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Article

Modeling Height—Diameter Relationship Using Artificial Neural Networks for Durango Pine Species in Mexico

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Abstract: The total tree height (h) and diameter at breast height (dbh) relationship is an essential tool in forest management and planning. The height—diameter (h - dbh) relationship had been studied with several approaches and for several species worldwide. The nonlinear mixed effect modeling (NLMEM) has been extensively used and lately the resilient backpropagation artificial neural network (RBPANN) approach has been a trend topic for modeling this relationship. The artificial neural network (ANN) is a computing system based in artificial intelligence and inspired in biological neural network for supervised learning. In this study the NLMEM and RBPANN approaches were used for modeling the h - dbh relationship for Durango pine species (*Pinus durangensis* Martínez) in mixed-species forest from Mexico. The total dataset considered 1,000 (11,472 measured trees) randomly selected from 14,390 temporary forest inventory plots and the dataset was randomly divided into two parts; 50% for training and 50% for testing. An unsupervised clustering analysis was used to grouped the dataset into 10 clusters based on k-means clustering method and plot-variables like density, basal area, mean dbh , mean h , quadratic mean diameter, altitude and aspect. The RBPANN was performed for tangent hyperbolicus (RBPANN-tanh), softplus (RBPANN-softplus), and logistic (RBPANN-logistic) activation functions for functions in cross product of the covariate or neurons and the weights for the ANN analysis. For both training and testing, 10 classical statistics (e.g., RMSE, R^2 , AIC, BIC, logLik) were computed for the residual values and assess the approaches for h - dbh relationship. For training and testing, the ANNs approach outperformed the NLMEM approach, and the RBPANN-tanh has the best performance in both training and testing phases.

Keywords: artificial intelligence; artificial neural network; height-diameter relationship; nonlinear mixed effect modeling

1. Introduction

Artificial Intelligence (AI) refers to the simulation of human intelligence in machines, wherein they are programmed to think and learn in a manner similar to humans. The AI has been used in forest modelling for different purposes and objectives. The machine learning (ML) is a subset of AI that focuses on the development of algorithms and models that enable computer to learns from a specific dataset and make predictions or take actions without being explicitly programmed [1,2]. One of the most ML techniques is the artificial neural network (ANN) and the resilient backpropagation artificial neural network (RBPANN) perform supervised ML in multi-layer perceptron, and the main principle is to eliminate the harmful influence of the size of the partial derivative on the weight step [3–5]. The ANNs are computational models inspired in the natural neurons and they represent a

generalization of mathematical models of human cognition or neural biology [1,6,7]. In ANNs, the training and testing datasets are used to train and evaluate the performance of the network for a specific randomly selected dataset. The training dataset is used to train the neural network. It consists of a set of input data points and their corresponding target output values, while, the testing dataset is a separate dataset that is used to evaluate the performance of the trained neural network [1,8,9].

One of the most important relationships in forest modeling is the total tree height and diameter at breast height (*h-dbh*) relationship, and this relationship is usually applied in forest inventory or for height estimation in forest management and planning [10]. The knowledge of *h-dbh* relationship is a fundamental both developing and applying many growth and yield models [11,12]. This relationship has mainly studied with nonlinear mixed effect modeling (NLMEM) with fix and random parameters for several species and grouping level or ecological conditions [10,11,13-16]. Lately, this relationship has been studied with AI, and the ML thought ANN has been used [7,17,18]. Also, other variables as basal area [19], crown width [20], biomass [21], volume [22], forest fire [23], and annual radial growth with competition indices [24] have been studied with different ML algorithms. Occasionally, the clustering analysis based in unsupervised ML has been included in to group similar data point together based on their inherent characteristics or similarities [1,25-27]. The unsupervised clustering analysis could identify patterns or structures in dataset to improve the fitted models in forest modeling.

