigenia fenestrata	34,040	11.7					
			Monoraphidium arcuatum	nia fenestrata 34,040 ohidium arcuatum 396						
			Monoraphidium irregulare	7,687	(mg/m ³) 1.7 0.1 0.8 8.3 0.7 2.0 7.1 5.3 48.8 2.5 2.8					
			Monoraphidium komarkovae	87,120	21.1					
		Chlorophyceae	Mougeotia sp.	144	0.4					
			<i>Oocystis</i> sp.	12,765	4.9					
			<i>Roya</i> sp.	4,356	4.0					
			Spondylosium planum	1,548	1.2					
0	FF0 F		Staurodesmus sp.	6,878	163.5					
September	FF2-5		Teilingia excavata	594	(mg/m³) 1.7 0.1 0.8 8.3 0.7 2.0 7.1 5.3 48.8 2.5 2.8 11.7 0.1 1.1 2.5 2.8 11.7 0.1 1.1 21.1 0.4 4.9 4.0 1.2 163.5 0.2 12.0 3.3 5.5 2.7 89.3 5.0 69.0 4.6 10.4 3.3					
			Chrysocapsella planctonica	29,785						
			Dinobryon bavaricum	1,890	8.3 0.7 2.0 7.1 5.3 48.8 2.5 2.8 11.7 0.1 1.1 21.1 0.4 4.9 4.0 1.2 163.5 0.2 12.0 3.3 5.5 2.7 89.3 5.0 69.0 4.6 10.4 3.3 5.0 69.0 4.6 10.4 3.3					
			Dinobryon divergens	2,700	0.1 0.8 8.3 0.7 2.0 7.1 5.3 48.8 2.5 2.8 11.7 0.1 1.1 21.1 0.4 4.9 4.0 1.2 163.5 0.2 12.0 3.3 5.5 2.7 89.3 5.0 69.0 4.6 10.4 3.3 5.0 69.0 4.6 10.4 3.3 0.2 5.1 1.0					
		Chrysophyceae	Mallomonas insignis	4,255						
			Ochromonas sp.	489,319	2.5 2.8 11.7 0.1 1.1 21.1 0.4 4.9 4.0 1.2 163.5 0.2 12.0 3.3 5.5 2.7 89.3 5.0 69.0					
			Uroglenopsis americana	21,600	0.8 8.3 0.7 2.0 7.1 5.3 48.8 2.5 2.8 11.7 0.1 1.1 21.1 0.4 4.9 4.0 1.2 163.5 0.2 12.0 3.3 5.5 2.7 89.3 5.0 69.0 4.6 10.4 3.3 5.0 5.1					
			Cryptomonas ovata	6,375	69.0					
		Cryptophyceae	Cryptomonas sp.	6,382	4.6					
			Komma caudata	70,207	10.4					
			Aphanocapsa sp.	594 0.2 29,785 12.0 1,890 3.3 2,700 5.5 4,255 2.7 489,319 89.3 21,600 5.0 6,375 69.0 6,382 4.6 70,207 10.4						
			Aphanothece paralleliformis	21,600	0.2					
		Cyanobacteria	Leptolyngbya sp.	326,700	5.1					
			Merismopedia sp.	37,224	1.0					
		Euglenophyceae	Trachelomonas sp.	2,127	6.5					

Table A-6	Phytoplankton Abundance and Biomass in Lac de Gras, Far-Field 2 Area, 2014
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Notes: Samples were analyzed by EcoAnalysts, Inc.

cells/L = number of cells per litre; mg/m³ = milligrams per cubic metre; Late Spring = July; Summer = August; Fall = September; sp. = a single species; spp. = multiple species.

a) Replicate 1.

b) Replicate 2.



									Relative Abundance (%) State Relative (%)							Relative Biomass (%)									
Basin	Waterbody	Station	Sampling Period	Replicate	Total Abundance (cells/L)	Total Biomass (mg/m³)	Taxonomic Richness (No. of Taxa)	Bacillariophyceae	Chlorophyceae	hrysophyce	Cryptophyceae	Cyanobacteria	uglenophyce	Dinophyceae	Xanthophyceae	Bacillariophyceae	Chlorophyceae	Chrysophyceae	Cryptophyceae	Cyanobacteria	Euglenophyceae	Dinophyceae	Xanthophyceae		
		Aa-1	Summer		1,565,282	545	33	1.6	5.3	62.0	18.6	11.6	0.9	0.0	0.0	7.7	4.7	69.2	12.8	2.4	3.3	0.0	0.0		
		Ad-1	Fall		896,782	1,515	18	15.1	4.0	68.5	4.8	4.5	0.0	3.1	0.0	27.5	1.3	32.9	6.0	0.0	0.0	32.3	0.0		
			Late Spring		457,734	244	17	3.6	4.4	77.4	4.2	8.2	0.0	2.3	0.0	16.0	0.7	27.0	10.8	2.0	0.0	43.5	0.0		
		Ab-1	Summer		1,209,854	271	30	1.1	5.2	65.7	10.1	17.7	0.2	0.0	0.0	13.3	11.3	64.3	8.8	1.6	0.7	0.0	0.0		
			Fall		502,497	704	23	16.3	13.0	57.2	10.3	0.5	0.0	2.6	0.0	49.4	7.0	15.9	2.1	0.1	0.0	25.5	0.0		
			Late Spring		539,871	248	27	2.1	2.8	84.4	3.3	4.8	2.6	0.0	0.0	12.4	6.3	41.0	11.1	5.2	23.9	0.0	0.0		
		Ac-1	Summer		795,246	451	48	5.1	17.0	41.7	26.9	6.4	0.4	2.3	0.2	23.5	10.3	30.5	10.0	0.6	9.1	15.8	0.1		
		1.0	Fall	Rep 1	866,090	1,065	20	20.6	5.1	60.7	5.4	6.3	0.0	2.0		43.1	3.1	34.0	2.0	2.1	0.0	15.7	0.0		
			Fall	Rep 2	933,088	1,032	24	18.6	4.6	60.8	6.2	7.3	0.0	2.5	0.0	50.7	2.3	28.3	2.0	3.2	0.0	13.5	0.0		
			Late Spring		638,033	282	30	1.0	4.9	62.5	4.0	26.4	0.0	1.1	0.0	4.8	7.1	29.5	6.8	31.7	0.0	20.1	0.0		
	Lac du Sauvage	Ac-4	Summer		726,728	283	32	3.9	10.1	62.9	11.6	9.0	0.2	2.3	0.0	38.7	11.5	34.5	1.7	2.1	0.3	11.2	0.0		
Lac du Sauvage			Fall		618,797	308	22	5.5	3.2	70.5	12.3	7.5	0.0	1.0	0.0	21.3	1.6	47.5	14.7	1.4	0.0	13.4	0.0		
			Late Spring		721,233	483	21	1.2	3.4	74.9	4.0	15.4	0.0	1.0	0.0	3.0	3.4	30.1	15.2	3.8	0.0	44.3	0.3		
		Ac-7	Summer		914,129	508	40	4.4	18.8	49.3	18.4	5.5	0.4	3.1	0.0	22.8	16.8	33.5	7.5	0.7	0.7	18.0	0.0		
			Fall		549,905	294	19	2.3	5.4	75.3	7.0	8.6	0.3	1.2	0.0	14.9	2.5	56.7	4.6	7.3	0.6	13.4	0.0		
			Late Spring		789,188	198	19	1.3	4.9	88.1	1.5	4.0	0.0	0.3	0.0	8.0	4.3	66.2	14.5	5.6	0.0	1.3	0.0		
		Ad-1	Summer		676,103	473	37	2.5	15.3	51.0	13.8	15.2	0.5	1.7	0.0	11.1	18.8	58.8	2.1	0.3	0.3	8.6	0.0		
			Fall		512,757	284	18	7.6	1.0	80.3	7.7	3.0	0.0	0.4	0.0	37.9	0.9	48.3	2.6	2.5	0.0	7.8	0.0		
			Late Spring		955,784	295	25	3.2	2.0	79.0	1.7	13.1	0.0	1.0	0.0	12.1	4.5	26.0	16.8	28.5	0.0	11.4	0.6		
		Ae-1	Summer		1,643,909	447	32	1.5	12.7	67.4	6.6	11.5	0.0	0.3	0.0	12.2	11.9	61.7	5.2	2.9	0.0	6.2	0.0		
			Fall		692,645	747	21	7.9	4.0	76.7	4.9	3.2	0.0	3.2	0.0	17.3	1.9	33.6	8.1	0.9	0.1	38.0	0.0		
	Duchess Lake	Af-1	Summer		3,287,508	1,327	34	1.3	6.0	78.4	8.2	5.4	0.0	0.8	0.0	5.5	3.2	73.8	3.9	1.5	0.1	11.9	0.0		
			Fall		954,969	512	16	3.7	6.0	71.2	14.5	4.1	0.5	0.0	0.0	8.3	16.0	64.6	7.2	0.1	3.9	0.0	0.0		

Table A-7 Summary of Phytoplankton Community Data for Lakes in the Jay Project Area, 2014

