Modelling lake trophic state: A random forest approach

Jeffrey W. Hollister * 1 W. Bryan Milstead 1 Betty J. Kreakie 1

¹ US Environmental Protection Agency, Office of Research and Development, National Health and Environmental Effects Research Laboratory, Atlantic Ecology Division, 27 Tarzwell Drive Narragansett, RI, 02882, USA

 * corresponding author: hollister.jeff@epa.gov

Abstract

Productivity of lentic ecosystems is well studied and it is widely accepted that as nutrient inputs increase, productivity increases and lakes transition from lower trophic state (e.g., oligotrophic) to higher trophic states (e.g., eutrophic). These broad trophic state classifications are good predictors of ecosystem condition, services (e.g., recreation and aesthetics), and disservices (e.g., harmful algal blooms). While the relationship between nutrients and trophic state provides reliable predictions, it requires in situ water quality data in order to parameterize the model. This limits the application of these models to lakes with existing and, more importantly, available water quality data. To address this, we take advantage of the availability of a large national lakes water quality database (i.e., the National Lakes Assessment), land use/land cover data, lake morphometry data, other universally available data, and apply data mining approaches to predict trophic state. Using these data and random forests, we first model chlorophyll a, then classify the resultant predictions into trophic states. The full model estimates chlorophyll a with both in situ and universally available data. The mean squared error and adjusted R² of this model was 0.09 and 0.8, respectively. The second model uses universally available GIS data only. The mean squared error was 0.22 and the adjusted R² was 0.48. The Kappa coefficient of the trophic state classifications derived from the chlorophyll a predictions were 0.57 for the full model and 0.29 for the "GIS only" model. Random forests extend the usefulness of the class predictions by providing prediction probabilities for each lake. This allows us to make trophic state predictions and also indicate the level of uncertainty around those predictions. For the full model, these predicted class probabilities ranged from 0.42 to 1. For the GIS only model, they ranged from 0.33 to 0.96. It is our conclusion that in situ data are required for better predictions, yet GIS and universally available data provide trophic state predictions, with estimated uncertainty, that still have the potential for a broad array of applications. The source code and data for this manuscript are available from https://github.com/USEPA/LakeTrophicModelling.

Keywords: Harmful Algal Blooms; Cyanobacteria; Open Science; Nutrients; National Lakes

Assessment

₁₆ 1 Introduction

Productivity in lentic systems is often categorized across a range of trophic states (e.g., the trophic continuum) from early successional (i.e., oligotrophic) to late successional lakes (i.e., hypereutrophic) with lakes naturally occurring across this range (Carlson 1977). Oligotrophic lakes occur in nutrient poor areas or have a more recent geologic history, are often found in higher elevations, have clear water, and are usually favored for drinking water or direct contact recreation (e.g., swimming). Lakes with higher productivity (e.g., mesotrophic and eutrophic lakes) have greater nutrient loads, tend to be less clear, have greater density of aquatic plants, and often support more diverse and abundant fish communities. Higher primary productivity is not necessarily a predictor of poor ecological condition as it is natural for lakes to shift from lower to higher trophic states but this is a slow process (Rodhe 1969). However, at the highest productivity levels (hypereutrophic lakes) biological integrity is compromised (Hasler 1969, Smith et al. 1999, Schindler and Vallentyne 2008).

Monitoring trophic state allows for rapid assessment of a lakes biological productivity and identification of lakes with unusually high productivity (e.g., hypereutrophic). These cases are indicative of lakes under greater anthropogenic nutrient loads, also known as cultural eutrophication, and are more likely to be at risk of fish kills, beach fouling, and harmful algal blooms (Smith 1998, Smith et al. 1999, 2006). Given the association between trophic state and many ecosystem services and disservices, being able to accurately model trophic state could provide a first cut at identifying lakes with the potential for harmful algal blooms (i.e., from cyanobacteria) or other problems associated with cultural eutrophication. This type of information could be used for setting priorities for management and allow for more efficient use of limited resources.

As trophic state and related indices can be best defined by a number of *in situ* water quality parameters (modeled or measured), most models have used this information as predictors

(Imboden and Gächter 1978, Salas and Martino 1991, Carvalho et al. 2011, Milstead et al. 2013).

This leads to accurate models, but these data are often sparse and not always available, thus limiting the population of lakes for which we can make predictions. A possible solution for this issue is to build models that use widely available data that are correlated to many of the *in situ* variables. For instance, landscape metrics of forests, agriculture, wetlands, and urban land in contributing watersheds have all been shown to explain a significant proportion of the variation (ranging from 50-86%, depending on study) in nutrients in receiving waters (Jones et al. 2001, 2004, Seilheimer et al. 2013). Building on these previously identified associations might allow us to use only landscape and other universally available data to build models. Identifying predictors using this type of ubiquitous data would allow for estimating trophic state in both monitored and unmonitored lakes. Furthermore, being able to classify a large number of lakes would have implications for the management of lakes. A broader discussion of ecological classification and resource management is beyond the scope of this paper, but see (Carpenter 1999) for more information on this topic.

Many published models of nutrients and trophic state in freshwater systems are based on linear modelling methods such as standard least squares regression or linear mixed models (Jones et al. 2001, 2004). While these methods have proven to be reliable, they have limitations (e.g., independence, distribution assumptions, and outlier sensitivity). Using data mining approaches, such as random forests, avoids many of the limitations, may reduce bias, and often provides better predictions (Breiman 2001, Cutler et al. 2007, Peters et al. 2007, Fernández-Delgado et al. 2014). For instance, random forests are non-parametric and thus the data do not need to come from a specific distribution (e.g., Gaussian) and can contain collinear variables (Cutler et al. 2007). Second, random forests work well with very large numbers of predictors (Cutler et al. 2007). Lastly, random forests can deal with model selection uncertainty as predictions are based upon a consensus of many models and not just a single model selected with some measure of goodness of fit.

The research presented here builds on past work in three areas. First, we built, assessed, and compared two random forest models of chlorophyll a with 1) in situ and universally available GIS data and then 2) universally available GIS data only. Second, we converted the chlorophyll a estimates, for both models, to trophic state and assessed prediction accuracy and uncertainty. Third, we examined the important predictors for both models. Lastly, to promote transparency in our work, the analysis code and data are available as an R package from https://github.com/USEPA/LakeTrophicModelling.

