Article

The Use of Three-Dimensional Convolutional Neural Networks to Interpret LiDAR for Forest Inventory

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**Abstract:** As light detection and ranging (LiDAR) technology becomes more available, it has become common to use these datasets to generate remotely sensed forest inventories across landscapes. Traditional methods for generating these inventories employ the use of height and proportion metrics to measure LiDAR returns and relate these back to field data using predictive models. Here we employ a three-dimensional convolutional neural network (CNN), a deep learning technique that scans the LiDAR data and automatically generates useful features for predicting forest attributes. We test the accuracy in estimating forest attributes using the three-dimensional implementations of different CNN models commonly used in the field of image recognition. Using the best performing model architecture, we compared CNN performance to models developed using traditional height metrics. The results of this comparison show that CNNs produced 12 % less prediction error when estimating biomass, 6 % less in estimating tree count, and 2 % less when estimating the percentage of needleleaf trees. We conclude that CNNs can be a more accurate means of interpreting LiDAR data for forest inventories than standard approaches.

**Keywords:** Deep learning, Artificial neural network, Machine learning, ALS, LiDAR, Enhanced forest inventory, Area-based

1. Introduction

Over the past two decades, airborne Light Detection and Ranging (LiDAR) has become an invaluable tool for remotely quantifying the three-dimensional structure of the forest canopy. This has enabled scientists to estimate forest attributes over large areas traditionally only measurable with an intensive field campaign. Attributes predicted with LIDAR include estimates of tree count, height, species, stem volume, and aboveground biomass [1-3]. Such enhanced forest inventories (EFIs) have proven useful for research and applications in forest and wildlife ecology, forest carbon cycling, and sustainable forest management [4-6].

The typical methodology used for developing EFIs is called the *Area Based* (AB) approach. The AB approach utilizes predictive modelling to associate plot-based field measurements with explanatory variables derived from various measures of a LiDAR dataset of the same forest area. These models are then applied to make estimates of forest attributes for new areas without field measurements [7]. Numerous statistical techniques have been used to develop AB models, among which linear mixed modelling and random forest imputation are common modelling tools [8, 9].

Model explanatory variables derived from LiDAR data can take several forms, but mostly constitute either measurements of point height or distribution along the vertical stratum [1, 2, 10]. Height measurements typically constitute summary statistics that calculate the mean, maximum, and percentile heights of points within each grid cell. Distribution metrics quantify the proportion of points found above certain height thresholds. Since their introduction in the early 2000s, these traditional height metrics (THMs) have been proven to be effective predictors for modeling forest attributes [11, 12]. Common software packages currently used for LiDAR EFI modeling, including FUSION [13] and rLiDAR [14], are based on extracting THMs from point clouds.

Despite the well-established effectiveness of THMs, these metrics come with several drawbacks. Many THMs are highly correlated with one another, and so care must be taken during model development to avoid issues of multicollinearity [15]. Furthermore, THMs generated from one LiDAR data set are sometimes not stable when applied to another due to variation in acquisition parameters such as laser penetration, pulse density, and scan angle [16-18]. Finally, THMs do not quantify measures of horizontal complexity, and so the nuance of distinct tree crown shapes, easily observable by the human eye, is lost. This may partly account for the difficulty in estimating tree count from THMs, as reported in previous studies [19-21].

Like LiDAR modelers, computer vision scientists have struggled to develop meaningful tools for quantifying remotely sensed data. Early computer vision scientists developed a series of metrics for estimating an image’s content using summary statistics of color histograms, detected edges, and blobs [22]. This methodology is known as feature engineering, and bears some resemblance to LiDAR’s THMs. Feature engineering, however, has largely since been discarded by the computer vision field in favor of deep learning, which can learn to self-identify important spatial features in a dataset [23].

Deep learning is a form of machine learning, and refers primarily to artificial neural networks of a sufficient complexity so as to interpret raw data without the need for human-derived explanatory variables [23]. This differs from simpler neural networks (such as perceptrons) which impute attributes using features such as THMs, and have been used to estimate forest attributes in remote sensing for many years [24-26]. Convolutional neural networks (CNNs), such as those employed here, are a subdivision of deep learning and are distinct in that they interpret spatial data by scanning it using a series of trainable moving windows. The values of these windows are initially randomized. During training, the CNN uses an optimizer to tune these values to identify useful features and objects for estimating the response variable. This is accomplished without user input.

The first CNN was developed in 1995 to classify images of hand-written digits [27]. However, the technique was largely underrepresented in scientific literature for the following decade, partially owing to computational and software constraints [23]. This changed in 2012, when a deep CNN (consisting of many layered convolutions) won the ImageNet image classification competition by a wide margin [28, 23]. Since then, CNNs of increasing complexity and depth have consistently outperformed models based upon feature extraction for computer vision [29-32]. Other variants, such as inception and residual models, have also been developed to extract greater numbers of meaningful features by using windows of varying sizes and preserve useful input data as it passes through the model [33].

Though CNNs have been mostly developed for computer vision with two-dimensional images, those in other fields are beginning to apply these algorithms to novel, three-dimensional problems. A few studies have begun using deep learning for measuring and analyzing forest attributes. For example, Guan et al. [34] used a segmentation technique to isolate tree crowns, and then used a neural network to classify species based on point distribution. In another study, Ghamisi et al. [35] applied a 2D CNN to estimate forest attributes from rasterized LiDAR and hyperspectral data. A 2D CNN is designed to scan two-dimensional images, and is only capable of identifying spatial features along two axes. A 3D CNN uses three-dimensional windows to scan volumetric space, and identify spatial features in the X, Y, and Z axes. In this context, a 2D CNN is only capable of scanning a height map derived from a LiDAR point cloud, whereas a 3D CNN is capable of scanning the entire cloud for 3D features such as tree crowns, stems, or branches.

