Modelling Lake Trophic State: A Random Forest Approach

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Abstract

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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, and disservices (e.g. recreation, aesthetics, and 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 modern data mining approaches to predict trophic state. Using this 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 (i.e. GIS only) uses universally available GIS data only. The mean squared error was 0.22 and the adjusted R² was 0.48. The accuracy of various trophic state classifications derived from the chlorophyll a predictions ranged from 69% to 87% for the full model and from 49% to 75% 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 probabilites 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.

$_{\scriptscriptstyle 8}$ 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
- 37 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 also requires data that 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.

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 and 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.

To build on past work, we have identified several areas in which this research contributes. First, we build, asses, and compare two random forest models of chlorophyll a 1) in situ and universally available GIS data and then 2) universally available GIS data only. Second, we examine the important predictors for both models. Third, we examine the predictions for spatial patterns. Lastly, this paper, the code, and the data used in the models is made available as an R package from https://github.com/USEPA/LakeTrophicModelling.

$_{ ext{ iny 83}}$ 2 Methods

84 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 data were collected during the summer of 2007 and the final data were released in 2009

(USEPA 2009 for detailed description of methods). 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 water quality measurements and total cyanobacteria abundance from the National Lakes Assessment (USEPA 2009).

Adding to the monitoring data collected via the NLA, we use the 2006 NLCD data to examine landscapelevel drivers of trophic status in lakes. The NLCD is a national land use/land cover dataset that also
provides estimates of impervious surface. We calculated total proportion of each NLCD land use land
cover class and total percent impervious surface within a 3 kilometer buffer surrounding each lake
(Homer et al. 2004, Xian et al. 2009). A three kilometer buffer was selected as an intermediate measure
of the adjacent neighborhood; the three kilometer buffer size is greater than the immediate parcel but
smaller than regional and whole-basin measures.

To account for unique aspects of each lake and characterize lake productivity, we also used measures of lake morphometry (i.e. depth, volume, fetch, etc.). As these data are difficult to obtain for large numbers of lakes over broad regions, we used modeled estimates of lake morphometry (Hollister and Milstead 2010, Hollister et al. 2011, Hollister 2014). These included: surface area, shoreline length, Shoreline Development, Maximum Depth, Mean Depth, Lake Volume, Maximum Lake Length, Mean Lake Width, Maximum Lake Width, and Fetch.

110 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. A detailed discussion of the benefits of a random forest approach is beyond the scope of this paper. To find out more 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 be difficult 119 to interpret, thus selecting the most important variables is an important first step. Several method 120 have been proposed to do this with random forest. For instance, this is a problem often faced in gene 121 selection and in that field, a variable selection method based on random forest has been successfully 122 applied and implemented in the R Language as the varSelRF package (Díaz-Uriarte and De Andres 123 2006), but this is limited to classification problems. Additionally, others have suggested alternative 124 variable importance measures, but this is only needed with a large number of categorical variables which 125 are selected against with traditional random forest approach (Strobl et al. 2007). 126

In our case, we are predicting a continuous variable, chlorophyll a, directly thus varSelRF, does not 127 apply, and all of our variables are continuous so the approach suggested by Strobl (2007) is not necessary. 128 Thus we developed an approach, similar to varSe1RF but applied to random forest with regression trees. 129 With this approach we fit a full random forest model that includes all variables and a large number of 130 trees. We then rank the variables using the increase in mean square error, which has been shown to be 131 a less biased metric of importance than the mean decrease in the gini coeffecient (Strobl et al. 2007). Using this ranking, we then iterate through the variables and create a random forests with the top two 133 variables and record mean square error and adjusted R² of the resultant random forest. We then repeat 134 this process by adding the next most important variable in order of importance. With this information 135 we identify the top variables and the point at which adding variables does not improve the fit of the 136 overall model. These variables are selected and used as the "reduced model." With this method, a 137 minimum set of variables that maximizes model accuracy is provided. This allows us to start with a 138 full suite of predictor variables from which to select a minimum, easier to interpret set of variables. 139

140 2.3 Model Details

Using randomForest R package we ran models the predict chlorphyll a with two sets of predictors (Liaw and Wiener 2002); all precitors (in situ and universally available GIS predictors) and the GIS only variables (i.e. no in situ information). A listing 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 between models.