94 2 Methods

95 2.1 Data and Study Area

We utilized three primary sources of data for this study, the National Lakes Assessment (NLA), the National Land Cover Dataset (NLCD), and lake morphometery modeled from the NHDPlus and National Elevation Data Set (Homer et al. 2004, USEPA 2009, Xian et al. 2009, Hollister and Milstead 2010, Hollister et al. 2011, Hollister 2014). All datasets are national in extent and provide a unique snapshot view of the condition of lakes in the conterminous United States during the summer of 2007.

The NLA dataset was collected during the summer of 2007 and the final datasets were released in 2009 (USEPA 2009). With consistent methods and metrics collected at over 1000 locations across the conterminous United States (Figure 1), the NLA provides a unique opportunity to examine broad scale patterns in lake productivity. The NLA collected data on biophysical measures of lake water quality and habitat as well as an assessment of the phytoplankton community. For this analysis, we only use the various water quality measurements from the National Lakes Assessment (USEPA 2009). Additionally, the NLA included ecological regions as defined in the Wadeable Streams Assessment (Figure 2) (Omernik 1987, USEPA 2006).

Adding to the monitoring data collected via the NLA, we used the 2006 NLCD data to examine landscape-level drivers of trophic status in lakes. The NLCD is a national land use/land cover 111 dataset that also provides estimates of impervious surface. We calculated total proportion of each 112 NLCD land use land cover class and total percent impervious surface within a 3 kilometer buffer 113 surrounding each lake (Homer et al. 2004, Xian et al. 2009). We chose this buffer distance for 114 several reasons. First, in some preliminary efforts we tried a variety of scales (300 m, 1.5 km, and 115 3 km), and they had little impact on prediction accuracy. Second, since we also include local lake 116 specific variables (see below) as well as the broader scale ecoregions, we chose the 3km buffer as it 117 made intuitive sense as representative of land use impacts that would not be accounted for these 118 other variables. While many regional classifications and scales have been shown to be effective 119 (e.g., Cheruvelil et al. 2013), we chose a three kilometer buffer as it represented an intermediate 120 scale that is greater than immediate parcels but smaller than regional and whole-basin measures. 121 Local, lake specific characteristics have been show to be important (Read et al. 2015). Thus to 122 account for this, we used measures of lake morphometry (i.e., depth, volume, fetch, etc.). As 123 these data are difficult to obtain for large numbers of lakes over broad regions, we used modeled 124 estimates of lake morphometry (Hollister and Milstead 2010, Hollister et al. 2011, Hollister 2014). 125 These included: surface area, shoreline length, Shoreline Development, Maximum Depth, Mean 126 Depth, Lake Volume, Maximum Lake Length, Mean Lake Width, Maximum Lake Width, and 127 Fetch. 128

2.2 Predicting Trophic State with Random Forests

Random forest is a machine learning algorithm that aggregates numerous decision trees in order to obtain a consensus prediction of the response categories (Breiman 2001). Bootstrapped sample data are recursively partitioned according to a given random subset of predictor variables and a predetermined number of decision trees are developed. With each new tree, the sample data subset is randomly selected and with each new split, the subset of predictor variables are randomly selected. For a more detail description of random forests see Breiman (2001) and Cutler et al. (2007).

Random forests are able to handle numerous correlated variables without a decrease in prediction accuracy; however, one possible shortcoming of this approach is that the resulting model may 138 be difficult to interpret, thus selecting the most important variables is an important first step. 139 Several methods have been proposed to do this with random forest. For instance, this is a problem 140 often faced in gene selection and in that field, a variable selection method based on random forest 141 has been successfully applied and implemented in the R Language as the varSelRF package 142 (Díaz-Uriarte and De Andres 2006), but this is limited to classification problems. Additionally, 143 others have suggested alternative variable importance measures, but this is only needed with a 144 large number of categorical variables which are selected against with traditional random forest 145 approach (Strobl et al. 2007). 146

In our case, we predicted a continuous variable, chlorophyll a, directly thus varSelRF, does not apply, and nearly all of our variables are continuous so the approach suggested by Strobl (2007) 148 is not necessary. Thus we developed an approach, similar to varSelRF but applied to random 149 forest with regression trees. With this approach we fit a full random forest model that includes 150 all variables and a large number of trees. We then rank the variables using the increase in mean 151 square error, which has been shown to be a less biased metric of importance than the mean 152 decrease in the Gini coefficient (Strobl et al. 2007). Using this ranking, we then iterate through 153 the variables and create a random forest with the top two variables and record mean square error 154 and adjusted R² of the resultant random forest. We then repeat this process by adding the next 155 most important variable in order of importance. With this information we identify both the top 156 variables and the point at which adding variables does not improve the fit of the overall model. 157 These variables are selected and used as the "reduced model." With this method, a minimum set 158 of variables that maximizes model accuracy is provided. This allows us to start with a full suite 159

of predictor variables from which to select a minimum, easier to interpret set of variables.

$_{\scriptscriptstyle 161}$ 2.3 Model Details

We used the randomForest package in R to build predictive models of chlorophyll a with two sets of predictors (Liaw and Wiener 2002). The first included in situ and universally available GIS predictors. We refer to this as the "All variables" model. For the second model we used just the universally available data (i.e., no in situ information). This is referred to as the "GIS only" model. A list of all considered variables is in Appendix 1. Our separation of predictors was chosen so that we could highlight the additional predictive performance provided by adding the in situ water quality variables on top of the GIS only variables. Lastly, we used only complete cases (i.e., missing data were removed) so the total number of observations varied among models.

Our modelling work flow was as follows:

- 171 1. Identify a minimal set of variables from the full suite of variables (Appendix 1) that
 172 maximize accuracy of the random forest algorithm. This minimal set of variables, the
 173 reduced model, is calculated for each of the models.
- 2. Using R's randomForest package, we develop two random forest models with 5000 trees ("All variables" and "GIS only").
- 3. Assess model performance for both the predicted chlorophyll *a* and for categorical trophic state classifications. Trophic state was defined using the NLA chlorophyll *a* trophic state cut offs (Table 1).
- 4. Examine importance and partial dependence of the most important variables.