Presently 3D CNNs have not been used to interpret LiDAR data for forest inventory; however there are examples of their use in other fields. One study implemented a 3D CNN for use in airborne LiDAR to identify helicopter landing zones in real time [36]. Others have used them in conjunction with terrestrial LiDAR to map obstacles for autonomous cars [37, 38]. One common application has been to identify malignancies using 3D medical scans [39-41]. Several studies have also used 3D CNNs for household object detection [42-44].

The goal of this study is to adapt common 2D CNN implementations to scan the 3D volumetric pixels, or voxels, derived from a LiDAR point cloud to estimate aboveground biomass, tree count, and the percent of needleleaf stems. Several CNN architectures were adapted, beginning with the least complex (LeNet) [27], and working towards deeper, more contemporary architectures (Inception V3) [45]. We compare these architectures against one another, noting performance with model complexity and type. The best performing CNNs were then tested against random forest models trained using THM.

2. Materials and Methods

3.2. *Field Data*

Deep learning is renowned for requiring extremely large datasets in order to calibrate properly. Training datasets often consist of thousands, if not millions, of training examples [46]. To meet this requirement, forest inventories from eight different sites across the Northern New England/ Acadian forest region were aggregated (Table 1). Each site often contained several inventories conducted by different agencies. We note that the models we developed are not intended to represent the entire landscape, and serve only as comparisons between model types.

**Table 1.** Field inventory sites used for model training as well as the origin of the accompanying LiDAR.

|  |  |  |  |
| --- | --- | --- | --- |
| **Site Name** | **Location** | **Number of plots** | **LiDAR Acquisition** |
| Acadia National Park | 44° 20' N 68° 16' W | 9 | G-LIHT |
| Bartlett Experimental Forest | 44° 2' N 71° 17' W | 46 | NEON |
| University of Maine’s Cooperative Forest Research Unit | 45° 11' N 69° 42' W | 90 | G-LIHT |
| University of Maine’s Dwight B. Demeritt Forest | 44° 55' N 68° 40' W | 344 | G-LIHT |
| Harvard Forest | 42° 31' N 72° 11' W | 13,470 | G-LIHT and NEON |
| Holt Research Forest | 43° 52' N 69° 46' W | 2,002 | G-LIHT |
| Howland Research Forest | 45° 12' N 68° 45' W | 1,014 | G-LIHT |
| Penobscot Experimental Forest | 44° 52' N 68° 37' W | 301 | G-LIHT |

All inventories consisted of stem-maps of varying sizes, on which all trees with diameters greater than 11.4 cm were measured. In some inventories, heights were measured on only a subset of trees, and so localized non-linear height-diameter models were generated following a Chapman-Richards model form to impute tree height. Uncommon trees were grouped with similar species. Some inventories were measured up to five years prior to the LiDAR being acquired over that site. In these cases, trees were projected forward in time using the Forest Vegetation Simulator’s Acadian Variant [47]. Table 2 contains summary statistics of field inventory data. Several plots lay entirely within canopy gaps or non-forested wetlands, and so had no trees. These were retained to better constrain the models.

**Table 2.** Summary statistics of the field inventory data used in this study.

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | Min | Mean | Max | SD |
| Aboveground Biomass (Mg/ha) | 0 | 185.7 | 715.7 | 88.5 |
| Tree Count (#) | 0 | 6.93 | 25 | 3.65 |
| Percent Needleleaf (%) | 0 | 42.9 | 100 | 28.9 |

The stem maps were segmented into 10 x 10 m grid cells, a size small enough to amplify the amount of training data, and large enough to include several whole tree crowns. We chose to model aboveground biomass, tree count, and the percent of needleleaf stems (including cedars) as the performance of these models may be indicative of other attributes which measure size, number, and types of trees. Biomass was estimated for each tree using the component ratio method developed for the US Forest Inventory and Analysis program [48]. This method makes use of species-specific localized stem volume and component equations, which are then converted to biomass using specific gravity and a site modifier.

In order to account for edge effects given the small plot size, regional diameter to crown width equations were used to project the area of each tree’s crown in space [49]. Tree-level biomass was then multiplied by the proportion that each tree’s crown overlapped the plot. Thus, trees were treated as areas containing biomass, rather than points with a binary in/out classification. All field-measured plots were larger than the 10 m grid cells used in this study, so the biomass proportion of trees with stems outside of the grid cells were also accounted for, assuring unbiased predictions across multiple cells.

One common technique for increasing sample size in deep learning model applications is to transform or rotate input images multiple times, and then calibrate a model using all of these quasi-novel inputs [50, 51]. In a similar vein, there were instances in which two LiDAR acquisitions overlapped a single plot, and we included both in the models. The precise configuration of LiDAR returns will always vary from one LiDAR acquisition to the next. In some instances plots were allowed to overlap one another by a maximum of 25 %. This allowed more 10 m cells to fit into many of the stem mapped areas, while retaining novel assemblages of trees. A total of 17,537 field plots were generated for training and analysis in this study.