2014 Plankton Supplemental Baseline Report Jay Project Appendix A, 2014 Plankton Taxonomy Data April 2015



							Relative Abundance (%) Relative Biomass (%)									%)							
Basin	Waterbody	Station	Sampling Period	Replicate	Total Abundance (cells/L)	Total Biomass (mg/m³)	Taxonomic Richness (No. of Taxa)	Bacillariophyceae	Chlorophyceae	Chrysophyceae	Cryptophyceae	Cyanobacteria	Euglenophyceae	Dinophyceae	Xanthophyceae	Bacillariophyceae	Chlorophyceae	Chrysophyceae	Cryptophyceae	Cyanobacteria	Euglenophyceae	Dinophyceae	Xanthophyceae
			Late Spring		808,528	313	17	0.9	3.5	86.0	7.7	0.5	0.0	1.3	0.0	6.8	2.0	38.9	42.6	0.1	0.0	9.6	0.0
		S2	Summer		1,298,240	898	33	4.1	9.8	44.7	9.5	31.3	0.6	0.0	0.0	49.5	5.7	39.1	3.2	0.5	2.1	0.0	0.0
			Fall		1,626,511	593	25	0.8	3.9	46.8	5.5	42.9	0.0	0.0	0.0	5.6	3.9	84.1	5.0	1.4	0.0	0.0	0.0
			Late Spring		585,574	406	14	1.4	4.8	88.0	5.7	0.0	0.0	0.1	0.0	2.8	1.7	90.6	2.1	0.0	0.0	2.8	0.0
		S3	Summer	Rep 1	1,761,738	626	34	0.7	9.1	41.7	6.5	41.8	0.2	0.0	0.0	13.6	3.8	76.5	4.4	0.6	0.9	0.1	0.1
			Summer	Rep 2	1,500,885	871	35	4.0	3.9	42.9	9.3	38.6	0.0	1.2	0.0	60.9	2.1	28.3	2.3	0.3	0.0	6.1	0.0
	Slipper Bay		Fall		798,203	417	16	1.5	8.9	60.3	18.1	11.1	0.0	0.0	0.0	11.8	3.8	70.4	13.4	0.6	0.0	0.0	0.0
			Late Spring	Rep 1	603,217	592	14	3.3	3.4	89.1	2.6	1.1	0.0	0.4	0.0	5.7	4.9	83.3	3.2	0.0	0.0	2.5	0.4
		S5	Late Spring	Rep 2	534,247	576	18	3.6	4.7	86.5	2.4	2.0	0.0	0.7	0.1	7.5	3.7	80.4	4.2	0.0	0.0	3.5	0.7
			Summer		1,032,607	322	33	1.4	3.9	65.0	8.8	20.9	0.0	0.1	0.0	24.0	2.2	61.1	5.9	0.3	0.0	6.5	0.0
			Fall		306,342	59	13	0.2	5.7	49.8	21.0	23.3	0.0	0.0	0.0	0.3	5.5	72.5	18.6	3.2	0.0	0.0	0.0
		S6	Late Spring		209,568	298	12	1.2	2.8	90.7	4.8	0.0	0.0	0.4	0.1	4.4	2.7	84.2	3.6	0.0	0.0	4.8	0.2
		30	Summer		948,606	310	31	0.3	4.5	57.8	9.6	27.1	0.0	0.6	0.0	7.0	13.8	67.1	9.8	0.3	0.0	2.2	0.0
			Late Spring		1,168,780	814	23	2.5	6.1	61.1	3.0	26.0	0.2	1.0	0.0	4.9	1.1	68.6	5.1	0.5	0.2	19.4	0.2
Lac de Gras		FF2-1	Summer		1,764,661	511	39	2.3	11.0	20.3	6.0	60.3	0.0	0.0	0.0	41.5	13.9	37.5	4.0	1.4	0.0	1.7	0.0
Lac de Gras			Fall		815,498	344	25	2.7	16.3	36.9	7.8	36.2	0.0	0.1	0.0	16.3	46.0	16.8	7.2	1.2	0.8	11.6	0.0
			Late Spring		1,065,176	1,993	23	1.8	2.6	90.2	4.1	0.4	0.0	0.8	0.0	1.9	1.8	81.0	6.0	0.0	0.0	9.2	0.0
		FF2-2	Summer		1,319,902	567	36	3.3	8.7	23.5	9.9	54.3	0.0	0.4	0.0	66.6	3.3	22.3	6.8	0.7	0.1	0.2	0.0
			Fall		605,545	348	24	2.8	17.2	55.0	9.4	15.7	0.0	0.0	0.0	16.5	22.5	42.0	18.8	0.2	0.0	0.0	0.0
			Late Spring		1,033,300	1,226	17	3.2	3.3	73.6	9.3	9.1	0.0	1.6	0.0	12.3	4.0	53.4	7.1	0.3	0.0	23.0	0.0
		FF2-3	Summer		1,391,157	768	36	5.2	9.9	42.0	6.6	36.1	0.2	0.0	0.0	58.8	4.4	30.2	2.2	0.5	2.7	1.3	0.0
	Far-field 2		Fall		759,109	351	26	5.0	8.5	44.3	8.5	33.5	0.0	0.3	0.0	35.1	16.6	29.8	12.8	1.1	0.0	4.6	0.0
			Late Spring		1,096,940	1,006	25	4.8	6.8	65.0	11.1	10.2	2.1	0.0	0.0	1.9	6.8	55.9	14.9	0.3	20.1	0.0	0.1
		FF2-4	Summer	Rep 1	1,104,847	348	38	2.4	16.6	5.2	1.6	74.2	0.0	0.0	0.0	57.1	11.7	25.2	3.3	1.5	1.3	0.0	0.0
		112-4	Summer	Rep 2	1,507,572	1,032	36	5.7	5.5	32.2	6.8	49.6	0.0	0.2	0.0	67.0	2.7	25.9	2.6	0.4	0.0	1.4	0.0
			Fall		1,514,140	718	28	1.6	11.7	22.3	4.9	59.5	0.0	0.0	0.0	11.6	36.0	13.9	36.9	1.6	0.0	0.0	0.0
			Late Spring	Rep 1	918,388	1,086	29	2.1	6.9	75.3	4.7	9.8	0.0	1.2	0.1	10.5	2.3	52.6	6.6	0.6	0.0	27.3	0.2
		FF2-5	Late Spring	Rep 2	1,041,809	1,547	21	2.2	5.3	78.2	5.7	7.9	0.0	0.7	0.1	10.3	1.0	52.5	4.9	0.1	0.0	31.1	0.2
		112-3	Summer		1,375,229	2,006	42	13.9	8.9	31.9	5.8	39.0	0.0	0.4	0.0	86.8	1.3	10.1	1.0	0.2	0.1	0.4	0.1
			Fall		1,426,239	494	25	2.0	12.6	38.5	5.8	41.0	0.1	0.0	0.0	12.8	43.2	23.8	17.0	1.9	1.3	0.0	0.0

Table A-7 Summary of Phytoplankton Community Data for Lakes in the Jay Project Area, 2014

Notes: Samples were analyzed by EcoAnalysts, Inc. Total abundance and biomass values are rounded to the nearest whole number.

cells/L = number of cells per litre; mg/m³ = milligrams per cubic metre; No. = number; % = percent; Late Spring = July; Summer = August; Fall = September.

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Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.34	1.52
		Cladagara	Chydorus sphaericus	0.04	0.07
		Cladocera	Daphnia longiremis	3.74	14.29
			Holopedium gibberum	1.65	12.85
			Conochilus unicornis	1.26	0.03
			Filinia terminalis	1.10	0.01
			Kellicottia longispina	35.92	0.37
		Rotifera	Keratella cochlearis	8.98	0.03
July	Ab-1		Gastropus stylifer	0.79	0.03
			Ploesoma truncatum	0.16	0.00
			Polyarthra remata	0.16	0.01
		Copepoda - nauplii	Nauplii	4.73	1.89
		Calanoida	Epischura sp.	0.04	0.78
			Copepodites	0.39	0.95
		Cyclopoida	Cyclops strenuus	0.04	0.68
			Diacyclops thomasi	1.22	7.39
			Copepodites	2.36	3.02
		Cladocera	Bosmina longirostris	0.49	0.43
			Chydorus sphaericus	0.03	0.03
			Daphnia longiremis	1.43	3.59
			Holopedium gibberum	1.16	9.02
			Conochilus unicornis	2.37	0.04
			Filinia terminalis	0.32	0.00
			Kellicottia longispina	27.09	0.28
la de c	A = 1	Datifara	Keratella cochlearis	13.38	0.04
July	Ac-1	Rotifera	Ascomorpha ecaudis	0.11	0.00
			Gastropus stylifer	0.97	0.03
			Ploesoma truncatum	0.11	0.01
			Polyarthra remata	1.62	0.08
		Copepoda - nauplii	Nauplii	2.48	0.99
		Copepodites	Copepodites	0.35	0.87
		Cuelopaida	Diacyclops thomasi	1.35	8.84
		Cyclopoida	Copepodites	2.19	5.30



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.20	1.44
		Cladocera	Daphnia longiremis	3.97	6.28
			Holopedium gibberum	2.27	17.63
			Conochilus unicornis	1.64	0.09
			Filinia terminalis	0.38	0.00
			Kellicottia longispina	24.57	0.25
		Rotifera	Keratella cochlearis	9.20	0.03
L.L.	A = 1		Gastropus stylifer	1.64	0.06
July	Ac-4		Polyarthra remata	0.88	0.04
			Synchaeta stylata	0.13	0.01
		Copepoda - nauplii	Nauplii	5.17	2.07
		Oslavsida	Epischura sp.	0.06	0.36
		Calanoida	Copepodites	0.32	0.52
		Cyclopoida	Cyclops strenuus	0.06	0.63
			Diacyclops thomasi	2.39	13.02
			Copepodites	2.39 3.91	7.61
			Bosmina longirostris	0.33	0.18
		Cladocera	Chydorus sphaericus	0.07	0.03
			Daphnia longiremis	1.19	1.86
			Holopedium gibberum	0.35	2.09
			Collotheca pelagica	0.09	0.00
			Conochilus unicornis	3.37	0.07
			Filinia terminalis	0.56	0.01
			Asplanchna priodonta	0.09	0.12
			Kellicottia longispina	37.20	0.38
L.L.	A . 7	Rotifera	Keratella cochlearis	10.21	0.03
July	Ac-7		Gastropus stylifer	1.12	0.04
			Ploesoma hudsoni	0.09	0.04
			Ploesoma truncatum	0.19	0.01
			Polyarthra remata	2.44	0.12
			Trichotria tetractis	0.09	0.00
		Copepoda - nauplii	Nauplii	3.94	1.57
			Epischura sp.	0.02	0.18
		Calanoida	Copepodites	0.63	1.25
		Quality in	Diacyclops thomasi	1.22	6.89
		Cyclopoida	Copepodites	2.39	3.88



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	0.23	0.16
		Cladocera	Daphnia longiremis	0.59	1.08
			Holopedium gibberum	0.26	1.61
			Conochilus unicornis	1.14	0.02
			Filinia terminalis	4.34	0.04
			Kellicottia longispina	13.60	0.14
			Keratella cochlearis	4.65	0.02
		Datifana	Keratella hiemalis	0.03	0.00
July	Ad-1	Rotifera	Gastropus stylifer	0.31	0.01
			Ploesoma hudsoni	0.07	0.01
			Ploesoma truncatum	0.17	0.01
			Polyarthra remata	1.01	0.05
			Synchaeta stylata	0.03	0.00
		Copepoda - nauplii	Nauplii	2.46	0.99
		Copepodites	Copepodites	0.07	0.26
		Qualanaida	Diacyclops thomasi	0.62	4.04
		Cyclopoida	Copepodites	2.05	4.27
		Cladocera	Bosmina longirostris	0.84	0.78
			Chydorus sphaericus	0.16	0.13
			Daphnia longiremis	2.36	6.26
			Holopedium gibberum	1.48	9.45
			Conochilus unicornis	3.04	0.08
			Filinia terminalis	0.48	0.01
			Asplanchna priodonta	0.16	0.18
			Kellicottia longispina	22.73	0.23
July	Ae-1	Rotifera	Keratella cochlearis	10.72	0.03
			Gastropus stylifer	0.16	0.00
			Ploesoma hudsoni	0.08	0.01
			Ploesoma truncatum	0.24	0.01
			Polyarthra remata	2.00	0.10
		Copepoda - nauplii	Nauplii	2.64	1.06
		Copepodites	Copepodites	0.52	0.81
		Quelesside	Diacyclops thomasi	1.20	6.69
		Cyclopoida	Copepodites	2.52	2.89