2.4 Measures of Model Performance and Variable Importance

We assessed the performance of the random forest two ways. First we compared the root mean 181 square error and the adjusted R² of the models. Second, we examined the accuracy of the model 182 predictions when converted to trophic states classes via a confusion matrix (Table 1). A confusion 183 matrix shows agreement and disagreement in a tabular form with predicted values forming the columns of the matrix and observed values, the rows. From this tabulated information we 185 calculated the total accuracy (i.e., percent correctly predicted) and the Kappa coefficient, which 186 takes into account the error expected by chance alone (i.e., the off diagonal values of the matrix) 187 (Cohen 1960, Hubert and Arabie 1985). The Kappa coefficient can range from -1 to 1 with 0 188 equaling the agreement expected by chance alone. Values greater than 0 represent agreement 189 greater than would be expected by chance. A Kappa coefficient greater than approximately 0.6 190 is considered "substantial" agreement (Landis and Koch 1977). Negative values are rare and 191 would indicate no agreement between the predicted and observed values. We use Kappa as a 192 means of comparison across models as well as within subsets of a given model. Additionally, 193 random forest builds each tree on bootstrapped, random subsets of the original data, thus, a 194 separate independent validation dataset is not required and random forest error estimates are 195 expected to be unbiased (Breiman 2001). 196

Random forests explicitly measure variable importance with two metrics: mean decrease in Gini 197 and percent increase in mean squared error. These measure the impact on the overall model 198 when a particular variable is included and thus can be used to assess importance (Breiman 2001). 199 The Gini Index has been shown to have a bias (Strobl et al. 2007), thus, we used percent increase 200 in mean squared error to assess variable importance. Lastly, partial dependence plots provide a 201 mechanism to examine the partial relationship between individual variables and the response 202 variable (Jones and Linder 2015). We examined these plots for the top variables as assigned by 203 percent increase in mean squared error for each the reduced models. 204

2.5 Trophic State Probabilities

One of the powerful features of random forests is the ability to aggregate a very large number of 206 competing models or trees. Each tree provides an independent prediction or vote for a possible 207 outcome. In the context of our chlorophyll a models, we have 5,000 estimates of chlorophyll 208 a for each lake. We convert these values to trophic states (Table 1) then count up total votes for each class and divide by total possible votes to get an estimate of the probability that a lake is in a given trophic state. For instance, for a single lake (National Lake Assessment ID = 211 NLA06608-0005), the vote probabilities for the "All variables" model were 95% for oligotrophic, 212 5% for mesotrophic, 0% for eutrophic, and 0% for hypereutrophic. The maximum probability 213 provides the predicted class, in this case oligotrophic, and suggests little uncertainty in this 214 prediction. We refer to this value as the "prediction probability." 215

Further, we might expect higher total accuracy for lakes that have more certain predictions. This should be evident by looking at the Kappa coefficient of lakes given their prediction probability is at or above a certain probability. To test this we use an approach similar to one outlined by Paul and MacDonald (2005) and implemented by Hollister et al. (2008) and examine the change in Kappa coefficient as a function of the prediction probability for both models.

3 Results

Our complete dataset included 1148 lakes; however 5 lakes did not have chlorophyll *a* data. Thus, the base dataset for our modelling was conducted on data for 1143 lakes. The lakes were well distributed across the four trophic state categories (Table 1) and spatially throughout the United States (Figure 1).

226 3.1 Models: All Variables

The model built with all predictors used 1080 total observations, had a mean squared error of 0.09 and and R² of 0.8. The accuracy of the four trophic states was 68.7% and the Kappa coefficient was 0.57 (Table 2). The variable selection process identified a reduced model with 20 variables (Figure 3). The six most important variables were turbidity, total phosphorus, total nitrogen, elevation, total organic carbon, and N:P ratio (Figures 4). The role that each played in predicting chlorophyll a varied (Figure 5).

$_{233}$ 3.2 Models: GIS Only Variables

The GIS only model was built using 1138 total observations, had a mean squared error of 0.22 and and R² 0.48. Four trophic states were predicted with a total accuracy of 49% and had a Kappa coefficient of 0.29 (Table 3). The variable selection process for this model produced a reduced model with 15 variables (Figure 6). The six most important variables were ecoregion, percent cropland, elevation, latitude, percent evergreen forest, and mean lake depth (Figures 7 & 5).

40 3.3 Trophic State Probabilities

The "All variables" model provides more certain model predictions with a median prediction probability of 0.81 versus 0.72 for the "GIS only" model (Figure 9). Additionally, the Kappa coefficient of the predictions is a function of this uncertainty. Lakes with more certain predictions were more accurately classified and had higher Kappa coefficients (Figure 10). For both models, when prediction probabilities are approximately 0.8 or higher, the models had a Kappa coefficient of ~1. This represents 55% of the lakes for the "All variables" model and 22% of the lakes for the "GIS only" model. A Kappa coefficient of 0.6 or higher is considered "substantial" agreement

²⁴⁸ (Landis and Koch 1977). For the "GIS only" model this is seen with 52% of the lakes. Lastly, as ²⁴⁹ prediction probabilities increased, the difference in Kappa coefficient between the two models ²⁵⁰ decreased (Figure 10 & Tables 4 & 5).