Specifically, in the Mexican Forestry the *h-dbh* relationship has been extensively studied with NLMEM for local and generalized models and occasionally the unsupervised cluster analysis was included in modelling [12,28,29]. The NLMEM are better than fitted models by ordinary least squares method and those use random parameter to explain the variability between groups, sites, or ecological regions. Lately, the ML algorithms are taken attention in forestry research and the results outperform the NLMEM approach for the *h-dbh* relationship. In ANN analysis is convenient to separate the dataset in two parts, one for training and other one for testing or validation [7,17]. The main used model for NLMEM has been the Chapman-Richards model [30], which is base in a sigmoid relationship growth based on age [31].

Considering the above schemes and the context of AI in forestry research, this study aim the *h-dbh* relationship for Durango Pine species (*Pinus durangensis* Martínez) by NLMEM and ANN for unsupervised clustered dataset for training and testing sets. The algorithms were compared in both training and testing phases and some conventional statistics like root mean square error, coefficient of determination, Akaike's information criterion, Bayesian information criterion, and loglikelihood were uses to perform the approaches. The resilience backpropagation of ANN (RBPANN) was employed, and three activation functions were computed and evaluated. The activation functions were tangent hyperbolicus (RBPANN-tanh), softplus (RBPANN-softplus), and logistic (RBPANN-logistic), and those were trained by resilience backpropagation and maximum likelihood was used.

2. Materials and Methods

2.1. Study area

The study was developed in a forest community in Norther Mexico, specifically in Durango state. The forest community is called San Diego de Tezains, and the total area is around 61,098 ha, which 30,000 ha are used for forest management and timber harvesting. The main applied silvicultural treatments are based on continuous cover forestry (CCF) and rotation forest management (RFM) [32]. The silvicultural treatments for CCF area base on selection, while three thinning and shelterwood cutting treatments for RFM with 15-years of forest cycle cutting [33]. The location of study area is showed in Figure 1. The mean annual temperature ranges from 5 to 18 °C, and the lowest temperature occurs in January (− 6 °C) and the hottest in May (28 °C). The altitude varies from 1,500 to 3,033 m. The mixed-species stands are represented by seven genera: *Pinus*, *Quercus*, *Juniperus*, *Cupressus*, *Pseudotsuga*, *Arbutus*, and *Alnus*. The main species are *Pinus durangensis* Martínez and *Quercus sideroxyla* Bonpl. Lately, the Improve forest management combine the forest

management and credit carbon offsets according to the Mexican Protocol developed by The Climate Action Reserve [34].

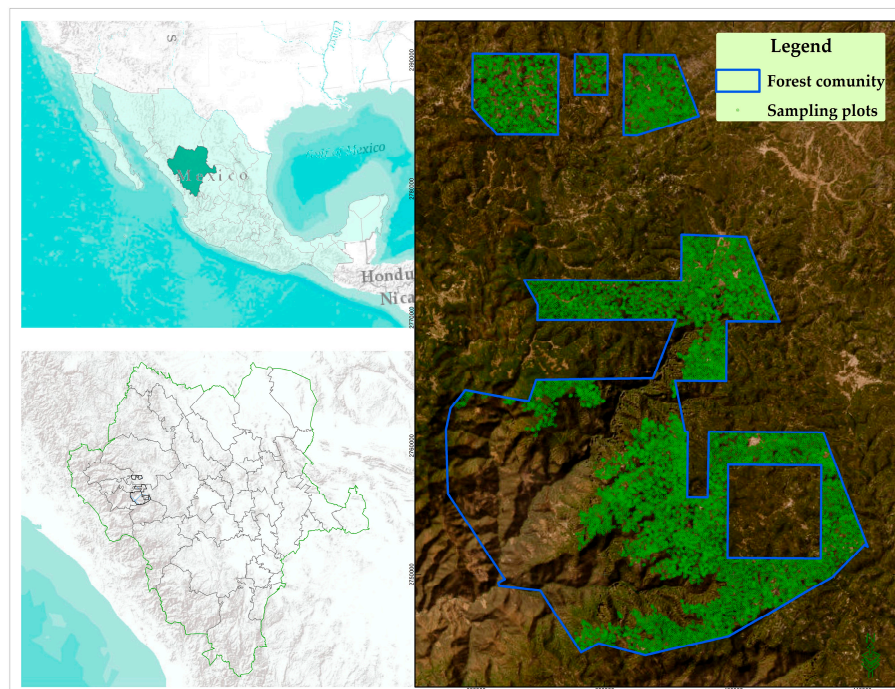


Figure 1. Study area location in Northern Mexico.