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	9.49	6.14
		Cladocera	Daphnia longiremis	2.71	4.23
			Holopedium gibberum	4.07	19.04
			Collotheca sp.	0.45	0.01
			Conochilus unicornis	6.10	0.14
			Filinia terminalis	4.18	0.04
			Asplanchna priodonta	0.23	0.25
			Kellicottia longispina	31.08	0.32
		Rotifera	Keratella cochlearis	10.06	0.03
A	A - 4		Ascomorpha ecaudis	0.11	0.00
August	Aa-1		Gastropus stylifer	2.03	0.07
			Ploesoma hudsoni	0.11	0.02
			Ploesoma truncatum	0.11	0.00
			Polyarthra remata	0.90	0.04
		Copepoda - nauplii	Nauplii	5.76	2.31
		Calanoida	Leptodiaptomus minutus	0.11	1.15
			Leptodiaptomus pribilofensis	0.11	0.87
			Epischura sp.	0.11	1.99
		Qualar aida	Diacyclops thomasi	1.70	9.26
		Cyclopoida	Copepodites	3.73	5.99
			Bosmina longirostris	3.00	3.55
		Cladocera	Daphnia longiremis	6.28	9.18
			Holopedium gibberum	3.23	27.55
			Collotheca pelagica	1.04	0.00
			Conochilus unicornis	3.80	0.09
			Filinia terminalis	0.12	0.00
			Kellicottia longispina	18.44	0.19
			Keratella cochlearis	2.42	0.01
		Rotifera	Ascomorpha ecaudis	0.12	0.00
August	Ab-1		Ascomorpha ovalis	0.12	0.00
			Gastropus stylifer	0.81	0.03
			Monostyla lunaris	0.12	0.00
			Polyarthra major	1.27	0.06
		Copepoda - nauplii	Nauplii	4.27	1.71
			Leptodiaptomus minutus	0.12	1.20
		Calanoida	Copepodites	0.06	0.39
			Diacyclops thomasi	2.19	13.00
		Cyclopoida	Copepodites	1.90	4.46



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.67	0.92
		Cladocera	Daphnia longiremis	5.83	10.80
			Holopedium gibberum	3.65	25.71
			Collotheca pelagica	0.64	0.01
			Conochilus unicornis	7.43	0.16
			Filinia terminalis	0.26	0.00
			Asplanchna priodonta	0.26	0.50
		Datifara	Kellicottia longispina	25.23	0.26
		Rotifera	Keratella cochlearis	2.18	0.01
August	Ac-1		Ascomorpha ovalis	0.26	0.00
			Gastropus stylifer	0.64	0.03
			Monostyla lunaris	0.13	0.00
			Polyarthra major	1.02	0.05
		Copepoda - nauplii	Nauplii	7.94	3.18
		Calanoida	Leptodiaptomus minutus	0.32	2.71
			Epischura nevadensis	0.06	1.64
			Copepodites	0.06	0.44
		Quality side	Diacyclops thomasi	1.54	10.17
		Cyclopoida	Copepodites	1.41	2.09
			Bosmina longirostris	1.13	0.95
		Cladocera	Eubosmina longispina	0.14	0.21
			Daphnia longiremis	13.00	19.00
			Holopedium gibberum	3.11	28.11
			Collotheca pelagica	0.85	0.00
			Conochilus unicornis	5.51	0.11
			Asplanchna priodonta	0.28	0.32
		Rotifera	Kellicottia longispina	26.42	0.27
August	Ac-4		Keratella cochlearis	2.83	0.01
			Gastropus stylifer	2.12	0.08
			Polyarthra major	0.99	0.05
		Copepoda - nauplii	Nauplii	11.59	4.63
			Leptodiaptomus minutus	0.28	3.18
		Calanoida	Leptodiaptomus pribilofensis	0.14	1.83
			Copepodites	0.14	0.80
		Qualizzaida	Diacyclops thomasi	4.66	26.37
		Cyclopoida	Copepodites	5.93	11.02



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.79	1.81
		Cladocera	Daphnia longiremis	3.63	6.16
			Holopedium gibberum	1.21	8.92
			Collotheca pelagica	0.63	0.00
			Conochilus unicornis	8.07	0.15
			Filinia terminalis	0.27	0.00
			Asplanchna priodonta	0.27	0.29
			Kellicottia longispina	19.19	0.20
		Rotifera	Keratella cochlearis	0.72	0.00
. .			Ascomorpha ovalis	0.09	0.00
August	Ac-7		Gastropus stylifer	1.17	0.05
			Ploesoma hudsoni	0.09	0.01
			Polyarthra major	0.27	0.01
		Copepoda - nauplii	Nauplii	5.83	2.33
		Calanoida	Leptodiaptomus minutus	0.31	2.61
			Leptodiaptomus pribilofensis	0.04	0.43
			Epischura nevadensis	0.13	4.00
			Copepodites	0.22	1.54
		Quality side	Diacyclops thomasi	1.48	9.61
		Cyclopoida	Copepodites	1.79 3.63 1.21 0.63 8.07 0.27 0.27 0.172 0.09 1.17 0.09 0.27 5.83 0.31 0.04 0.13 0.22 1.48 1.79 5.44 3.01 0.38 6.79 0.51 31.38 6.40 0.13 1.28 0.26 2.18 9.48 0.26 2.17	3.08
			Bosmina longirostris	1.79	1.28
		Cladocera	Daphnia longiremis	5.44	11.42
			Holopedium gibberum	3.01	22.48
			Collotheca pelagica	0.38	0.00
			Conochilus unicornis	6.79	0.13
			Asplanchna priodonta	0.51	0.34
			Kellicottia longispina	31.38	0.32
		Rotifera	Keratella cochlearis	6.40	0.02
August	Ad-1		Ascomorpha ovalis	0.13	0.00
			Gastropus stylifer	1.28	0.05
			Ploesoma truncatum	0.26	0.01
			Polyarthra major		0.11
		Copepoda - nauplii	Nauplii	9.48	3.79
			Leptodiaptomus minutus	0.26	2.37
		Calanoida	Copepodites	0.26	1.15
			Diacyclops thomasi		10.61
		Cyclopoida	Copepodites	3.71	4.56

 Table A-8
 Zooplankton Abundance and Biomass in Lac du Sauvage, 2014



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	4.58	3.91
		Cladocera	Daphnia longiremis	13.56	18.97
			Holopedium gibberum	11.02	84.53
			Collotheca pelagica	0.85	0.00
			Conochilus unicornis	14.41	0.24
			Kellicottia longispina	31.87	0.33
		Rotifera	Keratella cochlearis	1.19	0.00
		Rolliela	Ascomorpha ovalis	0.17	0.00
August	Ae-1		Gastropus stylifer	1.53	0.06
August	Ae-1		Ploesoma truncatum	0.17	0.01
			Polyarthra major	1.36	0.07
		Copepoda - nauplii	Nauplii	7.63	3.05
			Leptodiaptomus minutus	0.51	4.90
		Colonaida	Leptodiaptomus pribilofensis	0.51	4.08
		Calanoida	Epischura nevadensis	0.17	3.37
			Copepodites	0.51	1.81
		Qualan sida	Diacyclops thomasi	4.07	24.51
		Cyclopoida	Copepodites	0.85	1.77
		Cladocera	Bosmina longirostris	3.73	5.72
			Eubosmina longispina	0.47	1.41
			Chydorus sphaericus	0.38	0.35
			Daphnia longiremis	1.04	2.87
			Holopedium gibberum	0.05	0.23
			Collotheca pelagica	0.19	0.00
			Conochilus unicornis	11.06	0.27
			Filinia terminalis	3.69	0.03
			Asplanchna priodonta	9.92	6.29
Quarteration	A = 4	Datifana	Kellicottia longispina	14.56	0.15
September	Aa-1	Rotifera	Keratella cochlearis	5.77	0.02
			Ascomorpha ecaudis	0.28	0.00
			Ascomorpha ovalis	1.51	0.02
			Gastropus stylifer	0.76	0.03
			Polyarthra major	0.28	0.01
		Copepoda - nauplii	Nauplii	7.28	2.91
			Leptodiaptomus minutus	0.09	0.84
		Calanoida	Copepodites	0.05	0.19
			Diacyclops thomasi	2.79	14.56
		Cyclopoida	Copepodites	4.68	7.25

 Table A-8
 Zooplankton Abundance and Biomass in Lac du Sauvage, 2014



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
Period			Bosmina longirostris	3.58	5.63
			Eubosmina longispina	0.11	0.39
		Cladocera	Acroperus harpae	0.04	0.07
			Chydorus sphaericus	0.18	0.14
			Daphnia longiremis	0.21	0.36
			Collotheca pelagica	0.70	0.00
			Conochilus unicornis	12.23	0.26
			Filinia terminalis	0.42	0.00
September	Ab-1		Asplanchna priodonta	3.37	2.52
		Rotifera	Kellicottia longispina	10.54	0.11
			Keratella cochlearis	3.37	0.01
			Ascomorpha ovalis	0.56	0.01
			Gastropus stylifer	0.56	0.02
			Polyarthra major	1.41	0.07
		Copepoda - nauplii	Nauplii	16.16	6.47
		Quality slide	Diacyclops thomasi	1.55	9.31
		Cyclopoida	Copepodites	3.55	3.90
			Bosmina longirostris	1.38	2.52
		Cladocera	Eubosmina longispina	0.28	1.02
			Chydorus sphaericus	0.35	0.33
			Daphnia longiremis	0.57	1.26
			Asplanchna priodonta	1.55	1.33
			Collotheca pelagica	0.57	0.00
			Conochilus unicornis	12.43	0.26
			Filinia terminalis	0.28	0.00
			Kellicottia longispina	7.91	0.08
		Rotifera	Keratella cochlearis	2.40	0.01
September	Ac-1 ^(a)		Ascomorpha ecaudis	0.14	0.00
			Ascomorpha ovalis	0.57	0.01
			Gastropus stylifer	0.28	0.01
			Monostyla lunaris	0.14	0.00
			Polyarthra major	2.83	0.14
		Copepoda - nauplii	Nauplii	13.85	5.54
			Leptodiaptomus minutus	0.04	0.50
		Calanoida	Epischura sp.	0.04	1.18
			Cyclops strenuus	0.04	0.27
		Cyclopoida	Diacyclops thomasi	1.77	8.54
			Copepodites	5.16	6.71