²⁵¹ 4 Discussion

2 4.1 Trophic State Probabilities

Not surprisingly, lakes with more certain predictions (i.e., higher prediction probabilities) were 253 more accurately predicted (Figure 10). The fact that the difference in accuracy (as measured by 254 the Kappa coefficient) between the two models decreased as certainty in the prediction increased 255 suggests that models with lower overall accuracy, such as the "GIS only" model, may have 256 acceptable accuracy for many individual cases (Tables 4 & 5). Additionally, the prediction 257 probabilities may be mapped for each of the four classes (Figure 11). The spatial patterns show 258 little variability between the "All variables" and "GIS only" models, thus we only show the 259 results from the more broadly applicable "GIS only" model (Figure 11). 260

This map provides several insights. First, since low uncertainty is associated with high accuracy, 261 this map shows the broad spatial patterns of lake trophic state across the United States (i.e darker 262 colors more likely to be correctly predicted). Hypereutrophic lakes are much more commonly 263 predicted in the Midwest and southeastern United States. Clear, oligotrophic lakes are in the 264 northwestern United States, through the western mountains and in the northeastern united 265 states. The middle trophic states are more evenly distributed across the country. Lastly, this 266 particular map is very similar to simply mapping the raw data. However, it highlights what 267 could be done if the "GIS only" model were used to map data without measured chlorophyll a 268 values which would provide probabilities of given trophic states for all lakes in the United States. 269

4.2 Partial dependencies of explanatory variables

In line with past predictive modelling of chlorophyll a concentrations the "All variables" model 271 selected the water quality variables (turbidity, total organic carbon, total nitrogen, total phos-272 phorus, and N:P ratios) as important variables (Downing et al. 2001). While there is variation in the response of chlorophyll a to changes in nutrient concentrations, the general pattern suggests that limiting nutrients have predictable impacts. If we examine the partial dependencies of these 275 variables we see a general linear increase in log chlorophyll a with nitrogen, phosphorus and 276 organic carbon concentrations (Figure 5). This relationship holds until nutrient concentrations 277 become saturated. The partial dependency plots (Figure 5) for the nitrogen:phosphorus ratio 278 is more complicated, indicating that for ratios less than ~ 14 chlorophyll a increases but after 279 ~14 there is marked decrease. The effect of the nitrogen phosphorus ratio on chlorophyll has been the subject of considerable research and our results are consistent with the majority of the 281 findings suggesting that at low ratio values nitrogen is limiting (Downing and McCauley 1992, 282 Smith and Schindler 2009). Conversely, at higher ratios the phosphorus levels may be limiting. 283 This would be a cause for concern with linear models; however, linearity is not an assumption of 284 tree-based modelling approaches such as random forest. 285

Turbidity was selected as the most important variable in the "All variables" model. The partial dependency analysis shows that, similar to the nutrients discussed above, log chlorophyll a increases with increased turbidity. At first this may seem counter intuitive since we might expect productivity to decrease as turbidity increases, and therefore light availability decreases (Tilzer 1988, Bilotta and Brazier 2008). However, algal biomass can contribute heavily to measures of turbidity and we expect greater productivity to lead to increased turbidity (Hansson 1992). We interpret this pattern as indicating that as chlorophyll a concentrations increase we see a concomitant increase in turbidity due to increased algal cell densities.

Elevation was selected as an important predictive variable in both the all variables and the GIS

only models; the partial dependencies (Figures 5 & 8) indicate a negative relationship between elevation and chlorophyll a concentration that is probably due to fact that the location of mountains in the United States is the spatial inverse of the distribution of agricultural and urban lands. As elevation increases we expect decreased loads due to smaller watershed contributing areas. In contrast lower elevation sites will have larger drainage areas and greater potential for increased nutrient loads from urban and agricultural sources.

The variables in the "GIS only" model captured the large scale spatial pattern of the trophic 301 status gradient of lakes across the United States. In addition to elevation, mentioned above, the 302 model was most sensitive to latitude and ecoregion. In general, chlorophyll a concentrations are 303 highest in the Southern portions of the study area where temperatures can be higher (a known 304 driver of productivity), elevations lower, and agricultural impacts more pronounced. Likewise 305 ecoregion (see Figures 2 & 8) has a pronounced affect indicting continental scale effects of land 306 use and geography. Agriculturally dominated landscapes such as the Temperate Plains, Southern 307 Plains, and Coastal Plains show the highest levels of Chlorophyll a. Whereas high elevation 308 zones (Western Mountains), arid lands (Xeric), Northern habitats (Upper Midwest) have lower 309 concentrations. 310

Further evidence for the role of land use/land cover variables, and similar to results from Read et. al. (2015), is shown by the selection of the percent cropland and percent evergreen forest variables. As indicated by the partial dependency plots (Figure 8), chlorophyll a increases with cropland and decreases with evergreen cover. It is not surprising that croplands were selected given the overwhelming impact of agriculture on the eutrophication process. Evergreens and chlorophyll a concentrations show a negative association (Figure 8). As the percent of evergreens increases we are likely to see increased elevation and soil differences that limit agriculture.

Lastly, morphometry (e.g., depth) also proved to be important in the prediction of lake trophic state (Genkai-Kato and Carpenter 2005). As morphometry shows little to no broad scale spatial pattern and is unique to a given lake, these data are likely illuminating the local, lake scale

drivers such as in-lake nutrient processing and residence time.

5 Conclusions

Our research goals were to explore the utility of a widely used data mining algorithm, random forests, in the modelling of chlorophyll a and lake trophic state. Further, we hoped to examine the utility of these models when built with only ubiquitous GIS data, which allows estimation of trophic state for all lakes in the United States. The "All variables" model had an RMSE of 0.09 and an adjusted R^2 of 0.8 whereas, the GIS only models had an RMSE of 0.22 and the adjusted R^2 was 0.48. Our total accuracy in predicting chlorophyll a based trophic states was 69% for the "All variables" model and 49% for the "GIS only" model.

While the "GIS only" model showed lower prediction accuracies than the "All variables" model, the association between the uncertainty of prediction and total accuracy (Figure 10 and Tables Tables 4 & 5) suggest that the "GIS only" model will provide reasonable estimates of trophic state for many lakes across the United States. Furthermore, we can map the uncertainty of the predictions, thus, we know the spatial patterns and location of the lakes for which we are certain, or not, of their predicted trophic state. Given this and that these models may be applied to any lake in the United States we can recommend using this model.

Future iterations of this modelling effort may be able to utilize modeled predictions of nutrients to improve accuracy and also maintain broad applicability (Milstead et al. 2013). Changes such as these have several advantages. First, this would allow for estimating changes to chlorophyll a and trophic state as a function of changing nutrient loads, which are expected due to climate change (Adrian et al. 2009, Jeppesen et al. 2011, Moss et al. 2011, Jones and Brett 2014). Second, with the ability to make predictions for most lakes in the United States, the "GIS only" models could be used as a source of information on national scale phenomena. For example,

predictions of chlorophyll *a*, with measures of uncertainty, could be used in efforts to scale up
the contributions from lakes to broad scale estimates of gross primary production.