148 Our modelling work flow was as follows:

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- 1. Identify a minimal set of variables that maximize accuracy of the random forest algorithm. This minimal set of variables, the reduced model, is calculated for each of the models.
 - 2. Using R's randomForest package, we develop two random forest models.
- 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 and the two combinations of trophic state were used to highlight the possible error caused by misclassification of adjacent classes, such as mesotrophic and eutrophic (Table 1).

2.4 Measures of Model Performance and Variable Importance

We assessed the performance of the random forest two ways. First we compare the root mean square error and the adjusted R² of the models. Second, we examine the accuracy of the model when converted 158 to two different classifications of trophic state: a standard 4 class trophic state and 2 classes that 159 combine the high and low trophic states together (Table 1). We compare the two classifications via 160 a confusion matrix. A confusion matrix shows agreement and disagreement in a tabular form with 161 predicted values forming the columns of the matrix and observed values, the rows. From this tabulated 162 information we calculate the total accuracy (i.e. percent correctly predicted) and the kappa coefficient, 163 which takes into account the error expected by chance alone (i.e. the off diagonal values of the matrix) 164 (Cohen 1960, Hubert and Arabie 1985). The kappa coefficient can range from -1 to 1 with 0 equalling 165 the agreement expected by chance alone. Values greater than 0 represent agreement greater than would be expected by chance, with values greater than 0.61 considered "substantial" agreement (Landis and 167 Koch 1977). Negative values are rare and would indicate no agreement between the predicted and 168 observed values. Additionally, random forest builds each tree on bootstrapped, random subsets of the 169 original data, thus, a separate independent validation dataset is not required and random forest error

estimates are expected to be unbiased (Breiman 2001).

The random forest algorithm explicitly measures variable importance with two metrics:mean decrease 172 in Gini and percent increase in mean squared error. For each of these they measure the impact on 173 the overall model when that particular variable is included and thus can be used to assess importance 174 (Breiman 2001). The Gini Index has been shown to have a bias that is less apparent than with percent 175 increase in mean squared error (Strobl et al. 2007), thus, we use this metric to assess variable importance. 176 Lastly, partial dependence plots provide a mecahnism to examine the partial relationship between 177 individual variables and the response variable (Jones and Linder 2015). We examine these plots for the 178 top variables as assigned by percent increase in mean squared error of for each the reduced models. 179

180 3 Results

Our complete dataset includes 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 both across the four trophic state categories (Table 1) and spatially throughout the United States (Figure 1).

185 3.1 Models: All Variables

The model built with all predictors had a mean squared error of 0.21, 0.19, 0.17, 0.16, 0.15, 0.14, 0.13, 187 0.1, 0.1, 0.1, 0.1, 0.1, 0.09, 0.0188 0.09, 0.09189 0.09, 0.09190 0.09, 0.09191 0.09, 0.090.09, 0.09193 0.09, 0.09194 0.09, 0.09

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was 69\% and the kappa coefficient was 0.57 (Table 8). For the combined, two trophic states it was 0.87
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and 75% for accuracy and the kappa coefficient, respectively (Table 8).
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- 665 The reduced, all variables model was built using 1080 total observations. The variable selection process
- 666 identified 20 variables (Figure 2).
- The most important variables were ecoregion, growing degree days, and percent evergreen. (Figure ??).
- 668 MORE HERE.

669 3.1.1 Per Lake Probability

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671 3.2 Models: GIS Only Variables

The GIS only model had a mean squared error of 0.45, 0.44, 0.4, 0.37, 0.37, 0.36, 0.34, 0.34, 0.32, 0.32, 0.31, 0.3, 0.29, 0.29, 0.28, 0.28, 0.27, 0.27, 0.27, 0.27, 0.26, 0.26, 0.26, 0.26, 0.26, 0.26, 0.26, 0.25, 0.25, 0.25673 0.25, 0.25, 0.25, 0.25, 0.25, 0.25, 0.25, 0.24674 0.24, 0.24, 0.24, 0.23, 0.23, 0.23, 0.23, 0.23, 0.24, 0.24, 0.24, 0.24, 0.23675 $0.23,\ 0.23,\$ 0.23, 0.23677 0.23,
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The gis only variables model was built using 1138 total observations. The variable selection process for this model produced a reduced model with (Figure ??). The most important variables were ecoregion, growing degree days, and percent evergreen. (Figure ??). MORE HERE.