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.47	1.25
		Cladocera	Eubosmina longispina	0.28	0.97
		Cladocera	Chydorus sphaericus	0.28	0.25
			Daphnia longiremis	0.65	1.07
			Asplanchna priodonta	3.05	2.31
			Collotheca pelagica	1.02	0.01
			Conochilus unicornis	10.51	0.23
			Filinia terminalis	0.11	0.00
September	Ac-1 ^(b)	Rotifera	Kellicottia longispina	8.93	0.09
			Keratella cochlearis	3.05	0.01
			Ascomorpha ovalis	0.34	0.00
			Gastropus stylifer	0.79	0.03
			Polyarthra major	2.15	0.11
		Copepoda - nauplii	Nauplii	13.11	5.24
		Cyclopoida	Epischura nevadensis	0.06	1.66
			Diacyclops thomasi	1.95	9.38
			Copepodites	4.07	4.25
		Cladocera	Bosmina longirostris	1.19	1.15
			Eubosmina longispina	0.27	0.93
			Chydorus sphaericus	0.15	0.08
			Daphnia longiremis	1.34	3.06
			Collotheca pelagica	0.54	0.00
			Conochilus unicornis	10.84	0.24
			Filinia terminalis	0.08	0.00
Contombon	A = 4	Rotifera	Asplanchna priodonta	2.46	2.24
September	Ac-4	Rolliela	Kellicottia longispina	5.92	0.06
			Keratella cochlearis	1.77	0.01
			Gastropus stylifer	0.15	0.01
			Polyarthra major	0.46	0.02
		Copepoda - nauplii	Nauplii	15.98	6.39
		Calanoida	Epischura nevadensis	0.04	0.92
		Cuolonaida	Diacyclops thomasi	2.92	14.65
		Cyclopoida	Copepodites	4.23	4.77



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.61	1.65
			Eubosmina longispina	0.12	0.65
		Cladocera	Acroperus harpae	0.04	0.07
			Chydorus sphaericus	0.04	0.04
			Daphnia longiremis	1.31	2.07
			Collotheca pelagica	0.61	0.00
			Conochilus unicornis	10.30	0.22
			Asplanchna priodonta	1.92	1.89
September	Ac-7		Kellicottia longispina	5.53	0.06
		Rotifera	Keratella cochlearis	2.00	0.01
			Ascomorpha ovalis	0.38	0.00
			Gastropus stylifer	0.69	0.03
			Polyarthra major	0.46	0.02
		Copepoda - nauplii	Nauplii	14.83	5.93
		Calanoida	Epischura nevadensis	0.04	0.85
		Cyclopoida	Diacyclops thomasi	1.96	10.29
			Copepodites	3.84	5.10
		Cladocera	Bosmina longirostris	0.47	0.38
			Eubosmina longispina	0.09	0.42
			Chydorus sphaericus	0.09	0.07
			Daphnia longiremis	1.04	1.48
			Collotheca pelagica	1.04	0.00
			Conochilus unicornis	17.77	0.28
			Filinia terminalis	0.28	0.00
			Ascomorpha ovalis	0.28	0.00
September	Ad-1		Asplanchna priodonta	3.50	2.14
		Rotifera	Kellicottia longispina	5.58	0.06
			Keratella cochlearis	1.89	0.01
			Gastropus stylifer	1.23	0.05
			Monostyla lunaris	0.09	0.00
			Polyarthra major	1.23	0.06
		Copepoda - nauplii	Nauplii	15.69	6.28
			Diacyclops thomasi	3.73	15.48
		Cyclopoida	Copepodites	5.67	7.68

 Table A-8
 Zooplankton Abundance and Biomass in Lac du Sauvage, 2014



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	1.71	1.83
		Cladocera	Eubosmina longispina	0.68	3.28
			Daphnia longiremis	2.44	6.82
			Collotheca pelagica	2.34	0.01
			Conochilus unicornis	19.34	0.32
		Rotifera	Filinia terminalis	0.20	0.00
			Asplanchna priodonta	3.52	4.10
			Kellicottia longispina	13.09	0.13
September	Ae-1		Keratella cochlearis	1.17	0.00
			Gastropus stylifer	0.20	0.01
			Polyarthra major	1.56	0.08
		Copepoda - nauplii	Nauplii	21.88	8.75
			Leptodiaptomus minutus	0.05	0.38
		Calanoida	Leptodiaptomus pribilofensis	0.10	1.30
			Heterocope septentrionalis ^(c)	-	-
		Cualanaida	Diacyclops thomasi	3.91	20.10
		Cyclopoida	Copepodites	7.33	7.70

 Table A-8
 Zooplankton Abundance and Biomass in Lac du Sauvage, 2014

Notes: Samples were analyzed by EcoAnalysts, Inc. Values are rounded to the nearest whole number.

org/L = number of organisms per litre; mg/m³ = milligrams per cubic metre; sp. = species; - = not applicable.

a) Replicate 1.

b) Replicate 2.

c) Presence of *Heterocope septentrionalis* was noted in the fine fraction of this sample, but not enumerated or measured by the taxonomist. This taxon was counted towards taxonomic richness, but abundance and biomass could not be quantified.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	27.61	17.25
		Cladocera	Daphnia longiremis	2.80	6.22
			Holopedium gibberum	1.12	7.43
			Collotheca pelagica	1.12	0.01
			Conochilus unicornis	11.20	0.25
			Asplanchna priodonta	0.16	0.14
			Kellicottia longispina	5.36	0.06
			Keratella cochlearis	1.12	0.00
		Rotifera	Ascomorpha ecaudis	0.08	0.00
August	Af-1	Rolliela	Gastropus stylifer	0.40	0.01
			Lecane mira	0.08	0.00
			Ploesoma hudsoni	0.48	0.09
			Ploesoma truncatum	4.88	0.19
			Polyarthra major	0.48	0.02
			Synchaeta stylata	0.08	0.02
		Copepoda - nauplii	Nauplii	8.08	3.23
		Calanoida	Epischura nevadensis	0.08	2.93
		Cyclopoida	Diacyclops thomasi	0.16	1.80
		Cyclopolua	Copepodites	0.32	0.84
		Cladocera	Daphnia longiremis	8.48	7.58
		Claubcera	Holopedium gibberum	0.14	1.00
			Collotheca sp.	0.42	0.08
			Conochilus unicornis	14.13	0.34
			Asplanchna priodonta	5.93	4.09
		Rotifera	Kellicottia longispina	8.48	0.09
September	Af-1	Rolliera	Keratella cochlearis	7.63	0.03
			Ascomorpha ovalis	3.11	0.04
			Gastropus stylifer	3.67	0.14
			Polyarthra major	2.68	0.13
		Copepoda - nauplii	Nauplii	1.55	0.62
		Calanoida	Leptodiaptomus minutus	0.14	0.80
		Cyclopoida	Copepodites	4.66	6.55

Table A-9 Zooplankton Abundance and Biomass in Duchess Lake, 2014

Notes: Samples were analyzed by EcoAnalysts, Inc.

org/L = number of organisms per litre; mg/m³ = milligrams per cubic metre; sp. = species.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
		Cladocera	Daphnia longiremis	0.10	0.18
		Claudcera	Holopedium gibberum	0.29	1.83
			Conochilus unicornis	22.16	0.44
			Asplanchna priodonta	0.35	0.48
			Kellicottia longispina	28.23	0.29
		Datifara	Keratella cochlearis	2.45	0.01
hub e	S2	Rotifera	Gastropus stylifer	1.17	0.05
July	52		Cephalodella gibba	0.12	0.00
			Polyarthra remata	2.45	0.12
			Synchaeta stylata	0.35	0.03
		Copepoda - nauplii	Nauplii	0.93	0.37
		Copepodites	Copepodites	0.66	3.38
		Quelenside	Cyclops strenuus	1.01	14.39
		Cyclopoida	Copepodites	2.53	5.51
		Olesteere	Daphnia longiremis	0.06	0.27
		Cladocera	Holopedium gibberum	0.02	0.20
		Rotifera	Collotheca sp.	0.06	0.00
			Conochilus unicornis	3.52	0.07
			Kellicottia longispina	4.74	0.05
			Keratella cochlearis	2.69	0.01
			Keratella hiemalis	0.51	0.02
l. l.	00		Ascomorpha ecaudis	0.06	0.00
July	53		Gastropus stylifer	0.26	0.01
	S3		Polyarthra remata	1.09	0.05
			Trichotria tetractis	0.06	0.00
		Copepoda - nauplii	Nauplii	2.50	1.00
		Ostansida	Epischura sp.	0.02	0.41
		Calanoida	Copepodites	1.25	7.32
			Cyclops strenuus	0.45	4.89
		Cyclopoida	Copepodites	3.20	11.54
			Chydorus sphaericus	0.01	0.01
		Cladocera	Daphnia middendorffiana	0.08	0.54
			Holopedium gibberum	0.08	0.33
			Conochiloides natans	0.03	0.01
July	S5 ^(a)		Conochilus unicornis	2.33	0.06
		Detifere	Kellicottia longispina	4.14	0.04
		Rotifera	Keratella cochlearis	0.45	0.00
			Keratella hiemalis	0.26	0.01
			Ascomorpha ecaudis	0.05	0.00