For the "All variables" model, the *in situ* water quality variables drove the predictions. This is not surprising. For the "GIS only" model, the results were more nuanced. Three broad categories were routinely being selected as important: broad scale spatial patterns in trophic state, land use/land cover controls of trophic state, and local, lake-scale control driven by lake morphometry.

Our results raise three important considerations related to managing eutrophication. First, 351 the broad scale patterning, indicated by ecoregion as an important variable, suggests regional 352 trends. This is noteworthy because it suggests that efforts to monitor, model and manage 353 eutrophication and cyanobacteria should be undertaken at both national and regional levels. 354 This corroborates past findings that regional drivers are important for water quality (Cheruvelil 355 et al. 2013). Second, while direct control of water quality in lakes would have a large impact, 356 the land use/land cover drivers (i.e., non-point sources) of water quality are also important, and 357 better management of the spatial distribution of important classes such as forest and agriculture 358 can provide some level of control on trophic state and amount of cyanobacteria present. Third, 359 in-lake processes (i.e., residence time, nutrient cycling, etc.) are, as expected, important and 360 need to be part of any management strategy. Building on these efforts through updated models, 361 direct prediction of cyanobacteria, and additional information on the regional differences will 362 help us get a better handle on the broad scale dynamics of productivity in lakes and the potential 363 risk to human health from cyanobacteria blooms.

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Effects Research Laboratory, US Environmental Protection Agency.

³⁷³ 7 Figures

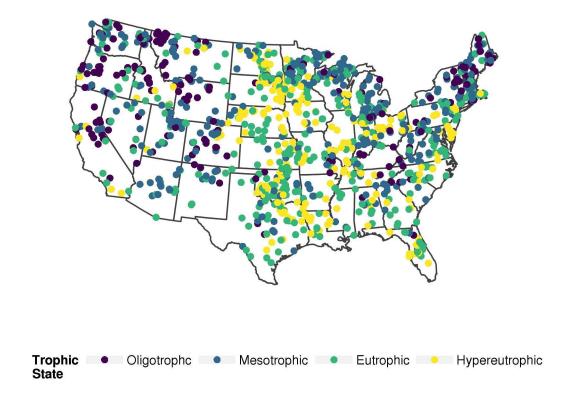


Figure 1: Map of the distribution of National Lakes Assesment Sampling locations

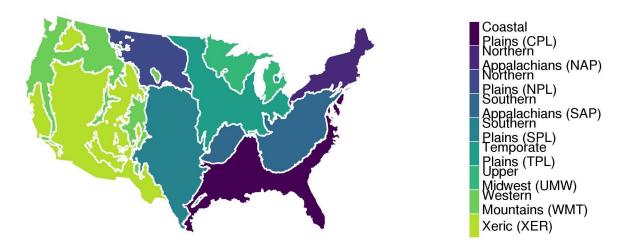


Figure 2: Wadeable Streams Assesment ecoregions

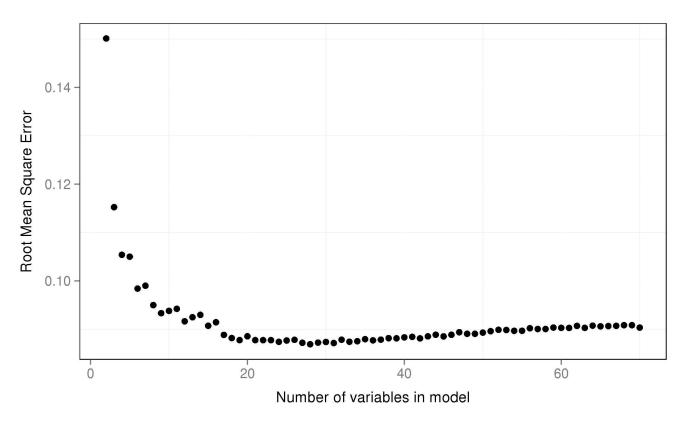


Figure 3: Variable selection plot for all variables. Shows percent increase in mean squared error as a function of the number of variables.

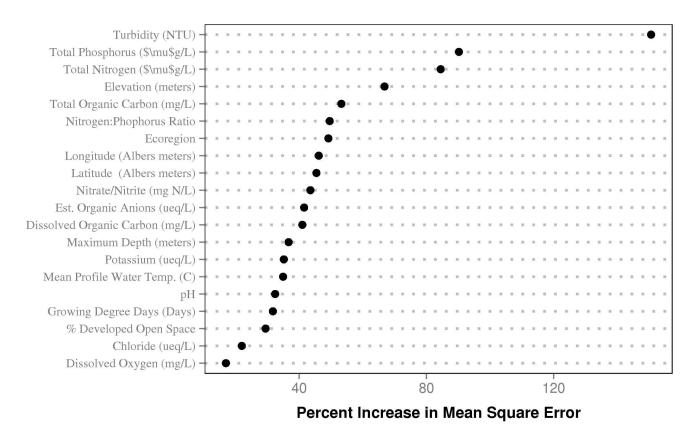


Figure 4: Importance plot for All Variables., shows percent increase in mean square error. Higher values of percent increase in mean squared error indicates higher importance.

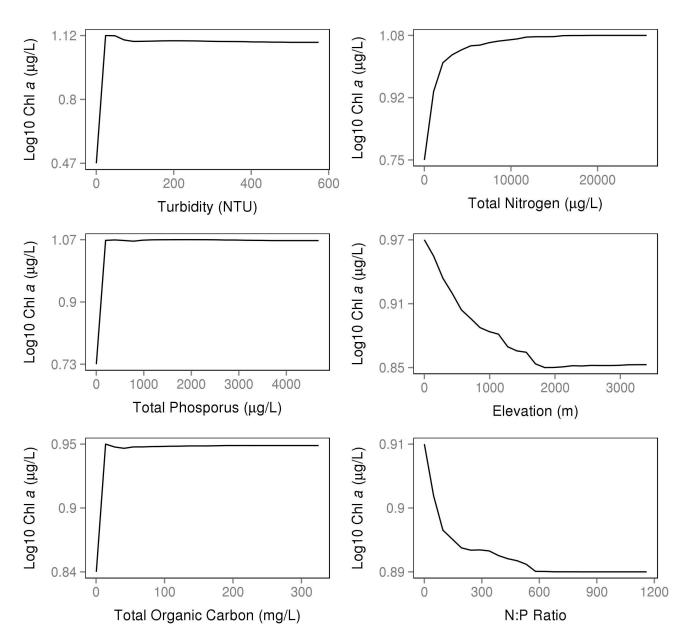


Figure 5: All Variables partial dependence plots for the top 5 most important variables.