2.2. Dataset description

The dataset came from temporary forest inventory plots with a random sampling design for framework of 30,000 ha. A total of 14,390 temporary forest inventory plots were considered and Durango pine species was selected. A random sample of 1,000 plots was selected in *sampling* R package [35], and 11,472 measured trees were considered. Firstly, the unsupervised clustering analysis was used for grouping the dataset [1,25] according to the k-Means Clustering of *kmeans* R package [35]. Ten clusters were generated according with density (N, trees per hectare), basal area (BA, m²), mean diameter (Dm, cm), mean total tree (Hm, m), quadratic mean diameter (QMD, cm), altitudes (A, m), slope (S, %), and aspect (As, categorical variable). All variables were standardized, which perform values bounded 0.0 and 1.0 [21,36,37]. The standardization was performed according to Milligan and Cooper [38] and the Equation 1.

$$Z = (x - \text{Min}(x)) / (\text{Max}(x) - \text{Min}(x)) \quad (1)$$

where Z is the standardized variable, x is the variable, Min and Max are the minimum and maximum values of x.

In the Clustering analysis, the cluster (k = 10) were generated, and the explained variance was 61.8812%. The data set is shown in Figure 2 for the total dataset and the clusters.

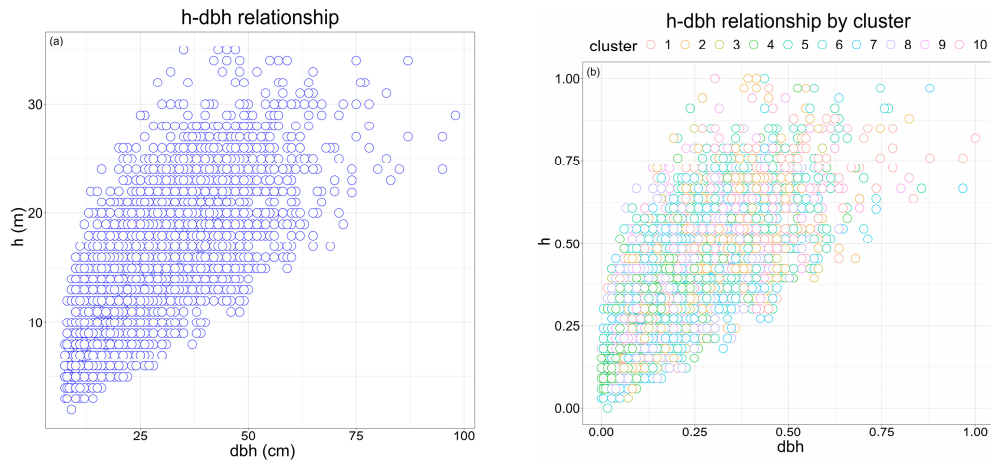


Figure 2. Scatter plot for *h-dbh* relationship for full dataset (a) and grouping dataset by cluster (b).

The site-specific variables for clustering analysis are recorded as a descriptive statistic in [Table 1](#). All variables were standardized with [Equation 1](#) to improve the clustering analysis [\[38\]](#).

Table 1. Descriptive statistics for plot-specific variables used in clustering analysis.

Variable	n	Statistic			
		Minimum	Mean	Maximum	SD
N	1000	1.0000	11.4720	57.0000	8.7717
BA	1000	0.0007	0.0193	0.0924	0.0137
Dm	1000	8.5000	22.9636	75.0000	7.6788
Hm	1000	4.0000	12.8062	35.0000	3.9813
QMD	1000	8.5147	24.7478	75.0000	8.0708
A	1000	2032.0000	2588.2170	2978.0000	137.3215
S	1000	0.0000	43.0499	96.0000	20.0551
As		1	5	9	2

N = density (trees per hectare); BA = basal area (m²); Dm = mean diameter (cm); Hm = mean total tree (m); QMD = quadratic mean diameter (cm); A = altitudes (m); S = slope (%); and As = aspect (categorical variable; 1 = Plain, 2 = N, 3 = S, 4 = E, 5 = W, 6 = NE, 7 = SE, 8 = NW, 9 = SW); n = observations; SD = standard deviation.