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Gastropus stylifer	0.10	0.00
			Polyarthra remata	0.42	0.02
		Copepoda - nauplii	Nauplii	1.15	0.46
			Diaptomidae	0.01	0.21
		Calanoida	Epischura sp.	0.01	0.31
		Calanoida	Heterocope septentrionalis	0.01	0.94
			Copepodites	1.49	4.96
		Quelenside	Cyclops strenuus	0.47	5.00
		Cyclopoida	Copepodites	0.89	2.62
			Daphnia longiremis	0.01	0.02
		Cladocera	Daphnia middendorffiana	0.04	0.23
			Holopedium gibberum	0.03	0.28
			Conochilus unicornis	2.59	0.07
			Kellicottia longispina	4.37	0.04
			Keratella cochlearis	0.58	0.00
			Keratella hiemalis	0.26	0.01
	o - (b)		Ascomorpha ecaudis	0.10	0.00
July	S5 ^(b)		Gastropus stylifer	(org/L) 0.10 0.42 1.15 0.01 0.01 0.01 1.49 0.47 0.89 0.01 0.04 0.03 2.59 4.37 0.58 0.26	0.01
			Polyarthra remata		0.01
		Copepoda - nauplii	Nauplii	1.52	0.61
			Leptodiaptomus sicilis	0.03	0.17
			Heterocope septentrionalis	1.49 0.47 0.89 0.01 0.04 0.03 2.59 4.37 0.58 0.26 0.10 0.18 0.16 1.52 0.03 0.01 2.12 0.52 1.01 0.06 0.02 0.04 3.85 4.53 0.40 0.24 0.08	0.35
			Copepodites	2.12	5.79
			Cyclops strenuus	0.52	5.43
		Cyclopoida	Copepodites	1.01	3.19
			Daphnia middendorffiana	0.06	0.76
		Cladocera	Holopedium gibberum	0.02	0.19
			Collotheca sp.	0.04	0.00
			Conochilus unicornis	(org/L) 0.10 0.42 1.15 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.01 0.03 2.59 4.37 0.58 0.26 0.10 0.18 0.16 1.52 0.03 0.01 2.12 0.52 1.01 0.06 0.02 0.04 3.85 4.53 0.40 0.28 2.47 0.06 0.02	0.11
			Kellicottia longispina		0.05
		Detifere	Keratella cochlearis	0.40	0.00
le de s	00	Rotifera	Keratella hiemalis	0.24	0.01
July	S6		Ascomorpha ecaudis	0.08	0.00
			Polyarthra remata	0.20	0.01
			Gastropus stylifer	0.28	0.01
		Copepoda - nauplii	Nauplii	2.47	0.99
			Leptodiaptomus sicilis	0.06	0.76
		Calanoida	Epischura sp.	0.02	0.39
			Copepodites	1.92	5.51



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
		Cyclopaida	Cyclops strenuus	0.35	3.91
		Cyclopoida	Copepodites	1.02	2.88
		Cladagara	Daphnia longiremis	(org/L) 0.35 1.02 0.12 1.46 0.09 0.26 4.55 12.16 0.87 0.70 2.19 5.34 0.12 0.17 0.03 - 1.78 0.06 0.23 1.92 0.08 0.09 1.48 4.40 0.09 1.48 4.40 0.09 0.49 0.63 2.32 0.07 0.08 0.02 0.05 0.93 0.01 0.06	0.64
		Cladocera	Holopedium gibberum	1.46	8.35
			Bdelloidea	0.09	0.00
			Collotheca pelagica	0.26	0.00
			Conochilus unicornis	4.55	0.10
		Rotifera	Kellicottia longispina	12.16	0.12
			Keratella cochlearis	0.87	0.00
			Gastropus stylifer	0.70	0.03
A	60		Polyarthra major	2.19	0.11
August	S2	Copepoda - nauplii	Nauplii	5.34	2.13
			Leptodiaptomus minutus	0.12	1.14
			Leptodiaptomus pribilofensis	0.17	2.43
		Calanoida	Epischura nevadensis	(org/L) 0.35 1.02 0.12 1.46 0.09 0.26 4.55 12.16 0.87 0.70 2.19 5.34 0.12 0.17 0.03 - 1.78 0.06 0.23 1.92 0.08 0.09 1.48 4.40 0.09 1.48 4.40 0.09 0.49 0.63 2.32 0.07 0.08 0.02 0.93 0.01	0.54
			Heterocope septentrionalis ^(c)		-
			Copepodites	1.78	14.05
			Acanthocyclops robustus	(org/L) 0.35 1.02 0.12 1.46 0.09 0.26 4.55 12.16 0.87 0.70 2.19 5.34 0.12 0.17 0.03 - 1.78 0.06 0.23 1.92 0.08 0.09 1.48 4.40 0.09 1.48 4.40 0.09 0.49 0.63 2.32 0.07 0.08 0.09 0.49 0.63 2.32 0.07 0.08 0.09 0.49 0.63 2.32 0.07 0.08 0.09 0.49 0.63 2.32 0.07 0.08 0.09 0.49 0.63 2.32 0.07 0.08 0.09 0.49 0.63 2.32 0.07 0.08 0.09 0.49 0.63 2.32 0.07 0.08 0.02 0.05 0.93 0.01 0.06 0.93 0.01 0.06 0.02 0.05 0.93 0.01 0.06	0.58
		Cyclopoida	Cyclops strenuus	0.23	2.53
			Copepodites	1.92	7.47
		Cladocera	Holopedium gibberum	0.08	0.69
			Collotheca pelagica	0.09	0.00
			Conochilus unicornis	1.48	0.04
		Datifana	Kellicottia longispina	4.40	0.05
		Rotifera	Keratella cochlearis	0.09	0.00
			Gastropus stylifer	0.49	0.02
			Polyarthra major	0.63	0.03
A		Copepoda - nauplii	Nauplii	2.32	0.93
August	S3 ^(a)		Leptodiaptomus minutus	(org/L) 0.35 1.02 0.12 1.46 0.09 0.26 4.55 12.16 0.87 0.70 2.19 5.34 0.12 0.17 0.03 - 1.78 0.06 0.23 1.92 0.08 0.09 1.48 4.40 0.09 1.48 4.40 0.09 0.49 0.63 2.32 0.07 0.08 0.02 0.05 0.93 0.01 0.06	0.75
			Leptodiaptomus pribilofensis		1.47
		Calanoida	Epischura nevadensis		0.64
			Heterocope septentrionalis		4.91
			Copepodites	0.93	7.01
			Acanthocyclops robustus	0.01	0.16
		Cyclopoida	Cyclops strenuus	0.06	0.80
		-	Copepodites	1.12	5.10



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
		Cladocera	Holopedium gibberum	0.07	0.48
			Collotheca pelagica	0.12	0.00
			Conochilus unicornis	(org/L) 0.07	0.04
		Datifara	Kellicottia longispina	4.05	0.04
		Rotifera	Keratella cochlearis	0.12	0.00
			Gastropus stylifer	0.47	0.02
			Polyarthra major	1.05	0.05
August	S3 ^(b)	Copepoda - nauplii	Nauplii	(org/L) 0.07 0.12 1.41 4.05 0.12 0.47 1.05 2.20 0.07 0.06 0.01 0.91 0.00 0.13 1.14 0.13 1.14 0.01 0.91 0.00 0.13 1.14 0.13 1.14 0.13 0.04 0.05 1.99 3.68 0.13 0.08 0.36 1.84 0.02 0.01 0.05 2.25 0.19 0.61 0.08 0.04 0.07 1.69 3.64	0.88
			Conochilus unicornisKellicottia longispinaKeratella cochlearisGastropus styliferPolyarthra majorNaupliiLeptodiaptomus minutusLeptodiaptomus pribilofensisHeterocope septentrionalisCopepoditesAcanthocyclops robustusCyclops strenuusCopepoditesDaphnia middendorffianaHolopedium gibberumConochilus unicornisKeratella cochlearisGastropus styliferPolyarthra majorNaupliiLeptodiaptomus pribilofensisCopepoditesDaphnia middendorffianaHolopedium gibberumCollotheca pelagicaConochilus unicornisKeratella cochlearisGastropus styliferPolyarthra majorNaupliiLeptodiaptomus minutusLeptodiaptomus pribilofensisEpischura nevadensisHeterocope septentrionalisCopepoditesCopepodites	0.07	0.71
	et S3 ^(b) Copepoda - nauplii Calanoida Cyclopoida Cladocera Rotifera		Leptodiaptomus pribilofensis	0.06	0.93
		Calanoida	Heterocope septentrionalis	0.01	1.17
			Copepodites	0.91	6.97
			Acanthocyclops robustus	0.00	0.00
			Cyclops strenuus	0.13	2.01
		Cyclopoida	Copepodites	1.14	5.32
			Daphnia middendorffiana	0.13	3.21
		Cladocera	Holopedium gibberum	0.04	0.49
		Rotifera	Collotheca pelagica	0.05	0.00
			Conochilus unicornis	1.99	0.06
			Kellicottia longispina	3.68	0.04
			Keratella cochlearis	0.13	0.00
			Gastropus stylifer	0.08	0.00
. .	0.5		Polyarthra major	0.36	0.02
August	\$5	Copepoda - nauplii	Nauplii	1.84	0.74
			Leptodiaptomus minutus	0.02	0.15
			Leptodiaptomus pribilofensis	0.02	0.33
		Calanoida	Epischura nevadensis	0.01	0.16
			Heterocope septentrionalis	0.05	4.70
			Copepodites	2.25	13.29
		Quality side	Cyclops strenuus	0.19	2.14
		Cyclopoida	Copepodites	0.61	2.35
		Cladarara	Daphnia middendorffiana	0.08	2.34
		Cladocera	Holopedium gibberum	0.04	0.35
			Collotheca pelagica	0.07	0.00
August	80		Conochilus unicornis	1.69	0.05
August	S6	Datif	Kellicottia longispina	3.64	0.04
		Rotifera	Keratella cochlearis	0.19	0.00
			Keratella hiemalis	0.05	0.00
			Gastropus stylifer	0.05	0.00



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Polyarthra major	0.16	0.01
		Copepoda - nauplii	Nauplii	3.10	1.24
			Leptodiaptomus minutus	0.01	0.15
		Calanoida	Leptodiaptomus pribilofensis	0.04	0.47
		Calanoida	Heterocope septentrionalis	0.02	1.72
			Copepodites	1.69	11.06
		Quelenside	Cyclops strenuus	0.20	2.23
		Cyclopoida	Copepodites	1.04	4.33
			Daphnia longiremis	0.04	0.41
		Cladocera	Daphnia middendorffiana	0.02	0.40
			Holopedium gibberum	0.63	8.54
			Collotheca sp.	0.36	0.06
			Kellicottia longispina	7.65	0.08
			Keratella cochlearis	1.12	0.00
		Rotifera	Ascomorpha ovalis	0.04	0.00
			Gastropus stylifer	(org/L) 0.16 3.10 0.01 0.04 0.02 1.69 0.20 1.04 0.02 0.63 0.36 7.65 1.12	0.01
			Monostyla lunaris	0.07	0.00
			Polyarthra major	0.72	0.04
September	S2	Copepoda - nauplii	Nauplii	5.30	2.12
			Leptodiaptomus ashlandi	0.04	0.29
			Leptodiaptomus minutus	0.34	2.78
			Leptodiaptomus pribilofensis	0.43	5.78
		Calanoida	Leptodiaptomus sicilis	0.78	9.97
			Epischura nevadensis	0.02	0.46
			Heterocope septentrionalis	0.07	5.76
			Copepodites	0.23	1.75
			Cyclops strenuus	0.14	1.20
		Cyclopoida	Copepodites	1.39	5.71
			Daphnia middendorffiana	0.13	4.81
		Cladocera	Holopedium gibberum	0.08	1.32
			Collotheca sp.	0.34	0.06
			Conochilus unicornis	1.24	0.03
		Detil	Kellicottia longispina	6.10	0.06
September	S3	Rotifera	Keratella cochlearis	0.23	0.00
			Gastropus stylifer	0.01 0.04 0.02 1.69 0.20 1.04 0.04 0.02 1.04 0.04 0.02 0.63 0.36 7.65 1.12 0.04 0.32 0.07 0.72 5.30 0.04 0.34 0.43 0.78 0.02 0.07 0.23 0.14 1.39 0.13 0.08 0.34 1.24 6.10 0.23 0.14 0.13 0.07	0.01
			Polyarthra major	0.14	0.01
		Copepoda - nauplii	Nauplii	3.19	1.28
			Leptodiaptomus ashlandi	0.07	0.43
		Calanoida	Leptodiaptomus minutus		2.09