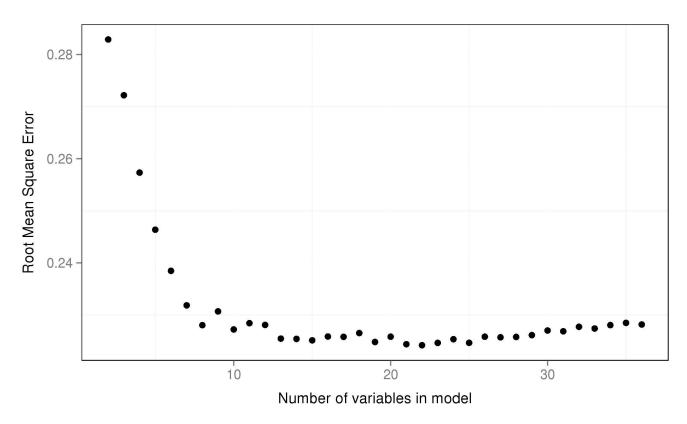


Figure 6: Variable selection plot for GIS only variables. Shows percent increase in mean squared error as a function of the number of variables.

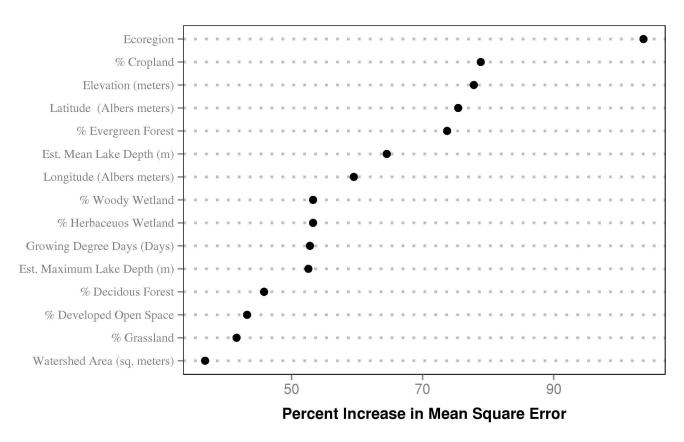


Figure 7: Importance plot for GIS Only Variables., shows percent increase in mean square error. Higher values of percent increase in mean squared error indicates higher importance.

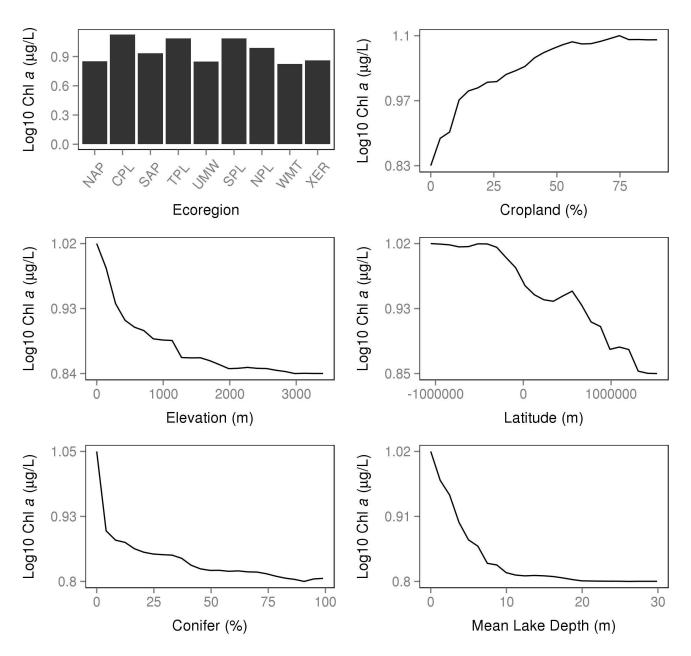


Figure 8: GIS Only Variables partial dependence plots for the top 5 most important variables.

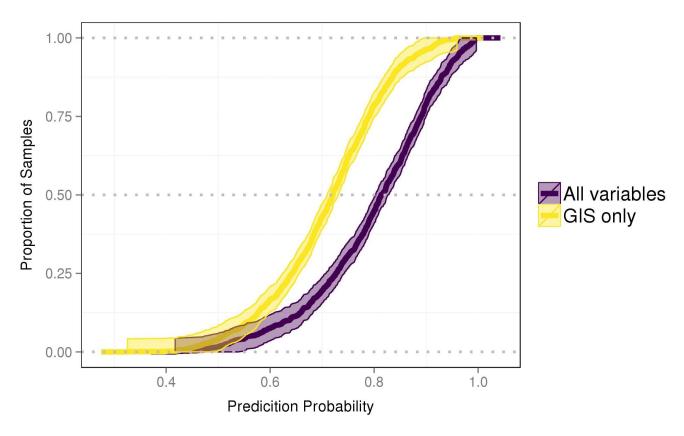


Figure 9: Prediction probabilities for the All Variables and GIS Only models.