3.1. *LiDAR Data*

Airborne LiDAR data were acquired over the study areas in a series of flights conducted by two entities. Data over 10,627 of the 17,537 field plots at seven sites were collected by NASA Goddard’s LiDAR, Hyperspectral, and Thermal Imager (G-LIHT) [52]. The G-LIHT sensor suite operates a Riegl VQ-480 LiDAR sensor with a 300 kHz scan rate and a 1550 nm wavelength. The average pulse density ranged from 12-15 pls/m2. The remaining 6910 plots were flown by the National Ecological Observatory Network (NEON) using an Optech Gemini with a 167 kHz scan rate and a 1064 nm wavelength. The average pulse density was 9 pls/m2. All LiDAR data were acquired between 2012 and 2016 in leaf-on conditions between June and August.

The LiDAR data were preprocessed first by normalizing points using a digital elevation model so as to measure height above ground level. Next, the point clouds were clipped to the extent of the field plots, representing the 10 x 10 m grid cells. The vertical space that we measured on each plot extended 35 m above the ground, an elevation slightly higher than the tallest tree in the field data. The point cloud was then binned into voxels measuring 25 x 25 x 33 cm, the dimensionality of the voxelized point cloud was thus 40 x 40 x 106. We used rectangular voxels in order to reduce the dimensionality of the vertical axis, and prevent horizontal features from being lost. The final dimensionality of the voxels was determined through qualitative testing of several sizes using the most simple model form (LeNet). Each voxel was assigned the value equal to the number of points occurring within that 3D space. The input to the neural network was therefore a matrix of those dimensions, upon which the three-dimensional kernels of a CNN could be passed over.

3.3. *Network Architecture*

Convolutional neural networks consist of a series of data transformations, or layers, which can be arranged like building blocks in order to form specific network architectures. The following layer types were used in various combinations for the construction of the CNNs tested in this study:

* *Convolutional layers* are a series of moving windows passed over data with spatial or temporal relationships. The values of these windows are used as multipliers and are initially randomized, but are defined over time as the network trains. Often these are used to identify features such as edges.
* *Dropout layers* are commonly applied following a convolution. These consist of randomized removals of data. This can prevent overfitting of the network by altering architecture during the training process, preventing specific pathways from becoming relied upon [53].
* *Batch normalization* is another layer frequently applied after a convolution, and is used to standardize input data. This can speed up model training by ensuring data fall upon the same scale from which weights are initially randomly derived. These shift subtly with each training step, and so are another way of preventing model overfitting [54].
* *Activation layers* are an essential component to any neural network. They act as thresholds for data passage onto the next layer, similar to the action potential in a living neuron. The most common type of activation function is the rectified linear unit (ReLU), which is a linear gateway allowing only data with a value greater than zero to pass.
* *Pooling layers* are spatial aggregations of data. These are used primarily to reduce dimensionality and condense important information. Maximum pooling is the most common type of pooling, which takes the largest data value inside a moving window [55].

We tested five CNN architectures across a range of model complexity, assessing their performance using the coefficient of determination (pseudo-R2), root mean squared error (RMSE), and bias. Each had to be adapted to measure 3D voxel space rather than 2D pixels. This conversion altered the dimensionality from each architecture’s native format, and so we attempted to replicate each architecture’s proportional changes in dimensionality using convolutions and pooling. For example, if an architecture called for a 5 x 5 kernel over a 30 x 30 image, we would scale our 3D window to approximately the same proportion relative to the input, 7 x 7 x 18. This was also balanced with the dimensionality required for the following layer.

1. The first network architecture we tested was *LeNet*, the earliest CNN [27]. LeNet consists of two layers of convolutions each followed by maximum pooling and dropout. These feed into a further two fully connected layers (consisting of a ReLU activator and dropout) before producing a prediction.
2. The second network we tested was *AlexNet* [28]. First introduced in 2012, this network is largely credited as the first ‘deep’ CNN, and it resulted in a dramatic increase in image classification accuracy. AlexNet consists of five layers of convolutions, each with an activator, with pooling following the first, second, and fifth convolutions. These lead into a further three fully connected layers and a final prediction.
3. Next we implemented *GoogLenet*, introduced in 2014 [29]. GoogLeNet improved upon image classification performance by introducing groups of convolutional layers called inception layers. Inception layers are bundles of convolutions of varying sizes and strides, useful for identifying different types of features. These are followed by batch normalization and activation, with the outputs concatenated. GoogLenet begins with three convolutional layers and pooling, and is followed by nine inception layers, with pooling after the second and seventh inception layer. The data are then funneled into a single fully connected output to achieve a prediction.
4. A series of networks succeeded GoogLenet, including *Inception-V3* [45]. This expanded upon GoogLenet’s inception layers, adding additional convolutions with four pathways by which the incoming data is analyzed before being concatenated. Inception-V3 begins with five convolution layers and two pooling layers, followed by eleven inception layers of varying compositions, and pooling layers interspersed along the way to continuously reduce dimensionality.
5. The final network architecture we tested was *ResNet-50* [31]. Rather than use inception layers, ResNet-50 uses residual layers. Data entering a residual layer is subject to several convolutions, batch normalization, and activation, the result of which is added to the original input. By retaining the input values, useful information from the previous residual layers is preserved and improved upon, reducing the potential for values running through the network to drift. Our implementation of ResNet-50 contained an initial convolution, ReLU, and pooling layer, followed by sixteen residual layers (each containing three convolutions) with three pooling layers interspersed to gradually reduce dimensionality.

Each of the neural networks were implemented using Google’s Tensorflow library [56]. Of the 17,537 total plots, we randomly withheld 2,000. As is typical in many deep learning applications, we segmented the data into three datasets. The largest dataset was comprised of 15,537 plots and was used to train the model. A validation dataset comprised of 1,000 withheld plots was used to determine the optimal stopping point of model training; which was the point at which mean squared error of these data no longer decreased. A third dataset called the testing data was comprised of another 1,000 withheld plots, and was used to assess model performance, as the independence of the validation dataset was compromised upon its use. In instances for which withheld plots had additional LiDAR flights over them, the additional LiDAR data were discarded.