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Leptodiaptomus pribilofensis	0.31	4.01
			Leptodiaptomus sicilis	0.75	10.75
			Epischura nevadensis	0.08	2.13
			Heterocope septentrionalis	0.07	5.91
			Copepodites	0.17	1.15
		Quelenside	Cyclops strenuus	0.04	0.33
		Cyclopoida	Copepodites	0.88	3.43
		Cladagara	Daphnia middendorffiana	0.43	21.05
		Cladocera	Holopedium gibberum	0.02	0.21
		Rotifera	Collotheca sp.	0.05	0.01
			Conochilus unicornis	0.27	0.01
			Kellicottia longispina	1.48	0.02
			Keratella cochlearis	0.02	0.00
			Gastropus stylifer	0.19	0.01
			Polyarthra major	0.03	0.00
September	S5	Copepoda - nauplii	Nauplii	2.36	0.94
			Leptodiaptomus ashlandi	0.06	0.31
			Leptodiaptomus minutus	0.35	2.20
		Colonaida	Leptodiaptomus pribilofensis	0.47	5.09
		Calanoida	Leptodiaptomus sicilis	1.14	10.15
			Heterocope septentrionalis	0.04	4.02
			Copepodites	0.82	4.83
		Quelenside	Cyclops strenuus	0.11	1.16
		Cyclopoida	Copepodites	0.76	2.74

Notes: Samples were analyzed by EcoAnalysts, Inc. Values are rounded to the nearest whole number.

org/L = number of organisms per litre; mg/m³ = milligrams per cubic metre; sp. = species; - = not applicable.

a) Replicate 1.

b) Replicate 2.

c) Presence of *Heterocope septentrionalis* was noted in the fine fraction of this sample, but not enumerated or measured by the taxonomist. This taxon was counted towards taxonomic richness, but abundance and biomass could not be quantified.



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
	Sampling Period Station July FF2-1 July FF2-2		Chydorus sphaericus	0.01	0.01
		Cladocera	Daphnia longiremis	0.01	0.19
			Holopedium gibberum	0.04	0.30
			Collotheca sp.	0.06	0.00
			Conochiloides natans	0.17	0.02
			Conochilus unicornis	1.17	0.03
			Kellicottia longispina	13.19	0.14
		Rotifera	Keratella cochlearis	3.17	0.01
July	FF2-1		Notholca laurentiae	0.89	0.03
			Gastropus stylifer	0.17	0.01
			Polyarthra remata	2.34	0.12
			Synchaeta stylata	1.06	0.09
		Copepoda - nauplii	Nauplii	1.06	0.42
			Diaptomidae	0.01	0.18
		Calanoida	Copepodites	0.86	3.66
		Quality side	Cyclops strenuus	0.64	7.28
		Cyclopoida	Copepodites	2.09	6.50
			Daphnia longiremis	0.02	0.02
		Cladocera	Holopedium gibberum	0.09	0.33
			Conochiloides natans	0.12	0.02
			Conochilus unicornis	2.66	0.06
			Filinia terminalis	0.06	0.00
			Kellicottia longispina	11.52	0.12
			Keratella cochlearis	1.61	0.01
		Rotifera	Keratella hiemalis	0.06	0.00
			Notholca laurentiae	0.37	0.01
July	FF2-2		Gastropus stylifer	0.12	0.01
			Polyarthra remata	2.66	0.13
			Synchaeta stylata	0.62	0.05
		Copepoda - nauplii	Nauplii	1.98	0.79
			Epischura sp.	0.03	1.07
		Calanoida	Copepodites	2.57	11.74
			Cyclops strenuus	0.73	7.28
		Cyclopoida	Diacyclops thomasi	0.02	0.18
			Copepodites	1.70	4.29



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	0.02	0.01
	eriod Station	Cladocera	Chydorus sphaericus	0.02	0.02
		Claubcera	Daphnia longiremis	0.11	0.32
			Holopedium gibberum	0.09	0.53
			Collotheca pelagica	0.09	0.00
			Conochiloides natans	0.15	0.02
			Conochilus unicornis	1.89	0.04
			Filinia terminalis	0.19	0.00
			Asplanchna priodonta	0.03	0.03
		Datifara	Kellicottia longispina	13.13	0.13
July	FF2-3	Rotifera	Keratella cochlearis	2.42	0.01
			Notholca laurentiae	0.09	0.00
			Gastropus stylifer	0.40	0.02
			Ploesoma truncatum	0.03	0.00
			Polyarthra remata	2.45	0.13
			Synchaeta stylata	0.99	0.08
		Copepoda - nauplii	Nauplii	2.29	0.92
		Calanoida	Copepodites	1.47	6.21
		Cyclopoida	Cyclops strenuus	0.62	6.57
			Diacyclops thomasi	0.06	0.54
			Copepodites	3.24	5.85
			Daphnia longiremis	0.07	0.07
		Cladocera	Holopedium gibberum	0.12	0.55
			Collotheca pelagica	0.28	0.00
			Conochilus unicornis	1.72	0.04
			Filinia terminalis	0.28	0.00
			Kellicottia longispina	12.03	0.12
			Keratella cochlearis	1.67	0.01
		Rotifera	Keratella hiemalis	0.14	0.01
L.L.	FF0 4		Notholca laurentiae	0.70	0.02
July	FF2-4		Gastropus stylifer	0.19	0.01
			Ploesoma truncatum	0.05	0.00
			Polyarthra remata	1.58	0.08
			Synchaeta stylata	1.67	0.19
		Copepoda - nauplii	Nauplii	1.25	0.50
			Epischura nevadensis	0.02	0.66
		Calanoida	Copepodites	1.35	6.12
		-	Cyclops strenuus	0.72	7.67
		Cyclopoida	Copepodites	2.56	5.25



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	0.05	0.18
		Cladagara	Daphnia longiremis	0.07	0.30
		Cladocera	Daphnia middendorffiana	0.02	0.13
			Holopedium gibberum	0.12	0.58
			Collotheca pelagica	0.42	0.01
			Conochiloides natans	0.19	0.03
			Conochilus unicornis	1.72	0.04
			Filinia terminalis	0.09	0.00
			Asplanchna priodonta	0.05	0.05
		Rotifera	Kellicottia longispina	11.52	0.12
July	FF2-5 ^(a)	Rolliela	Keratella cochlearis	3.21	0.01
			Keratella hiemalis	0.09	0.00
			Notholca laurentiae	0.33	0.01
			Gastropus stylifer	0.19	0.01
			Polyarthra remata	2.23	0.12
			Synchaeta stylata	0.88	0.10
		Copepoda - nauplii	Nauplii	1.58	0.63
		Copepodites	Copepodites	1.21	6.04
		Cyclopoida	Cyclops strenuus	0.84	9.33
			Diacyclops thomasi	0.05	0.37
			Copepodites	2.67	5.15
			Bosmina longirostris	0.01	0.01
		Cladocera	Daphnia longiremis	0.06	0.11
		Claudcela	Daphnia middendorffiana	0.00	0.00
			Holopedium gibberum	0.00	0.00
			Collotheca pelagica	0.14	0.00
			Conochiloides natans	0.09	0.01
			Conochilus unicornis	2.23	0.05
			Filinia terminalis	0.14	0.00
lubz	БЕ2 Б ^(b)		Asplanchna priodonta	0.05	0.03
July	FF2-3	Rotifera	Kellicottia longispina	10.87	0.11
			Keratella cochlearis	2.23	0.01
			Notholca laurentiae	0.19	0.01
			Gastropus stylifer	0.33	0.02
			Polyarthra remata	2.42	0.12
			Synchaeta stylata	0.79	0.06
		Copepoda - nauplii	Nauplii	2.23	0.89
		Copepodites	Copepodites	1.14	5.17
		Cyclopoida	Cyclops strenuus	0.56	6.54



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Diacyclops thomasi	0.03	0.24
			Copepodites	1.93	4.83
		Cladocera	Daphnia longiremis	0.15	0.79
		Cladocera	Holopedium gibberum	0.21	2.74
			Collotheca pelagica	0.17	0.00
			Conochilus unicornis	7.93	0.15
			Asplanchna priodonta	0.08	0.14
		Rotifera	Kellicottia longispina	17.61	0.18
		Rolliela	Keratella cochlearis	1.59	0.01
			Gastropus stylifer	1.09	0.04
			Ploesoma hudsoni	0.08	0.01
August	FF2-1		Polyarthra major	1.25	0.06
		Copepoda - nauplii	Nauplii	4.01	1.60
			Leptodiaptomus minutus	0.10	0.84
			Leptodiaptomus pribilofensis	0.08	1.26
		Calanoida	Epischura sp.	0.02	0.38
			Heterocope septentrionalis	0.02	1.46
			Copepodites	1.96	14.11
			Cyclops strenuus	0.44	4.33
		Cyclopoida	Diacyclops thomasi	0.02 1.96 0.44 0.13 2.67 0.30	0.79
			Copepodites	2.67	9.23
		Cladacara	Daphnia longiremis	0.30	1.51
		Cladocera	Holopedium gibberum	0.30	3.72
			Collotheca pelagica	0.15	0.00
			Conochilus unicornis	7.09	4.88
			Kellicottia longispina	22.16	0.23
		Rotifera	Keratella cochlearis	1.92	0.01
			Notholca laurentiae	0.07	0.01
			Gastropus stylifer	1.77	0.06
August	FF2 2		Polyarthra major	0.52	0.03
August	FF2-2	Copepoda - nauplii	Nauplii	2.51	1.00
			Leptodiaptomus minutus	0.26	2.52
			Leptodiaptomus pribilofensis	0.07	1.18
		Calanoida	Epischura sp.	0.02	0.24
			Heterocope septentrionalis ^(c)	-	-
			Copepodites	0.15 0.21 0.17 7.93 0.08 17.61 1.59 1.09 0.08 1.25 4.01 0.10 0.08 0.02 0.02 1.96 0.44 0.13 2.67 0.30 0.30 0.15 7.09 22.16 1.92 0.07 1.77 0.52 2.51 0.26 0.07	25.28
			Cyclops strenuus	0.48	5.48
		Cyclopoida	Diacyclops thomasi	0.06	0.27
			Copepodites	2.16	6.75