1204 3.2.1 Per Lake Probability

Historgram of Binned max prob.

1206 4 Discussion

1207 4.1 Trophic State Probabilities

One of the powerful features of random forests is the ability to aggregate a very large number 1208 of competing models or trees. Each tree provides an independent prediction or vote for a pos-1209 sible outcome. In the context of our trophic state models, we have 10,000 votes for each lake. 1210 These values may be interpreted as the probability that a lake is in a given trophic state. For 1211 instance, for a single lake (National Lake Assessment ID = NLA06608-0005), the vote prob-1212 abilities for Model 1 were "r 100 round(lakeVotes[lakeVotes\$COMID==23491387,][,4],2)"% for 1213 $oligotrophic, "r 100 \verb| round(lakeVotes[lakeVotes$COMID==23491387,][,5], 2)"\% ~for mesotrophic, "r 100 \verb| round(lakeVotes[lakeVotes[lakeVotes], 2)" ~for mesotrophic, "r 100 \verb| round(lakeVotes[lakeVotes], 2)" ~f$ 1214 $100 round (lake Votes | lake Votes | SCOMID = -23491387, | [,6], 2) \text{ ``\% for eutrophic, and '`r 100 } \text{ round (lake Votes | lake Votes | SCOMID | SCOMI$ 1215 for hypereutrophic. This suggests little uncertainty in the predicted oligotrophic state. 1216

Further, the maximum probability for each lake can be used as a measure of how certain the random 1217 forest model was of the prediction. We would expect higher total accuracy for lakes that had more 1218 certain predictions. Some lakes may have many votes for a single trophic state and few votes for 1219 other trophic states and these would thus have a large maximum probability and the random forest 1220 predictions would be more certain. Alternatively, the 10,000 votes could have been spread more equally 1221 across the trophic state classes for a lake and that lake would have a small maximum probability and 1222 the final predictions would be less certain. This should be evident by looking at the total classification 1223 accuracy of lakes given their maximum probability is above a certain point. To test this we can examine 1224 the accuracy of trophic state predictions across the full range of trophic state probabilities, similar to an 1225 approach outlined by Paul and MacDonald (2005) and implemented by Hollister et al. (2008). We utilize 1226 this approach and examine the change in total accuracy as a function of the maximum probability for 1227 each lake. As expected, lakes with higher maximum vote probabilities were more accurately predicted 1228 (Figure ??). The increasing trend suggests that even for models with lower overall accuracy there can 1229 also be a large number of individual cases that are predicted with high accuracy. 1230

1231 4.2 Variable Selection and Importance

There was a great deal of agreement on the important variables for each set of models. In line with past predictive modelling of cyanobacteria abundance and not surprisingly, the *in situ* models consistently select the water quality variables (turbidity, total nitrogen, total phosphorus, and N:P ratios) as important variables (Downing et al. 2001). While there is variation in the response of cyanobacteria to changes in relative nutrient concentrations, the general pattern suggests that limiting nutrients have considerable impact once amounts increase beyond expected levels.

The mechanistic role of turbidity on lake trophic state is more complex. Light availability in turbid waters is lower than in clear waters. This would suggest a negative relationship between turbidity and chlorophyll a. Second, chlorophyll a can also be a component of turbidity and lakes with higher chlorophyll a concentrations will also be more turbid. Last, chlorophyll a is not the only component of of turbidity and turbid waters can be caused by, for example, increased sediment loads or tannin. This would be a cause for concern with linear models; however, linearity is not an assumption of tree-based modelling approaches such as random forest.