Performance of each CNN was assessed by calculating the root mean squared error (RMSE) and bias against each independent variable of the testing dataset. In order to settle on the optimal architecture, we first trained each of the five models to estimate aboveground biomass. We selected the best performing model and then used the same architecture to train the tree count and percent needleleaf models. The assumption for this being that the architecture which best predicted biomass was also the architecture which best interpreted LiDAR features.

3.4. *Traditional Height Metrics Modelling*

Models trained using the optimal CNN architecture were compared to models trained using both linear mixed modelling and random forest in regression mode with THMs. We withheld the same 2,000 plots as before, once again separating out those for training, validation, and testing data. The THMs were generated for each plot using the rLiDAR package for the R programming language [14]. These metrics included measures of height percentiles, height summary statistics, and percentages of returns above thresholds..

For the purposes of model selection, we first generated small random forest models consisting of only 50 decision trees for each of the three forest attributes. We then generated importance values for each THM covariate using conditional variable importance, which is less prone to importance inflation due to correlated variables [57].

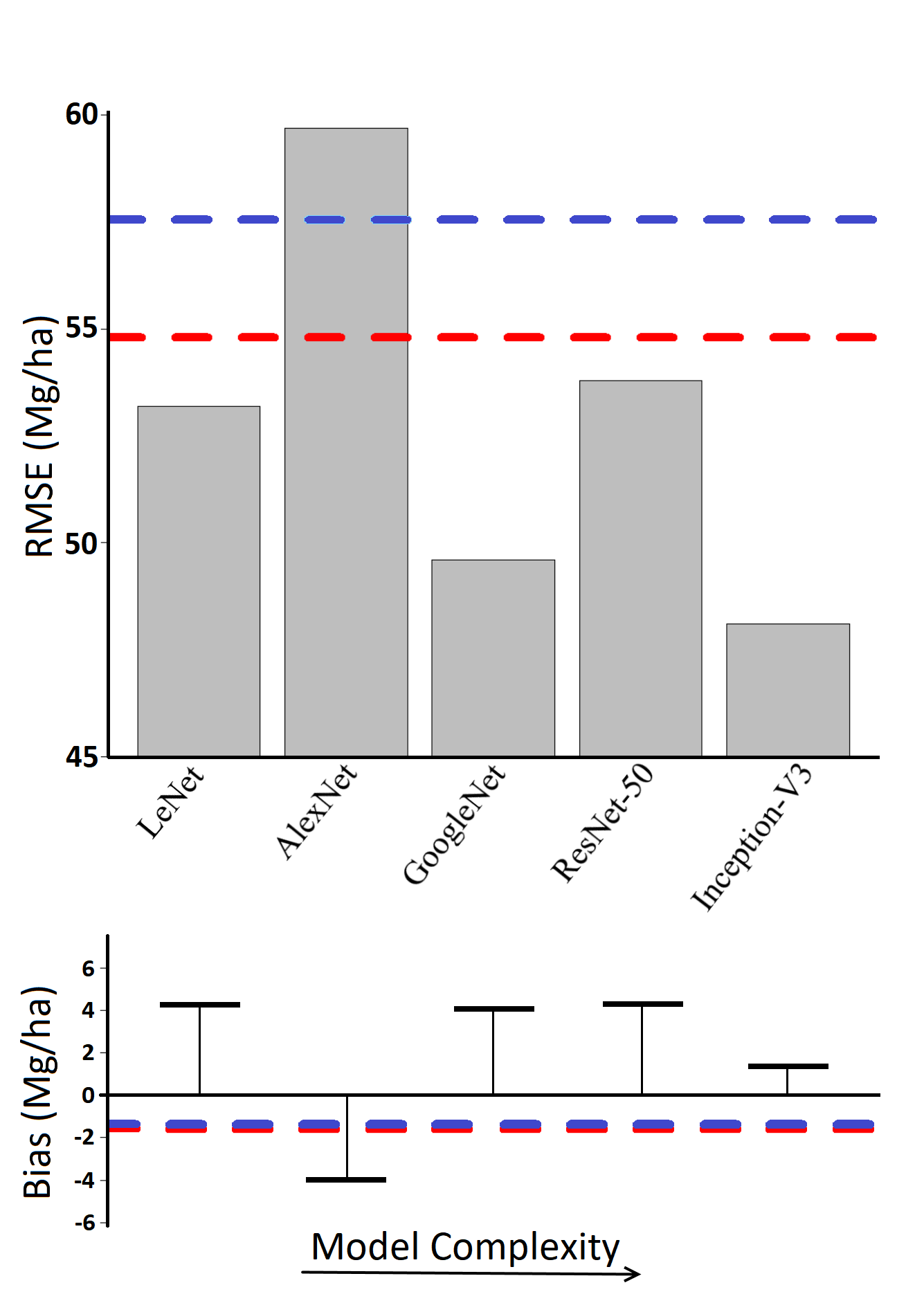
Using the top ten most important predictors of each attribute, we developed generalized linear mixed models with site as a random effect. We then performed reverse stepwise regression to select a final model form [58]. Percent needleleaf models were developed using a logistic form. The withheld testing data were used to assess model performance.