The total dataset (11,472 pairs of *h-dbh* relationship) was randomly divided in two sets; 50% for training and 50% for testing or validation. The main statistics for both training and testing dataset are shown in [Table 2](#) for the total tree height and diameter at breast height (*h-dbh*).

Table 2. This is a table. Tables should be placed in the main text near to the first time they are cited.

Dataset	Variable	n	Statistic			
			Minimum	Mean	Maximum	SD
Training	<i>h</i>	5736	7.5000	21.5362	95.0000	11.4394
	<i>dbh</i>	5736	3.0000	12.3900	35.0000	5.3217
Testing	<i>h</i>	5736	7.5000	21.3846	98.0000	11.5267
	<i>dbh</i>	5736	2.0000	12.1742	35.0000	5.2871

h = total tree height (m); *dbh* = diameter at breast height (cm); n = observations; SD = standard deviation.

2.3. Nonlinear mixed effect modeling (NLMEM)

The base growth model developed by [Richards \[30\]](#) was used to model the nonlinear *h-dbh* relationship. This model is based on a sigmoid curve, and it is represented by [Equation 2](#). This model had been extensively used in this kind of relationship [\[7,11,17,28\]](#).

$$h_{ij} = A_0 + \alpha_0(1 - e^{-\alpha_1 dbh_{ij}})^{\alpha_2} + e_{ij} \quad (2)$$

where h_{ij} = total tree height j in the plot i ; A_0 = lower asymptote parameter; α_0 , α_1 and α_2 = upper asymptote, growth rate, and slope of growth parameters; e = exponential function; dbh_{ij} = diameter at breast height j in the plot i ; e_{ij} = residual j in the plot i . In this case for A_0 parameter, the value 1.3 was fixed. This warranted the total tree height equal to 1.3 m when diameter at breast height is equal to 0 mentioned by [Fang and Bailey \[39\]](#).

For NLMEM, the parameter vector of nonlinear model was defined according to [Pinheiro, et al. \[40\]](#) and summarize as follows [\[7,11,13\]](#) (Equation 3):

$$\Phi_k = A_k \lambda + B_k b_k \quad (3)$$

where λ is the $p \times 1$ vector of fixed parameters (p is the number of fixed parameters in the model), b_k is the $q \times 1$ vector of random effect associated with lth cluster (q is the number of random parameters in the model), and A_k and B_k are the design matrices of size $r \times p$ and $r \times q$ (r is the total number of parameters in the model) for the fixed and random effects specific to each cluster. The residual vector (e_{ij}), and the random effect vector (b_k) are frequently assumed uncorrelated and normally distributed with mean zero and variance-covariance matrices R_k and D , respectively.

The upper asymptote parameter (α_o) was treated as a random parameter in the analysis for each cluster (α_{ol} $l = 1, 2, \dots, 10$), which explain about the maximum relationship between h and dbh . The random effect vector represent the variability between clusters for the asymptote parameter.

2.4. Artificial neural network (ANN)

The ANN are inspired by the early models of sensory processing by the brain. An ANN can easy be created by simulating a network of model neurons in a computer or specific programming language. Also, by applying mathematical algorithms that mimic the process of real neurons, we can make the network “learn” to solve many types of problems [\[41\]](#). The ANNs can learn by themselves. Because of they have similarities with the information processing features of the human brain (nonlinearity, high parallelism, capability to generalize), this modeling technique has the potential ability to solve problems that are difficult to formalize, such as problems of biological nature [\[7,41\]](#). The resilient backpropagation artificial neural network (RBPANN) is a logical adaptative learning scheme, which perform supervised batch learning in multi-layer perceptron. The basic principle of RBPANN is to eliminate the harmful influence of the size of the partial derivative on the weight steps [\[3,4,19\]](#).