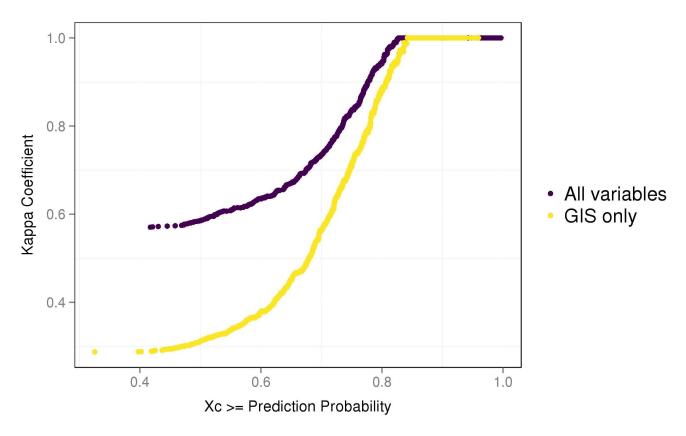


Figure 10: Accuracy of predictions as a function of lake prediction probability. The x-axis represents lakes with a prediction probability at a given level or higher.

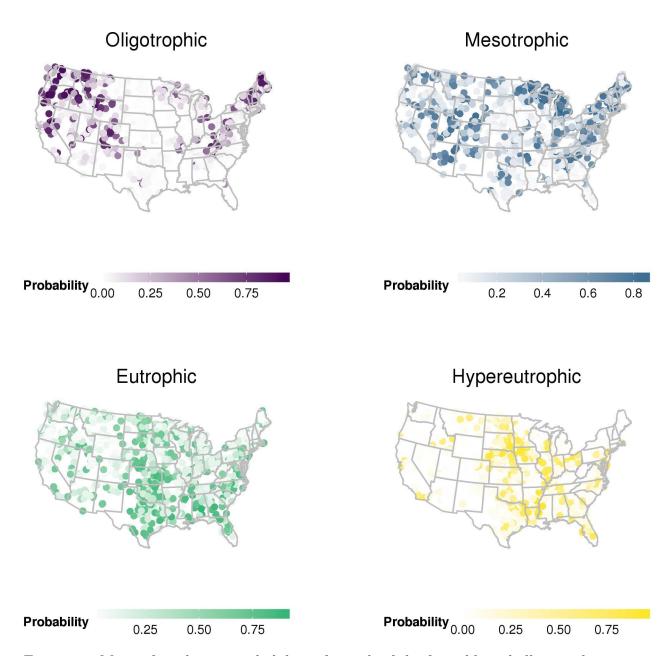


Figure 11: Maps of prediction probabilities for each of the four chlorophyll a trophic states

374 8 Tables

Table 1: Chlorophyll a based trophic state cut-offs.

| Trophic State (4 class) | Trophic State (2 class) | $\mu g/L$ Cut-off |
|-------------------------|--------------------------|-------------------|
| oligotrophic | oligotrophic/mesotrophic | <= 2 |
| mesotrophic | oligotrophic/mesotrophic | >2-7 |
| eutrophic | eutrophic/hypereutrophic | >7-30 |
| hypereutrophic | eutrophic/hypereutrophic | >30 |

Table 2: Random Forest confusion matrix for All Variables model converted to 4 trophic states. Columns show predicted values and rows show observed values. Agreement indicated on diagonal and accuracy for each trophic state indicated in 'Class Accuracy' column.

| | oligo | meso | eu | hyper | Class Accuracy (%) |
|-------|-------|------|-----|-------|--------------------|
| oligo | 115 | 31 | 0 | 0 | 78.77 |
| meso | 67 | 251 | 63 | 0 | 65.88 |
| eu | 7 | 61 | 217 | 75 | 60.28 |
| hyper | 0 | 5 | 29 | 159 | 82.38 |

Table 3: Random Forest confusion matrix for GIS Only model converted to 4 tropic states. Columns show predicted values and rows show observed values. Agreement indicated on diagonal and accuracy for each trophic state indicated in 'Class Accuracy' column.

| | oligo | meso | eu | hyper | Class Accuracy (%) |
|-------|-------|------|-----|-------|--------------------|
| oligo | 65 | 14 | 6 | 0 | 76.47 |
| meso | 101 | 213 | 98 | 18 | 49.53 |
| eu | 29 | 126 | 193 | 141 | 39.47 |
| hyper | 1 | 8 | 38 | 87 | 64.93 |

Table 4: Summary of relationship between prediction probabilities, total accuracy, and number of lakes for the All variables model.

| Prediction Prob. | Kappa Coefficient | Percent of Sample | Number of Samples |
|------------------|-------------------|-------------------|-------------------|
| All | 57 | 100 | 1080 |
| 0.50 | 59 | 98 | 1063 |
| 0.60 | 63 | 92 | 999 |
| 0.70 | 73 | 81 | 870 |
| 0.80 | 95 | 55 | 596 |
| 0.90 | 100 | 21 | 227 |

Table 5: Summary of relationship between prediction probabilities, total accuracy, and number of lakes for the GIS only model.

| Prediction Prob. | Kappa Coefficient | Percent of Sample | Number of Samples |
|------------------|-------------------|-------------------|-------------------|
| All | 29 | 100 | 1138 |
| 0.50 | 31 | 96 | 1091 |
| 0.60 | 38 | 83 | 949 |
| 0.70 | 56 | 57 | 651 |
| 0.80 | 88 | 22 | 247 |
| 0.90 | 100 | 4 | 43 |