Next we developed new random forest models. Random forest can occasionally benefit from variable selection in instances in which there are a large number unimportant variables [59]. Using the preliminary importance values, we removed 15 % of the variables with the lowest importance and then re-ran the model. We continued this process until all variables had been removed, and then used the validation data to select the model form with the lowest RMSE [59]. Several variables were removed from the biomass and tree count models, none were removed from the percent needleleaf model. New models were then trained using these variables with 1000 decision trees. The withheld testing data were again used to assess model performance.

All models were graphically examined using one-to-one plots between observed field measurements and model predictions. In addition to a one-to-one line, loess regression was to fit a moving trendline to this data. For this loess trend line, we used second order polynomials for the fit, and used 66 % of observation to smooth the line at each value [60, 61].

**3. Results**

We first compared results in terms of the RMSE and bias of aboveground biomass predictions among each of the five CNN architectures relative to the models trained using the THMs (Figure 1). Four of the five architectures we tested outperformed the both the linear mixed model with traditional height metrics (LMM-THM) and the random forest model with traditional height metrics (RF-THM) in terms of RMSE. Only the Inception-V3 CNN exhibited less absolute bias than the THM-RF model, and equal absolute bias of the LMM-THM.



**Figure 1.** A comparison of biomass model performance by architecture. Results are shown in terms of root mean square error (RMSE) and bias. Red lines represent the performance of the random forest model trained on traditional height metrics. Models are listed in order of complexity, which refers to the number of trainable parameters in the model. Note that the y-axis begins at a RMSE of 45 Mg/ha to better highlight differences in model performance.

The best performing CNN architecture was Inception-V3, which was also the most complex CNN, with the greatest number of layers and over 20 million trainable parameters. Inception-V3 had a RMSE of 48.1 Mg/ha, representing an 11 % decrease in error from 54.1 Mh/ha with the RF-THM model, and a 16.5 % decrease from the 57.6 Mg/ha RMSE in the LMM-THM model. The absolute bias of Inception-V3 was lower than the RF-THM and equal to the absolute bias of the LMM-THM model, at 1.3 Mg/ha. This was 68 % lower than the next best performing CNN architecture, GoogleNet with 4 Mg/ha bias. The architectures using inception layers (GoogleNet, and Incepton-V3) were the top two performing architectures in terms of lowest RMSE and bias.

Of the five CNN architectures, only AlexNet performed worse in terms of RMSE than the LMM-THM and RF-THM models. AlexNet’s RMSE and bias were 59.7 and -3.9 Mg/ha respectively. The next-worst performing CNN was ResNet-50, with a RMSE and bias of 53.8 and 4.2 Mg/ha, respectively. Models using residual layers have garnered some degree of popularity in recent years, however these results indicate that they may not be as effective at quantifying LiDAR, or at the very least require more fine tuning. LeNet performed slightly better than ResNet-50 in terms of error, and equally in terms of bias despite its relative simplicity (53.3 Mg/ha and 4.2 Mg/ha).

Based on these results for modeling biomass, Inception-V3 models were assumed to be the best for interpreting LiDAR data, and were developed for the other two desired forest attributes: tree count and percent of needleleaf trees. Results of those models are shown in Table 3. Each of the Inception-V3 CNNs outperformed the LMM-THM and RF-THM models in terms of RMSE, however the improvement in accuracy was less pronounced for both the tree count and percent needleleaf models than with the biomass estimation. Random forest models consistently had a lower RMSE and higher coefficient of determination (pseudo R2) than the linear mixed models, while the linear mixed models had slightly lower absolute bias than random forest.

The CNN model predicting tree count resulted in 6 % less error than the RF-THM model and 10 % less RMSE than the LMM-THM, with a RMSE of 2.78 trees. However, the CNN model’s bias was double that of THM-RF, at 0.2 and 0.1 trees, respectively. The LMM-THM exhibited almost no overall bias in estimating tree count (.03 trees). With a 10 x 10 m cell size, these biases represent between 3 - 20 trees/ha, which is a relatively low bias for all three models (0.5 – 2.8 %) when taking into consideration that the mean value of these plots was 714 trees/ha.