Our models with the GIS-only variables captured the large scale spatial pattern of the trophic status gradient of lakes across the United States. We reliably saw latitude and longitude and ecoregion selected as important variables. It is also possible that other variables selected as important are also capturing a portion of this trend. For instance, elevation and growing degree days both have obvious spatial components, but may also be accounting for variation in temperature.

The land use/land cover variables were also important in describing trophic state patterns. Like elevation and growing degree days, broad scale spatial patterns are inherent in the data. For instance, the relative continental position of mountains in the United States is the spatial inverse of the distribution of agricultural lands. However, it is known that forests are positively associated with lower nutrient loads where as agricultural land shows a negative association. These more local scale relationships with land use/land cover likely provide additional predictive power to the information in the broader scale data.

Lastly, morphometry (e.g. depth and volume) also proved to be important in the prediction of lake trophic state. 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 of trophic state. As only depth and volume were selected, this likely shows the importance of in-lake nutrient processing and residence time.

4.3 Associating Trophic State and Cyanobacteria

Cyanobacteria biomass should be closely associated with trophic state as cyanobacteria contribute to the chlorophyll concentration in a lake. If these associations are strong enough we may be able to expand models such as those reported here to also predict probability of cyanobacteria blooms. To test if trophic state can be used to differentiate cyanobacteria abundance, we examine distribution of cyanobacteria abundance for each trophic state and also explored linear associations between chlorohyll and cyanobacteria abundance.

The distribution of cyanobacteria abundance showed separation between all of the trophic state classifications (Figures ??, ??, and ??) and there was a significant linear relationship (r^2 =0.33) between chlorophyll a and cyanobacteria abundance (Figure ??). Furthermore, Yuan et al. (2014) used the 2007 NLA to demonstrate that total nitrogen and chlorophyll a concentrations were good predictors of World Health Organization microcystin (a toxin produced by some cyanobacteria) criteria exceedences. These results suggest that trophic state is indeed an acceptable proxy for cyanobacteria abundance and that in lakes with higher trophic state it is also reasonable to expect higher cyanobacteria.

₂₇₄ 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. We were able to successfully predict a variety of trophic state classes. With the GIS only models our total accuracy ranged from , and with the full suite of data our model accuracy had a minimum accuracy of %.

While some of the models (i.e. Model 4) showed relatively low prediction accuracies, another feature of

the random forest, votes, can provide additional information. In addition to providing a single estimate of trophic state for each lake, our models also indicated the probability that a lake was classified in any of the categories. These probabilities may be mapped directly to show the uncertainty of a given predicted class. Furthermore, as the certainty of prediction increases, so does overall trophic state classification accuracy (Figure ??). These results suggest that our models will provide reasonable estimates of trophic state across the United States.

There was great deal of agreement on the important variables for each set of models. For the combined 1288 in situ and GIS models, the in situ water quality variables drove the predictions. This is expected. For 1289 the GIS only models, the results were more nuanced with three broad categories routinely being selected 1290 as important: broad scale spatial patterns in trophic state, land use/land cover controls of trophic state. 1291 and local, lake-scale control driven by lake morphometry. Lastly, associations between trophic state and 1292 cyanobacteria showed that, at the broad scale of the 2007 NLA, there is a linear relationship between 1293 chlorophyll a and cyanobacteria abundance and that using trophic state as a proxy for cyanobacteria 1294 has potential. 1295

These broad categories and the association between trophic state and total cyanobacteria abundance raise 1296 three important considerations related to managing eutrophication. First, the broad scale patterning 1297 suggests regional trends. This is important because it suggests that efforts to monitor, model and 1298 manage eutrophication and cyanobacteria should be undertaken at both national and regional levels. 1299 Second, while direct control of water quality in lakes would have a large impact, the land use/land 1300 cover drivers (i.e. non-point sources) of water quality are also important, and better management of 1301 the spatial distribution of important classes such as forest and agriculture can provide some level of 1302 control on trophic state and amount of cyanobacteria present. Third, in-lake processes (i.e. residence 1303 time, nutrient cycling, etc.) are, as expected, very important and need to be part of any management 1304 strategy. Building on these efforts through updated models, direct prediction of cyanobacteria, and 1305 additional information on the regional differences will help us get a better handle on the broad scale 1306 dynamics of productivity in lakes and the potential risk to human health from cyanobacteria blooms. 1307