According to [Barbosa, Costa, Schons, Finger, Liesenberg and Bispo \[19\]](#), and [Anastasiadis, et al. \[42\]](#), the RBP for ANN employs a sign-based scheme to update the weights in order to eliminate harmful influences of the derivatives’ magnitude on the weight updates. The size of the update step along a weight direction is exclusively determined by a weight-specific “update-value” as follows:

$$\Delta w_{ij}^k = \begin{cases} -\Delta_{ij}^k \text{ if } \frac{\partial E(w^k)}{\partial w_{ij}} > 0 \\ +\Delta_{ij}^k \text{ if } \frac{\partial E(w^k)}{\partial w_{ij}} < 0 \\ 0 \text{ otherwise} \end{cases} \quad (4)$$

where $\partial E(w^k)$ denotes the partial derivative of bathc error with respect to weight w_{ij} at the kth iteration.

The second step of RBP learning is to determine the new update values [\[19\]](#), as follows:

$$\Delta_{ij}^k = \begin{cases} \eta^+ \Delta_{ij}^k \text{ if } \frac{\partial E(w^{k-1})}{\partial w_{ij}} \frac{\partial E(w^{k-1})}{\partial w_{ij}} > 0 \\ \eta^- \Delta_{ij}^k \text{ if } \frac{\partial E(w^{k-1})}{\partial w_{ij}} \frac{\partial E(w^{k-1})}{\partial w_{ij}} < 0 \\ \Delta_{ij}^{k-1} \text{ otherwise} \end{cases} \quad (5)$$

where $0 < \eta^- < 1 < \eta^+$.

The total number of parameters in *RBPANN* is five; (i) the increase factor is set to $\eta^+ = 1.2$; (ii) the decrease factor is set to $\eta^- = 0.5$; (iii) the initial update-value is set to $\Delta_o = 0.1$; (iv) the maximum step; (v) the minimum steps.

According to [Cartwright \[43\]](#), the first step in using *ANNs* is to determine a suitable topology, optimal if possible (number of inputs and outputs, number of hidden layers, and neurons in each layer) and optimal (weights, biases, and activation functions). The process of *ANNs* begins by setting up the weights as small random variables. Then, each input pattern undergoes a feedforward phase, where the input signal is received and transmitted to all nodes in the hidden layer. In *ANNs*, every hidden node calculates the sum of its weighted input signals, applies an activation function to determine its output signal, and transmits this signal to the output node. At the output node, the final output signal is computed using the received signals from the hidden nodes [\[7\]](#). Within the context of *RBPANN*, the associated error (δ_k) is computed, and this error is utilized to adjust the weights. The weights correction term is determined based on the error, and it is subsequently employed to update the corresponding weights. Additionally, the δ_k is transmitted to each hidden node. Each hidden node then calculates its error information term by summing the inputs received from the output node, multiplied by the derivative of its activation function [\[7,43,44\]](#). According to [Fausett \[45\]](#) and [Cartwright \[43\]](#), the general formulation for *RBPANN* could be as follows ([Equation 6](#)):

$$\Delta w_{ij}(t+1) = \alpha \delta_k Z_j + \mu \Delta w_{ij} \quad (6)$$

where w_{ij} is the bias on output unit k , α is the learning rate, δ_k is the ratio of error correction weight fitted for w_{ij} that is due to an error at output O_k , also the information about the error at unit O_k that is propagated back to the hidden units that feed into unit O_k , Z_j is the output activation of the hidden unit j and μ is the momentum parameter (refers to the contribution of the gradient calculated at the previous time step to the correction of the weights) [\[43\]](#).