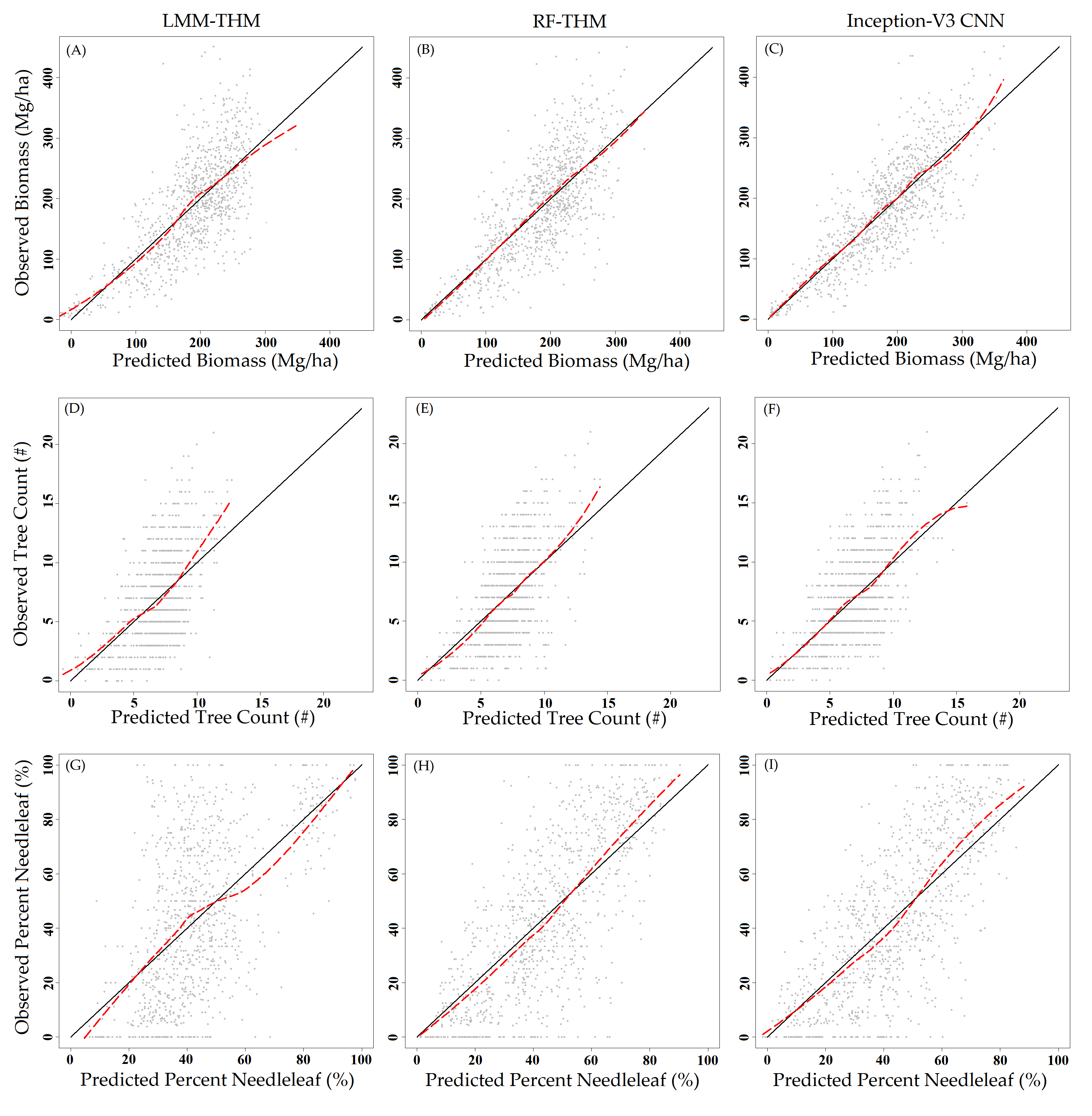
The CNN model predicting percent needleleaf had a RMSE of 18.7 %, which is 2 % less than the RF-THM model’s RMSE of 19.1 %, and 22 % less than the LMM-THM model’s RMSE of 24.1 %. The percent needleleaf CNN also had 60 % less bias than the RF-THM and 50 % less absulute bias than the LMM-THM, with a bias of 0.2 %. Once again it should be noted that the overall biases of all three percent needleleaf models are only several tenths of a percent, and thus are negligible in any practical context.

**Table 3.** Performance of each of the Inception-V3 models alongside random forest and linear mixed models (LMM) using traditional height metrics. Root Mean Square Error (RMSE) and bias were used to quantify performance. When possible, error and bias as a percent of each attribute’s mean are shown in parentheses.

|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
|  | Aboveground Biomass (Mg/ha) | | | Tree Count  (#) | | | Percent Needleleaf  (%) | | |
|  | Pseudo R2 | **RMSE**  **(%)** | **Bias**  **(%)** | Pseudo R2 | **RMSE**  **(%)** | **Bias**  **(%)** | Pseudo R2 | **RMSE** | **Bias** |
| THM with  LMM | 0.55 | 57.6  (31.3) | -1.3  (0.7) | 0.31 | 3.10  (44.9) | 0.03  (0.5) | 0.31 | 24.1 | -0.4 |
| THM with  Random Forest | 0.61 | 54.1  (29.5) | -1.6  (0.9) | 0.36 | 2.96  (43.0) | 0.1  (1.4) | 0.57 | 19.1 | 0.5 |
| Inception-V3  CNN | 0.69 | 48.1  (26.1) | 1.3  (0.7) | 0.43 | 2.78  (40.3) | 0.2  (2.8) | 0.58 | 18.7 | 0.2 |

In order to determine whether biases were consistent across plots, predicted versus observed values were plotted for each model type/forest attribute combination in Figure 2. In the estimating biomass, the LMM-THM appeared to slightly over predict in the highest biomass plots, while the CNN appeared to slightly under predict in the highest biomass plots (Figure 2 A, C). It should be noted however, that few plots fell within these extremes, and that loess regression is susceptible to outliers at the tail ends of data.

In predicting tree count, the LMM-THM and RF-THM both appeared to underestimate the number of trees in plots with high tree counts (Figure 2D, E). This trend is appears to be less substantial in the CNN model, despite having more slightly more overall bias than both the THM based models. In predicting percent needleleaf, the LMM-THM appeared to over predict in plots with more needleleaf stems (Figure 2G). Error was also greatest for this model around plots with a mix of species. The RF-THM and the CNN models both tended to slightly over predict percent needleleaf in plots with few needleleaf trees, and under predict in plots with a higher percentage of needleleaf trees. Overall however, there appeared to be no obvious trend in observed vs. fitted biases with model type.

** Figure 2.** Predicted vs. observed plots for each of the three model types estimating biomass, tree count, and percent needleleaf. The solid line is a one-to-one line, the dashed red line is a loess regression fit of the data. Left: linear mixed models with traditional height metrics (LMM-THM). Center: random forest models with traditional height metrics (RF-THM). Right: Inception-V3 convolutional neural networks (Inception-V3 CNN).