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
			Bosmina longirostris	0.02	0.04
		Cladocera	Daphnia longiremis	0.39	2.27
			Holopedium gibberum	0.35	4.47
			Collotheca pelagica	0.16	0.00
			Conochilus unicornis	6.76	0.16
		Datifana	Kellicottia longispina	28.52	0.29
		Rotifera	Keratella cochlearis	2.51	0.01
			Gastropus stylifer	1.26	0.05
Aure	FF0 0		Polyarthra major	2.36	0.12
August	FF2-3	Copepoda - nauplii	Nauplii	3.14	1.26
			Leptodiaptomus minutus	0.10	0.99
			Leptodiaptomus pribilofensis	0.06	0.86
		Calanoida	Epischura sp.	0.06	1.41
			Heterocope septentrionalis ^(c)	-	-
			Copepodites	1.87	13.26
		Cyclopoida	Cyclops strenuus	0.22	2.79
			Diacyclops thomasi	0.04	0.32
			Copepodites	2.12	7.79
		Cladadara	Daphnia longiremis	0.22	0.71
		Cladocera	Holopedium gibberum	0.15	2.12
			Ascomorpha ecaudis	0.15	0.00
			Collotheca pelagica	0.22	0.00
			Conochilus unicornis	11.29	0.27
		Rotifera	Kellicottia longispina	(org/L) 0.02 0.39 0.35 0.16 6.76 28.52 2.51 1.26 2.36 3.14 0.10 0.06 0.06 - 1.87 0.22 0.04 2.12 0.22 0.15 0.15 0.22	0.30
			Keratella cochlearis	1.83	0.01
			Gastropus stylifer	0.44	0.02
August	FF2-4 ^(a)		Polyarthra major	2.86	0.14
		Copepoda - nauplii	Nauplii	3.22	1.29
			Leptodiaptomus pribilofensis	0.07	1.09
		Ostansida	Leptodiaptomus sicilis	0.07	0.87
		Calanoida	Heterocope septentrionalis ^(c)	-	-
			Copepodites	3.30	26.97
			Cyclops strenuus	0.81	8.65
		Cyclopoida	Diacyclops thomasi	0.04	0.23
			Copepodites	3.63	13.73



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
		Cladocera	Daphnia longiremis	0.11	0.83
		Ciauocera	Holopedium gibberum	0.26	1.51
			Asplanchna priodonta	(org/L) 0.11	0.18
			Collotheca pelagica		0.00
			Conochilus unicornis	12.90	0.31
		Rotifera	Kellicottia longispina	34.59	0.36
			Keratella cochlearis	2.27	0.01
			Gastropus stylifer	1.17	0.04
August	FF2-4 ^(b)		Polyarthra major	2.20	0.11
		Copepoda - nauplii	Nauplii	4.91	1.96
			Leptodiaptomus minutus	0.11	0.88
		Calanaida	Leptodiaptomus pribilofensis	0.22	2.66
		Calanoida	Heterocope septentrionalis	0.04	3.95
			Copepodites	3.08	23.49
			Cyclops strenuus	0.59	6.54
		Cyclopoida	Diacyclops thomasi	0.15	0.83
			Copepodites	4.40	14.96
			Bosmina longirostris	0.03	0.08
		Cladocera	Eubosmina longispina	0.03	0.19
			Daphnia longiremis	0.34	1.44
			Holopedium gibberum	0.50	3.98
			Collotheca pelagica	0.19	0.00
			Conochilus unicornis	22.23	0.49
			Asplanchna priodonta	0.13	0.13
		Rotifera	Kellicottia longispina	15.15	0.16
			Keratella cochlearis	0.88	0.00
A			Gastropus stylifer	0.31	0.01
August	FF2-5		Polyarthra major	1.25	0.06
		Copepoda - nauplii	Nauplii	3.88	1.55
			Leptodiaptomus minutus	0.13	1.22
			Leptodiaptomus pribilofensis	0.16	2.23
		Calanoida	Epischura nevadensis	0.09	1.79
			Heterocope septentrionalis ^(c)	-	-
			Copepodites	2.97	17.58
			Cyclops strenuus	0.69	7.83
		Cyclopoida	Diacyclops thomasi	0.16	0.94
			Copepodites	3.51	11.83



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)			
			Bosmina longirostris	0.05	0.11			
		Cladocera	Daphnia longiremis	0.28	1.27			
			Holopedium gibberum	0.05	1.11			
			Collotheca pelagica	0.56	0.00			
			Conochilus unicornis	6.94	0.12			
			Filinia terminalis	0.19	0.00			
		Rotifera	Kellicottia longispina	10.84	0.11			
		Rolliela	Keratella cochlearis	0.68	0.00			
			Keratella serrulata	0.06	0.00			
September	FF2-1		Gastropus stylifer	0.25	0.01			
September			Polyarthra major	0.62	0.03			
		Copepoda - nauplii	Nauplii	6.88	2.75			
			Leptodiaptomus pribilofensis	0.39	5.49			
		Calanoida	Leptodiaptomus sicilis	0.85	11.37			
		Calanoida	Epischura nevadensis	0.08	1.71			
			Copepodites	0.15	1.19 5.70 1.37			
			Cyclops strenuus	0.68				
		Cyclopoida	Diacyclops thomasi	0.19				
			Copepodites	2.18	7.90			
			Bosmina longirostris	0.03	0.02			
		Cladagara	Eubosmina longispina	0.05	0.29			
		Cladocera	Daphnia longiremis	0.81	2.34			
			Holopedium gibberum	0.11 2.12				
			Collotheca pelagica	0.22	0.00			
			Conochilus unicornis	7.10	0.11			
			Asplanchna priodonta	0.05	0.08			
		Rotifera	Kellicottia longispina	10.73	0.11			
		Rolliela	Keratella cochlearis	1.46 0.00				
Sontombor	FF2-2		Ascomorpha ecaudis	0.05	0.00			
September	FF2-2		Gastropus stylifer	0.11	0.00			
			Polyarthra major	0.54	0.03			
		Copepoda - nauplii	Nauplii	7.53	3.01			
			Leptodiaptomus pribilofensis	0.68	9.27			
		Calanoida	Leptodiaptomus sicilis	1.25	15.15			
		Calanolua	Epischura nevadensis	0.03	0.60			
			Copepodites	0.03	0.23			
			Cyclops strenuus	0.65	5.82			
		Cyclopoida	Diacyclops thomasi	0.60	3.68			
			Copepodites	2.57	9.71			



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)
		Cladocera	Daphnia longiremis	0.28	0.90
			Collotheca pelagica	0.32	0.00
			Conochilus unicornis	6.61	0.12
		Rotifera	Asplanchna priodonta	0.16	0.22
		Rolliela	Kellicottia longispina	8.63	0.09
			Keratella cochlearis	1.85	0.00
			Polyarthra major	0.81	0.04
September	FF2-3	Copepoda - nauplii	Nauplii	6.85	2.74
Coptombol			Leptodiaptomus pribilofensis	0.38	5.50
		Calanaida	Leptodiaptomus sicilis	1.75	24.65
		Calanoida	Epischura nevadensis	0.04	0.95
			Copepodites	0.08	0.65
			Cyclops strenuus	0.50	5.56
		Cyclopoida Diacyclops thomasi			2.45
			Copepodites	2.38	8.64
			Bosmina longirostris	0.15	0.19
		Cladagara	Daphnia longiremis	0.76	2.31
		Cladocera	Daphnia middendorffiana	0.03	0.27
			Holopedium gibberum	0.06	0.60
			Collotheca pelagica	0.23	0.00
			Conochilus unicornis	6.33	0.11
			Filinia terminalis	0.12	0.00
			Kellicottia longispina	8.56	0.09
		Rotifera	Keratella cochlearis	1.17	0.00
			Gastropus stylifer	0.12	0.00
			Monostyla lunaris	0.06	0.00
September	FF2-4		Polyarthra major	0.23	0.01
			Trichotria tetractis	0.06	0.00
		Copepoda - nauplii	Nauplii	6.92	2.77
			Leptodiaptomus ashlandi	0.06	0.43
			Leptodiaptomus minutus	0.03	0.23
			Leptodiaptomus pribilofensis	0.29	3.72
		Calanoida	Leptodiaptomus sicilis	1.11	14.65
			Epischura nevadensis	0.03	0.63
			Copepodites	0.03	0.20
			Cyclops strenuus	0.56	5.46
		Cyclopoida	Diacyclops thomasi	0.41	2.25
			Copepodites	2.11	7.49



Sampling Period	Station	Major Taxonomic Group	Taxonomic Name	Abundance (org/L)	Biomass (mg/m³)	
			Eubosmina longispina	0.02	0.03	
		Cladocera	Daphnia longiremis	0.40	1.57	
			Holopedium gibberum	0.02	0.37	
			Collotheca pelagica	0.81	0.00	
			Conochilus unicornis	5.86	0.10	
			Asplanchna priodonta	0.07	0.05	
			Kellicottia longispina	10.55	0.11	
		Rotifera	Keratella cochlearis	1.32	0.00	
		Ascon	Ascomorpha ecaudis	0.07	0.00	
Contombor	FF2-5		Ascomorpha ovalis	0.07	0.00	
September	FF2-3		Gastropus stylifer	0.07	0.00	
			Polyarthra major	2.27	0.11	
		Copepoda - nauplii	Nauplii	6.38	2.55	
			Leptodiaptomus pribilofensis	0.13	1.73	
		Calanoida	Leptodiaptomus sicilis	1.34	17.75	
		Calanoida	Epischura nevadensis	0.04	0.73	
			Copepodites	0.07	0.67	
			Cyclops strenuus	0.27	2.44	
		Cyclopoida	Diacyclops thomasi	0.33	1.84	
			Copepodites	1.70	6.35	

Notes: Samples were analyzed by EcoAnalysts, Inc. Values are rounded to the nearest whole number.

org/L = number of organisms per litre; mg/m³ = milligrams per cubic metre; sp. = species; - = not applicable.

a) Replicate 1.

b) Replicate 2.

c) Presence of *Heterocope septentrionalis* was noted in the fine fraction of this sample, but not enumerated or measured by the taxonomist. This taxon was counted towards taxonomic richness, but abundance and biomass could not be quantified.