9 Appendix 1. Variable Definitions

| Variable Names | Description | Source | Mean | Std. Error |
|--------------------------|--------------------------------|--------|----------|------------|
| AlbersX | Longitude (Albers meters) | GIS | 126757.1 | 34305.5 |
| AlbersY | Latitude (Albers meters) | GIS | 436908.1 | 17367.2 |
| BarrenPer_3000m | % Barren | GIS | 0.7 | 0.1 |
| BASINAREA | Watershed Area (sq. meters) | GIS | 3208.5 | 788.1 |
| CropsPer_3000m | % Cropland | GIS | 13.3 | 0.6 |
| DDs45 | Growing Degree Days (Days) | GIS | 2750.0 | 41.0 |
| DeciduousPer_3000m | % Decidous Forest | GIS | 17.1 | 0.6 |
| DevHighPer_3000m | % High Intensity Development | GIS | 0.4 | 0.0 |
| DevLowPer_3000m | % Low Intensity Development | GIS | 3.0 | 0.2 |
| ${\rm DevMedPer_3000m}$ | % Medium Intensity Development | GIS | 1.4 | 0.1 |
| DevOpenPer_3000m | % Developed Open Space | GIS | 5.4 | 0.2 |
| ELEV_PT | Elevation (meters) | GIS | 607.6 | 20.1 |
| EvergreenPer_3000m | % Evergreen Forest | GIS | 12.2 | 0.6 |
| FetchE | Fetch from East (m) | GIS | 1652.8 | 80.3 |
| FetchN | Fetch from North (m) | GIS | 2009.6 | 106.9 |
| FetchNE | Fetch form Northeast (m) | GIS | 1645.0 | 80.9 |
| FetchSE | Fetch from Southeast (m) | GIS | 1642.0 | 80.5 |
| $GrassPer_3000m$ | % Grassland | GIS | 13.8 | 0.7 |
| $HerbWetPer_3000m$ | % Herbaceuos Wetland | GIS | 1.7 | 0.1 |
| $IceSnowPer_3000m$ | % Ice/Snow | GIS | 0.0 | 0.0 |
| LakeArea | Lake Surface Area (sq. meters) | GIS | 12.2 | 2.3 |
| LakePerim | Lake Perimeter (meters) | GIS | 33.6 | 4.5 |
| MaxDepthCorrect | Est. Maximum Lake Depth (m) | GIS | 8.4 | 0.3 |
| MaxLength | Maximum Lake Length (m) | GIS | 2972.1 | 137.2 |

| Variable Names | Description | Source | Mean | Std. Error |
|----------------------------|---------------------------------------|--------|-------------|------------|
| MaxWidth | Maximum Lake Width (m) | GIS | 1567.5 | 76.0 |
| ${\it MeanDepthCorrect}$ | Est. Mean Lake Depth (m) | GIS | 2.9 | 0.1 |
| MeanWidth | Mean Lake Width (m) | GIS | 1370.1 | 122.6 |
| ${\it MixedForPer_3000m}$ | % Mixed Forest | GIS | 3.8 | 0.3 |
| PasturePer_3000m | % Pasture | GIS | 7.7 | 0.3 |
| PercentImperv_3000m | % Impervious | GIS | 2.6 | 0.2 |
| ShoreDevel | Shoreline Development Index | GIS | 2.7 | 0.1 |
| $ShrubPer_3000m$ | % Shrub/Scrub | GIS | 10.4 | 0.6 |
| VolumeCorrect | Est. Lake Volume (cubic meters) | GIS | 101211909.9 | 27438696.4 |
| WaterPer_3000m | % Water | GIS | 4.1 | 0.2 |
| $WoodyWetPer_3000m$ | % Woody Wetland | GIS | 5.2 | 0.3 |
| WSA_ECO9 | Ecoregion | GIS | NA | NA |
| ANC | Acid Neutralizing Capacity (ueq/L) | NLA | 2584.2 | 171.7 |
| ANDEF2 | Anion Deficit (ueq/L) | NLA | -506.4 | 143.2 |
| ANSUM2 | Sum of Anions using ANC (ueq/L) | NLA | 8043.1 | 1197.9 |
| BALANCE2 | Ion Balance (%) | NLA | -0.7 | 0.1 |
| CA | Calcium (ueq/L) | NLA | 1388.3 | 54.0 |
| CATSUM | Sum of Cations (ueq/L) | NLA | 7536.7 | 1105.0 |
| CL | Chloride (ueq/L) | NLA | 1600.3 | 438.2 |
| COLOR | Color (PCU) | NLA | 16.1 | 0.5 |
| CONCAL2 | Calculated Conductivity (uS/cm) | NLA | 949.0 | 148.1 |
| COND | Conductivity (uS/cm) | NLA | 656.0 | 72.6 |
| CONDHO2 | D-H-O Calculated Conductivity (uS/cm) | NLA | 618.6 | 55.1 |
| DATE_COL | Date Samples Collected | NLA | NA | NA |
| DEPTHMAX | Maximum Depth (meters) | NLA | 9.6 | 0.3 |

| Variable Names | Description | Source | Mean | Std. Error |
|----------------|---------------------------------|--------|--------|------------|
| DO2_2M | Dissolved Oxygen (mg/L) | NLA | 7.9 | 0.1 |
| DOC | Dissolved Organic Carbon (mg/L) | NLA | 8.6 | 0.5 |
| Н | Hydrogen Ions (ueq/L) | NLA | 0.2 | 0.1 |
| K | Potassium (ueq/L) | NLA | 245.6 | 40.6 |
| MG | Magnesium~(ueq/L) | NLA | 2190.4 | 282.2 |
| Na | Sodium (ueq/L) | NLA | 3709.7 | 816.3 |
| NH4 | Ammonium (mg/L) | NLA | 2.9 | 0.2 |
| NH4ION | Calculated Ammonium (ueq/L) | NLA | 2.5 | 0.2 |
| NO3 | Nitrate (ueq/L) | NLA | 5.4 | 0.7 |
| NO3_NO2 | Nitrate/Nitrite (mg N/L) | NLA | 0.1 | 0.0 |
| NPratio | Nitrogen:Phophorus Ratio | NLA | 34.5 | 1.8 |
| NTL | Total Nitrogen ($\mu g/L$) | NLA | 1109.9 | 56.4 |
| ОН | Hydroxide (ueq/L) | NLA | 3.1 | 0.2 |
| ORGION | Est. Organic Anions (ueq/L) | NLA | 85.9 | 4.8 |
| PH_FIELD | рН | NLA | 8.1 | 0.0 |
| PTL | Total Phosphorus ($\mu g/L$) | NLA | 103.1 | 7.8 |
| SIO2 | Silica (mg/L) | NLA | 8.6 | 0.3 |
| SO4 | Sulfate (ueq/L) | NLA | 3853.4 | 935.7 |
| SOBC | Sum of Base Cation (ueq/L) | NLA | 7534.1 | 1105.0 |
| TmeanW | Mean Profile Water Temp. (C) | NLA | 24.1 | 0.1 |
| TOC | Total Organic Carbon (mg/L) | NLA | 9.6 | 0.6 |
| TURB | Turbidity (NTU) | NLA | 12.3 | 1.0 |

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