4. Discussion

Our results indicate that 3D CNNs can be used to develop an area-based forest inventory with less error and often less bias than a model based upon traditional height metrics. Model performance varied based on the specifics of the CNN architecture: those CNNs which made use of inception layers (GoogleNet and Inception-V3) outperformed those which did not. We also note that the deeper inception-based CNN, Inception-V3, outperformed the shallower inception-based GoogleNet. The CNN employing residual layers (ResNet-50) performed relatively poorly. We also evaluated ResNet-35 and ResNet 101 models, and obtained poorer and similar results, respectively. However, the top performing CNNs for image recognition presently make use of layers which combine residual and inception elements [45]; it is possible that such a model may outperform those tested here.

In general the RF-THM models outperformed the LMM-THM models. Random forest models resulted in lower RMSE and greater explained variance in terms of pseudo-R2. The linear mixed models often had slightly absolute lower biased than the random forest models, but this benefit was at times negated by greater bias in plots representing extremes. These finding match those of others who have likewise concluded that random forest performs equal to or better than linear modelling for LiDAR inventories [8, 60].

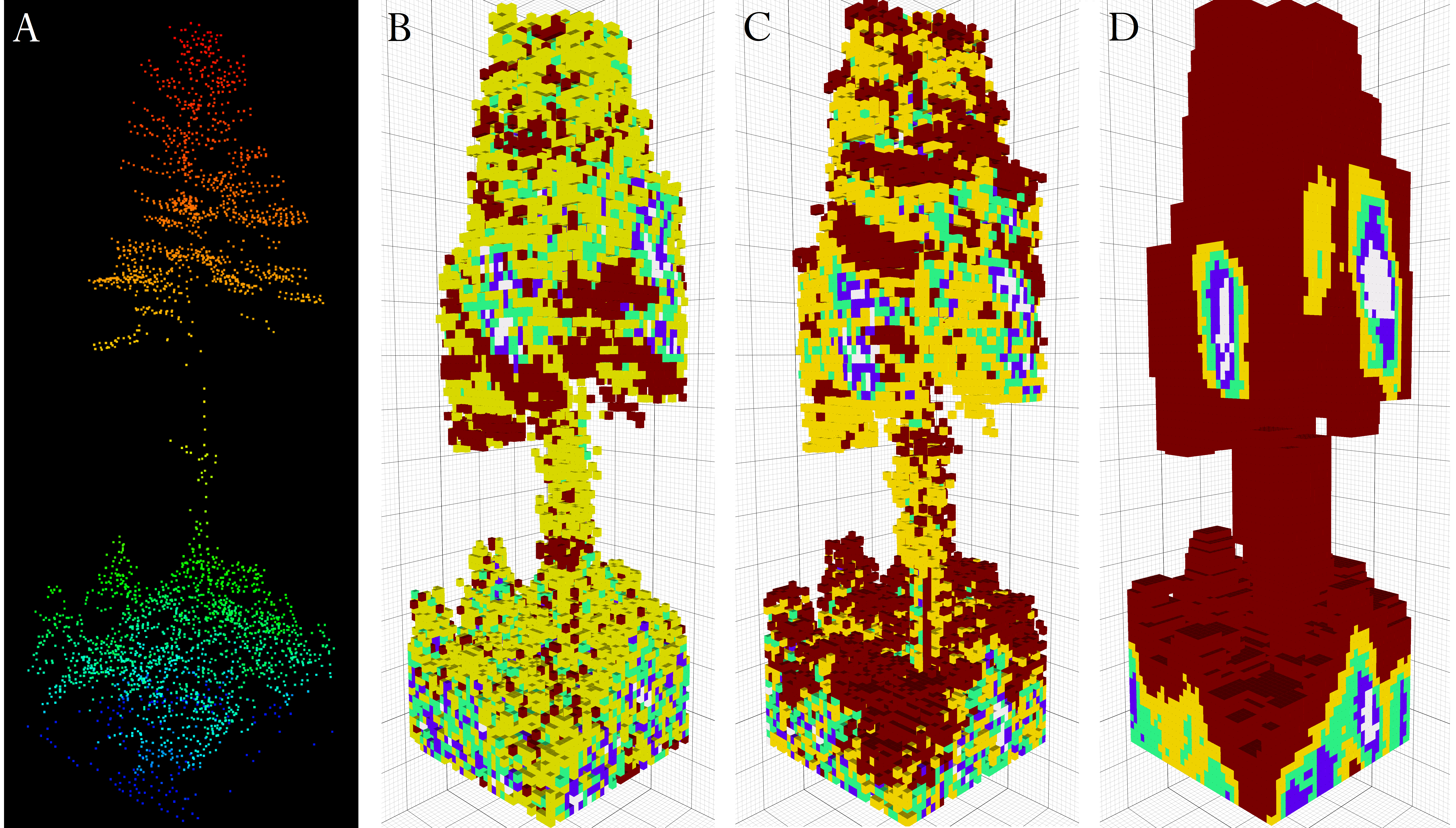
Performance of the CNNs relative to the THM based models varied by the forest attribute being predicted, though in every instance the CNNs produced a lower RMSE. In modelling biomass and percent needleleaf, the CNNs performed equal to or better than THM based models in terms of bias. The tree count CNN resulted in slightly more bias than both the THM models, however in practical terms this bias was minimal, and both the THM based models appeared to underestimate tree count to a greater extent in plots with greater numbers of trees.