Table A-12 Summary of Zooplankton Community Data for Lakes in the Jay Project Area, 2014

								Relat	ive Abundan	ce (%)	-		Rela	tive Biomas	s (%)		
Basin	Waterbody	Station	Sampling Period Re		Total Abundance (org/L)	Total Biomass (mg/m³)		Calanoida	Cladocera	Cyclopoida	Rotifera	Copepod nauplii	Calanoida	Cladocera	Cyclopoida	Rotifera	Copepod nauplii
		Aa-1	August		83	52	16	0.4	19.6	6.5	66.6	6.9	7.7	56.7	29.4	1.8	4.4
		Ad-1	September		69	43	15	0.2	8.3	10.9	70.0	10.6	2.4	24.5	50.6	15.8	6.7
			July		64	44	14	0.7	10.6	5.7	75.7	7.4	3.9	65.4	25.2	1.1	4.3
		Ab-1	August		49	61	14	0.4	25.4	8.3	57.3	8.7	2.6	65.6	28.4	0.6	2.8
			September		59	29	14	0.0	7.0	8.7	56.7	27.6	0.0	22.5	45.1	10.3	22.1
			July		55	30	13	0.6	5.6	6.4	82.9	4.5	2.9	44.2	47.8	1.6	3.4
		Ac-1	August		61	59	16	0.7	18.4	4.9	62.9	13.1	8.2	63.8	20.9	1.7	5.4
		AC-1	September	Rep 1	53	30	16	0.1	4.9	13.2	55.4	26.3	5.7	5.7 17.2 52.3	52.3	6.2	18.6
			September	Rep 2	52	27	16	0.1	5.2	11.6	57.8	25.3	6.2	13.2	50.7	10.3	19.5
			July		58	50	13	0.7	12.9	11.0	66.5	8.9	1.7	50.7	42.5	1.0	4.1
	Lac du Sauvage	Ac-4	August		79	97	12	0.7	22.0	13.4	49.3	14.6	6.0 49.8 38.6	38.6	0.9	4.8	
Lac du Sauvage			September		48	35	13	0.1	6.1	14.8	45.9	33.1	2.7	15.1	56.2	7.5	18.5
			July		66	19	16	1.0	3.0	5.5	84.5	6.0	7.6	22.2	57.4	4.4	8.4
		Ac-7	August		47	41	16	1.5	14.1	6.3	65.6	12.4	20.8	41.0	30.8	1.7	5.7
			September		46	29	14	0.1	6.8	12.7	47.9	32.5	3.0	15.5	53.3	7.7	20.5
			July		32	13	12	0.2	3.4	8.4	80.2	7.8	2.1	22.4	65.4	2.4	7.8
		Ad-1	August		64	44	14	0.7	10.6	5.7	75.7	7.4	3.9	65.4	25.2	1.1	Build Bodd .8 4.4 5.8 6.7 .1 4.3 .6 2.8 0.3 22.1 .6 3.4 .7 5.4 5.2 18.6 0.3 19.5 .0 4.1 0.9 4.8 7.5 18.5 .4 8.4 .7 5.7 .7 20.5 .4 7.8 .1 4.3 .6 18.3 .3 3.7 .5 2.0 .5 16.0 .0 8.0
			September		60	34	14	0.0	2.9	15.8	55.1	26.3	0.0	6.8	67.4	7.6	18.3
			July		51	29	13	1.0	9.4	7.2	77.2	5.1	2.8	57.9	33.4	2.3	3.7
		Ae-1	August		95	152	14	1.8	30.7	5.2	54.3	8.0	9.3	70.8	17.3	0.5	2.0
			September		80	55	12	0.2	6.1	14.1	52.1	27.5	3.1	21.8	50.7	8.5	16.0
	Duchoco Loko	Af-1	August		66	41	16	0.1	48.0	0.7	38.8	12.3	7.2	76.3	6.5	2.0	8.0
	Duchess Lake	AI-1	September		76	33	13	0.2	30.8	6.2	60.8	2.1	2.4	60.5	20.1	15.1	1.9

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									Relat	ive Abundan	ce (%)		Relative Biomass (%)					
Basin	Waterbody	Station	Sampling Period	Replicate	Total Abundance (org/L)	Total Biomass (mg/m³)	Taxonomic Richness (No. of Taxa)	Calanoida	Cladocera	Cyclopoida	Rotifera	Copepod nauplii	Calanoida	Cladocera	Cyclopoida	Rotifera	Copepod nauplii	
			July		63	27	11	1.1	0.6	5.6	91.2	1.5	12.5	7.4	73.5	5.2	1.4	
		S2	August		32	40	14	6.6	4.9	6.9	65.0	16.7	45.1	22.3	26.3	0.9	5.3	
			September		20	45	13	9.7	3.5	7.8	52.2	26.9	59.0	20.6	15.2	0.4	4.7	
			July		20	26	12	6.2	0.4	17.8	63.4	12.2	29.9	1.9	63.5	0.8	3.9	
		S3	August	Rep 1	12	23	12	9.6	0.7	10.0	60.3	19.4	65.4	3.1	26.8	0.6	4.1	
		53	August	Rep 2	12	19	10	8.9	0.6	10.7	61.1	18.7	52.5	2.6	39.4	0.8	4.7	
	Slipper Bay		September		14	38	12	12.2	1.5	6.4	57.5	22.4	70.0	16.2	9.9	0.4	3.4	
			July	Rep 1	12	16	14	12.8	1.4	11.4	64.8	9.6	41.4	5.6	49.1	1.0	3.0	
		S5	July	Rep 2	14	16	11	16.0	0.6	11.3	60.9	11.2	38.9	3.3	53.2	0.9	3.7	
		30	August		11	28	12	20.5	1.5	7.0	54.9	16.1	67.3	13.4	16.2	0.4	2.7	
			September		9	53	12	33.6	5.3	10.0	23.8	27.4	ED ED DO DO <thdo< th=""> DO DO DO<!--</td--><td>1.8</td></thdo<>	1.8				
		S6	July		16	16	12	12.9	0.5	8.8	61.9	15.9	42.8	6.1	43.6	1.2	6.3	
		30	August		12	24	11	14.6	1.0	10.3	48.4	25.7	55.9	11.2	27.3	0.4	5.2	
			July		27	19	14	3.3	0.3	10.1	82.4	3.9	20.2	2.6	72.6	2.3	2.2	
Lac de Gras		FF2-1	August		40	38	15	5.5	0.9	8.2	75.3	10.1 47.3	9.2	37.7	1.6	4.2		
Lac de Glas			September		32	40	14	4.6	1.2	9.6	63.1	21.6	49.1	6.2	37.2	0.7	6.8	
			July		27	26	14	9.6	0.4	9.1	73.5	7.4	49.1	1.3	45.0	1.6	3.0	
		FF2-2	August		43	53	14	8.6	1.4	6.2	77.9	5.8	1.5 12.5 7.4 73.5 5.2 16.7 45.1 22.3 26.3 0.9 26.9 59.0 20.6 15.2 0.4 12.2 29.9 1.9 63.5 0.8 19.4 65.4 3.1 26.8 0.6 18.7 52.5 2.6 39.4 0.8 22.4 70.0 16.2 9.9 0.4 9.6 41.4 5.6 49.1 1.0 11.2 38.9 3.3 53.2 0.9 16.1 67.3 13.4 16.2 0.4 27.4 50.4 40.3 7.4 0.1 15.9 42.8 6.1 43.6 1.2 25.7 55.9 11.2 27.3 0.4 3.9 20.2 2.6 72.6 2.3 10.1 47.3 9.2 37.7 1.6 21.6 49.1 6.2 37.2 0.7 7.4 49.1 1.3 45.0 1.6 5.8 29.2 5.2 12.5 5.2 21.8 48.0 9.1 36.5 0.6 7.7 29.0 4.1 60.4 2.2 6.3 45.8 18.8 30.2 1.7 22.1 60.5 1.7 31.7 0.9 4.8 31.9 2.9 60.6 2.3 5.6 51.3 5.0 40.1 1.3 7.3 52.8 4.0 38.1 1.7 </td <td>5.2</td> <td>1.0</td>	5.2	1.0			
			September		35	53	15	5.7	2.9	11.0	58.6	21.8	48.0	9.1	36.5	0.6	5.7	
			July		30	21	18	4.9	0.8	13.2	73.4	7.7	29.0	4.1	60.4	2.2	4.3	
		FF2-3	August		50	36	14	4.2	1.5	4.8	83.2	6.3	45.8	18.8	30.2	1.7	3.5	
	Far-field 2		September		31	53	11	7.3	0.9	10.5	59.3	22.1	60.5	1.7	31.7	0.9	5.2	
			July		26	21	14	5.2	0.7	12.4	76.9	4.8	31.9	2.9	60.6	2.3	2.4	
		FF2-4	August	Rep 1	57	56	13	6.0	0.6	7.8	80.0			5.0		1.3	2.3	
		112-4	August	Rep 2	67	59	13	5.1	0.5	7.6	79.4			-	38.1	1.7	3.4	
			September		29	41	16	5.3	3.4	10.5	57.4	23.5	47.9	8.1	36.7	0.5	6.7	
			July	Rep 1	28	23	16	4.4	0.9	12.9	76.0	5.7	26.0	5.1	64.0	2.2	2.7	
		FF2-5	July	Rep 2	25	18	15	4.5	0.3	9.9	76.6		28.4	0.6	63.7	2.4	4.9	
		112-5	August		53	52	15	6.4	1.7	8.3	76.3	7.4	44.3	11.0	40.0	1.7	3.0	
			September		32	36	15	5.0	1.4	7.3	66.4	20.0	57.3	5.4	29.2	1.1	7.0	

Summary of Zooplankton Community Data for Lakes in the Jay Project Area, 2014 Table A-12

Notes: Samples were analyzed by EcoAnalysts, Inc. Total abundance and biomass values are rounded to the nearest whole number. org/L = number of organisms per litre; mg/m³ = milligrams per cubic metre; No. = number; % = percent; Late Spring = July; Summer = August; Fall = September.

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