The used activation functions for smoothing the *h-dbh* relationship through *RBPANN* were tangent hyperbolicus (*RBPANN-tanh*), softplus (*RBPANN-softplus*), and logistic (*RBPANN-logistic*) functions [\[4,42\]](#), these activation functions occur between the hidden layers or between the input layer and hidden layer [\[17\]](#). These functions were defined for *RBPANN-tanh*, *RBPANN-softplus*, and *RBPANN-logistic* in [Equations 7, 8, and 9](#), respectively. Also, the derivatives are in [Equations 10, 11 and 12](#), respectively.

$$f(x) = \frac{e^x - e^{-x}}{e^x + e^{-x}} \quad (7)$$

$$f(x) = \frac{1}{1 + e^{-x}} \quad (8)$$

$$f(x) = \log(1 + e^x) \quad (9)$$

$$f'(x) = 1 - \left(\frac{e^x - e^{-x}}{e^x + e^{-x}} \right)^2 \quad (10)$$

$$f'(x) = \frac{e^{-x}}{(1 + e^{-x})^2} \quad (11)$$

$$f'(x) = \frac{e^x}{1 + e^x} \quad (12)$$

where $s = \sum w_i x_i$ is the information of the node transmits, in which w_i are the weights and x_i are the input values with $s \in [-1, 1]$, $s \in [0, \infty]$, and $s \in [0, 1]$ for *RBPANN-tanh*, *RBPANN-softplus*, and *RBPANN-logistic*, respectively.

In the *ANN* learning process, a different vector from 1 to 10 for each hidden layer was performed in a preliminary analysis, and the best results were obtained when the vector was 10 nodes for each hidden layer $c(10, 10, 10)$. In [Figure 3](#), the *ANNs* plot are presented for vectors $c(3, 3, 3)$, $c(5, 5, 5)$, and $c(10, 10, 10)$ in the hidden layer.

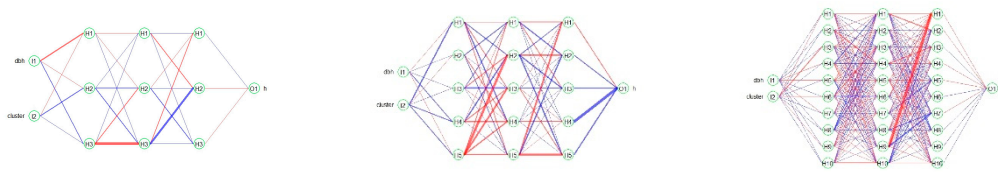


Figure 3. Plots of ANNs for vectors of $c(3, 3, 3)$ on the left, $c(5, 5, 5)$ on the center, and $c(10, 10, 10)$ on the right, in the hidden layer.

For both *RBPANN-tanh* and *RBPANN-logistic* functions, the topology was as follows: (i) two inputs (*dbh* and *cluster*); (ii) one output (*h*); (iii) a vector $c(10, 10, 10)$ hidden layers; and (iv) two nodes for the first layer, 11 nodes for the second, third and fourth layer layers (Bias node is included) and two nodes for the fifth layer. The ANNs for *RBPANN-tanh*, *RBPANN-softplus*, and *RBPANN-logistic* activation functions are presented in [Figure 4](#). Input variables are represented by "I" in nodes, hidden nodes are represented by "H", input variable is represented by "O", and Bias nodes are represented by "B".

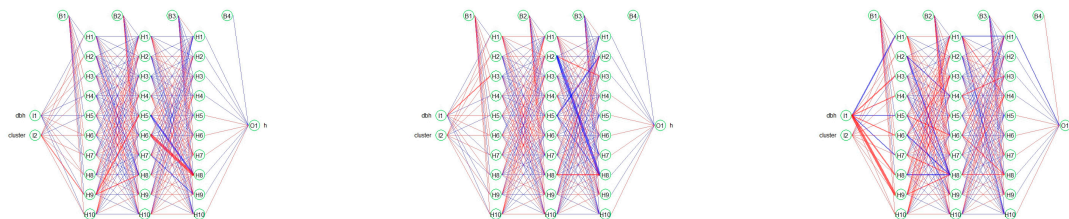


Figure 4. Plots of *RBPANN-tanh* (left), *RBPANN-softplus* (center), and *RBPANN-logistic* (right) for *h-dbh* relationship with unsupervised clustering analysis. Bias is included in nodes "B". Positive weight values in the visual representation are denoted by blue lines, while negative weight values are represented by red lines.