We note that the greatest performance gains of using CNNs over the THM based models in terms of RMSE and explained variance were achieved when estimating biomass. However, it should also be noted that we also spent a greater amount of time and effort modelling biomass, as it was the attribute used to decide upon a model architecture. Neural networks offer modelers a great number of user-specified hyperparameters and architecture decisions. We believe that model performance was at least somewhat related to the amount of time spent manually fine-tuning these during training. This usually amounts to more-or-less a game of trial and error, made harder given the lengthy time it takes to train these models. Some studies have automated this process through random searches or more sophisticated means of optimization [62].

Many in the literature have noted that CNNs require a very large quantity of training data to achieve optimal performance [63, 28]. This will likely relegate the use of CNNs to only those instances in which multiple forest inventories are combined, as was the case here, or when a large national forest inventory dataset is used. We did however demonstrate here that the number of plots used to train these models can be artificially inflated through the collection of coincident remote sensing data. This mirrors results found by other studies [64, 65], in which dataset size was increased via transformations and the inclusion of many images rapidly taken of the same object. Several studies have also achieved good results by generating artificial training data [66, 67], and we believe that it may be possible to do the same with LiDAR. For example, artificial plots could be generated by combining pieces of other plots, individual tree crowns, or crowns derived from allometry or forest modeling [68].

In terms of computational performance, we made no formal effort to assess the time it took to train each model included in this study. We did note, however, that training the more complex models (such as Inception V3 and Resnet-50), could take one or more days using GPU acceleration with an Nvidia Tesla k80. As settling upon the optimal model architecture and hyperparameters will likely take many attempts, the modelling process may take days or weeks. A process called “warm starting” may offer future modelers a potential solution by allowing them to reuse some or all parameters of a pre-trained 3D CNN [69]. We would also note that once the models were trained, predictions could be made very rapidly at a landscape level, perhaps offsetting the initial cost in training time.

Critics of deep learning have justifiably noted that the results of a CNN may be more difficult to explain given that these models lack human intuited covariates. This, combined with the size of the models, makes it nearly impossible to trace the route of data through these models and justify the result. It is possible however to visualize features identified by early layers of the model to get an understanding of the shapes and patterns being used by the model to make predictions. These visualizations are often referred to as feature maps [70, 71]. The raw values of a feature map have little biological relevance, although the relative values score features according to their utility in model predictions. The higher relative values on a feature map highlights features that are more likely to be retained by the model following an activation function. Figure 3 illustrates this with example feature maps generated over a plot. We note that some convolution layers appear to be identifying edges (Figure 3D), while others appear to be identifying surfaces and possibly branches (Figure 3B, C).



**Figure 3.** (**A)** The initial point cloud prior to being input to the CNN model, colored by height. **(B-D)** Feature maps resulting from the first layer of convolutions using LeNet.Red values represent areas of higher interest to the model, blue and white areas of lower interest. Note that each convolution detects different patterns and features in the voxelized point cloud.

In addition to area-based forest inventories, we believe that CNNs may also be able to address the issue of individual tree segmentation. A wide array of algorithms have been put forward for segmenting the crowns of individual trees from a LiDAR point cloud for the purpose of developing a tree-list inventory [72-74]. Concurrently, CNNs have been enormously effective in segmenting objects from photographic and video imagery [75, 76]. Most CNN-based segmentation algorithms work by identifying potential bounding boxes of objects, and then analyzing the interior of those bounding boxes to assess their validity. We believe that a similar algorithm could be adapted to identify the 3D bounding boxes of individual trees. Another CNN-based segmentation method known as semantic segmentation seeks to isolate individual pixels (or voxels) that represent a desired object [77]. We believe that this method could be of use in classifying objects or terrain in LiDAR point clouds.

Another intriguing potential use for CNNs is in the development of pseudo-LiDAR point clouds. A number of studies have demonstrated the ability of CNNs to essentially be trained in reverse, producing images from data [78, 79]. These types of CNNs are known as deconvolutional, inverse graphic, or transposed neural networks. For example, a CNN could be trained to produce voxelized point clouds over a forested area from either previous LiDAR, or a standard forest inventory. These could then be used to inform modelers as to which features their neural networks are making use of, to test ecological hypotheses, to aid in visualization, and perhaps even to project LiDAR point clouds forward in time.

We have demonstrated that deep-learning methods using CNNs to interpret LiDAR data sets can improve upon traditional methods for area-based predictions of forest attributes. However, these improvements come with some drawbacks. Large amounts of training data, time, and effort are required for any modeling application that uses deep learning. Ultimately it falls upon modelers to use their own best judgment to decide whether improvements in model performance are worth the effort involved in successfully training these models. That said, given our success and the prevalent adoption of deep learning in related fields, it is safe to assume that deep learning will play a large role in remote sensing in the future.

5. Conclusions

We demonstrate here that deep learning via 3D convolutional neural networks can be trained to interpret LiDAR point clouds for the estimation of forest attributes such as biomass, tree count, and the percent of needleleaf trees. Our results indicate that CNNs of greater complexity and those which make use of inception layers are most effective at prediction. Of the models tested, the optimal model form was a 3D variant of Google’s Inception-V3. Given enough training data and fine tuning, these models frequently outperformed traditional feature-based approaches for modeling LiDAR.

**Supplementary Materials:** Additional material including relevant computer code and pre-trained models can be found online at https://github.com/Eayrey.

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**Author Contributions:**

Elias Ayrey designed this study, developed the methodology, performed the analysis, and wrote the manuscript. Daniel J. Hayes provided extensive revisions, aided in conceptualization, and supplied laboratory and computing resources.

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