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7 Figures

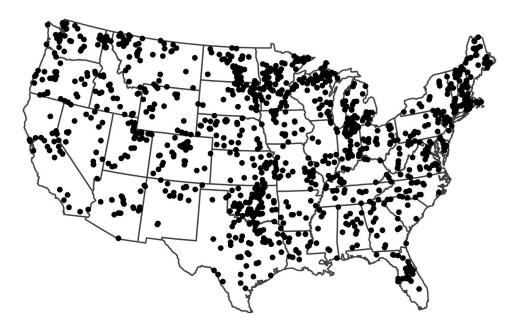


Figure 1: Map of the distribution of National Lakes Assesment Sampling locations

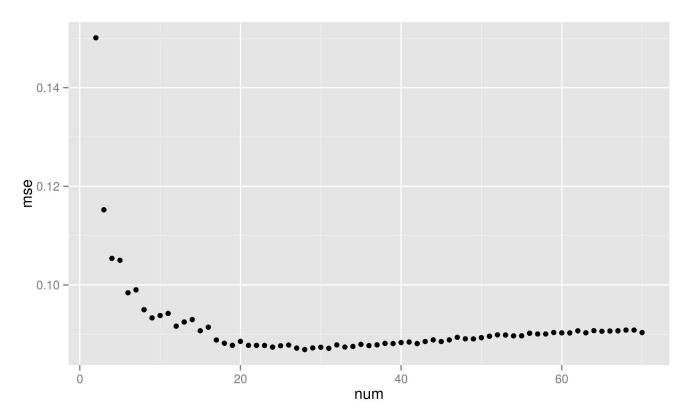


Figure 2: Variable selection plot for all variables. Shows percent increase in mean squared error as a function of the number of variables.

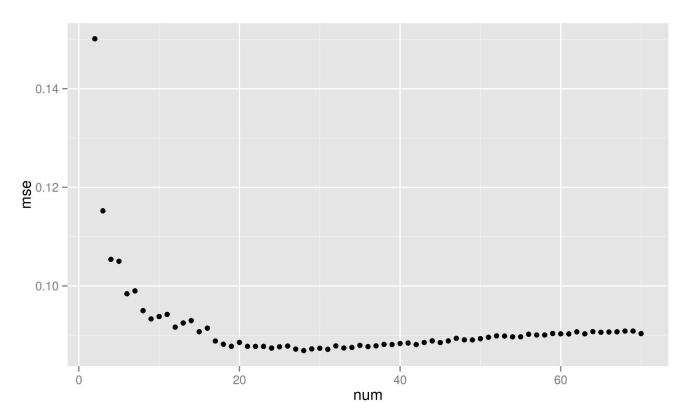


Figure 3: Variable selection plot for GIS only variables. Shows percent increase in mean squared error as a function of the number of variables.

1317 8 Tables

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

Trophic State (4 class)	Trophic State (2 class)	Concentration Cut-off
oligotrophic	oligotrophic/mesotrophic	<= 2
mesotrophic	oligotrophic/mesotrophic	>2-7
eutrophic	eutrophic/hypereutrophic	>7-30
hypereutrophic	eutrophic/hypereutrophic	>30

1318 :Random Forest confusion matrix for All Variables model converted to 4 trophic states. Columns show 1319 predicted values and rows show observed values. Agreement indicated on diagonal and accuracy for 1320 each trophic state indicated in 'Class Accuracy' column.

1321 :Random Forest confusion matrix for All Variables model converted to 2 trophic states. Columns show 1322 predicted values and rows show observed values. Agreement indicated on diagonal and accuracy for 1323 each trophic state indicated in 'Class Accuracy' column.

1324 :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.

1327 :Random Forest confusion matrix for GIS Only model converted to 2 tropic states. Columns show 1328 predicted values and rows show observed values. Agreement indicated on diagonal and accuracy for 1329 each trophic state indicated in 'Class Accuracy' column.

9 Appendix 1. Variable Definitions

variable_names	description	type
PercentImperv_3000m	Percent Impervious	GIS
$WaterPer_3000m$	Percent Water	GIS
$IceSnowPer_3000m$	Percent Ice/Snow	GIS
DevOpenPer_3000m	Percent Developed Open Space	GIS
DevLowPer_3000m	Percent Low Intensity Development	GIS
${\rm DevMedPer_3000m}$	Percent Medium Intensity Development	GIS
${\rm DevHighPer}_3000{\rm m}$	Percent High Intensity Development	GIS
BarrenPer_3000m	Percent Barren	GIS
${\bf DeciduousPer_3000m}$	Percent Decidous Forest	GIS
$Evergreen Per_3000 m$	Percent Evergreen Forest	GIS
${\it MixedForPer_3000m}$	Percent Mixed Forest	GIS
$ShrubPer_3000m$	Percent Shrub/Scrub	GIS
$GrassPer_3000m$	Percent Grassland	GIS
$Pasture Per_3000m$	Percent Pasture	GIS
${\rm CropsPer_3000m}$	Percent Cropland	GIS
$WoodyWetPer_3000m$	Percent Woody Wetland	GIS
$HerbWetPer_3000m$	Percent Herbaceuos Wetland	GIS
AlbersX	Longitude	GIS
AlbersY	Latitude	GIS
LakeArea	Lake Surface Area	GIS
LakePerim	Lake Perimeter	GIS
ShoreDevel	Shoreline Development Index	GIS
DATE_COL	Date Samples Collected	Water Quality
WSA_ECO9	Ecoregion	GIS
BASINAREA	Watershed Area	GIS
DEPTHMAX	Maximum Depth	Water Quality

variable_names	description	type
ELEV_PT	Elevation	GIS
DO2_2M	Dissolved Oxygen	Water Quality
PH_FIELD	pH	Water Quality
COND	Conductivity	Water Quality
ANC	Acid Neutralizing Capacity	Water Quality
TURB	Turbidity	Water Quality
TOC	Total Organic Carbon	Water Quality
DOC	Dissolved Organic Carbon	Water Quality
NH4	Ammonium	Water Quality
NO3_NO2	Nitrate/Nitrite	Water Quality
NTL	Total Nitrogen	Water Quality
PTL	Total Phosphorus	Water Quality
CL	Chloride	Water Quality
NO3	Nitrate	Water Quality
SO4	Sulfate	Water Quality
CA	Calcium	Water Quality
MG	Magnesium	Water Quality
Na	Sodium	Water Quality
K	Potassium	Water Quality
COLOR	Color	Water Quality
SIO2	Silica	Water Quality
Н	Hydrogen Ions	Water Quality
ОН	Hydroxide	Water Quality
NH4ION	Calculate Ammonium	Water Quality
CATSUM	Cation Sum	Water Quality
ANSUM2	Anion Sum	Water Quality
ANDEF2	Anion Deficit	Water Quality
SOBC	Base Cation Sum	Water Quality

variable_names	description	type
BALANCE2	Ion Balance	Water Quality
ORGION	Estimated Organic Anions	Water Quality
CONCAL2	Calculated Conductivity	Water Quality
CONDHO2	D-H-O Calculated Conductivity	Water Quality
TmeanW	Mean Profile Water Temperature	Water Quality
DDs45	Growing Degree Days	GIS
MaxLength	Maximum Lake Length	GIS
MaxWidth	Maximum Lake Width	GIS
MeanWidth	Mean Lake Width	GIS
FetchN	Fetch from North	GIS
FetchNE	Fetch form Northeast	GIS
FetchE	Fetch from East	GIS
FetchSE	Fetch from Southeast	GIS
MaxDepthCorrect	Estimated Maximum Lake Depth	GIS
VolumeCorrect	Estimated Lake Volume	GIS
MeanDepthCorrect	Estimated Mean Lake Depth	GIS
NPratio	Nitrogen:Phophorus Ratio	Water Quality

1331 References

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