For *RBPANN-tanh*, *RBPANN-softplus*, and *RBPANN-logistic* activation functions. The number of repetitions was 10, the maximum steps for training of the NN was 10^7 , and the threshold was 0.1, which is similar that arguments used by [Özcelik, Diamantopoulou, Crecente-Campo and Eler \[7\]](#) and [Shen, Hu, Sharma, Wang, Meng, Wang, Wang and Fu \[17\]](#). Also, the training algorithm for ANNs was the resilient backpropagation with weight backtracking [\[4,42,43\]](#)

2.5. Fitting modeling

For *NLME*, the total tree height and diameter at breast height (*h-dbh*) relationship was fitted in "nlme" R package [\[35\]](#) and used maximum likelihood estimation method [\[40\]](#) for fixed and random parameters withing cluster groups. While for ANN models, the "neuralnet" R package [\[35\]](#) was used. For ANN, the resilient backpropagation (RPROP) for tangent hyperbolicus (*tanh*), softplus (*softplus*), and logistic (logistic) functions were programed for smoothing the result of the cross product of the covariate or neurons and the weights [\[3,4,45\]](#). All functions about fitting statistics for both training and testing were programmed in R environment [\[35\]](#).

2.6. Models performance criteria

For both training and testing steps, the fitting statistics were obtained in two levels, first one for the entire dataset and second one for each cluster. The statistics were the root mean square error (RMSE), standard error of estimate (SEE), relative SEE (RSEE), fitting index (FI), mean error (E),

relative E (RE), Akaike information criterion (AIC), Bayesian information criterion (BIC) and the log-likelihood ($\log Lik$). The statistics were computed as follows:

$$RMSE = \sqrt{\frac{\sum_{i=1}^n (O_i - P_i)^2}{n}} \quad (13)$$

$$SEE = \sqrt{\frac{\sum_{i=1}^n (O_i - P_i)^2}{n - p}} \quad (14)$$

$$RSEE = \frac{SEE}{\bar{O}} \quad (15)$$

$$FI = 1 - \frac{\sum_{i=1}^n (O_i - P_i)^2}{\sum_{i=1}^n (O_i - \bar{O})^2} \quad (16)$$

$$E = \frac{\sum_{i=1}^n (O_i - P_i)}{n} \quad (17)$$

$$RE = \frac{E}{\bar{O}} \quad (18)$$

$$AIC = n \log \left(\frac{\sum_{i=1}^n (O_i - P_i)^2}{n} \right) + 2p \quad (19)$$

$$BIC = n \log \left(\frac{\sum_{i=1}^n (O_i - P_i)^2}{n} \right) + p \log n \quad (20)$$

$$\log Lik = -n \log \left(\frac{\sum_{i=1}^n (O_i - P_i)^2}{n} \right) \quad (21)$$

where O_i , P_i and \bar{O} are observed, predicted and average values of h variable; n = observations; p = number of parameters estimated; and \log = logarithm function.

In all cases, the residual values were obtained with the implementation of *NLMEM* or *RBPANN* models and the statistics were programmed in R environment [35]. Lately, the *NLMEM* and *ANNs* models were ranked based on the overall dataset and cluster-group for all fitting statistics. A ranking system of [Kozak and Smith \[46\]](#) was used. All fitted statistics were equally weighted and Rank 1 was used for the best model and 4 for the poorest.

3. Results

3.1. Training phase

3.1.1. NLMEM

The fitted growth equation for h - dbh relationship by *NLMEM* performed well and all parameters were significantly different to zero at 5% of significance level. The relationship between total tree height and diameter at breast height can be explained with fixed and random parameters. In [Table 3](#), the estimated parameters and their statistical properties can be found for the entire training dataset. Furthermore, the confidence interval for each parameter is recorded at a 95% confidence level.

Table 3. Estimated parameter for h - dbh relationship in Durango pine by *NLMEM*.

Parameter	Estimate	SE	DF	t-value	p-value	lower	upper
α_0	26.409060	1.100113	5724	24.005770	<0.00001	24.252985	28.565134
α_1	0.029786	0.002534	5724	11.754320	<0.00001	0.024820	0.034752
α_2	1.083133	0.040518	5724	26.732200	<0.00001	1.003723	1.162543
$sd(\alpha_0)$	1.928939	0.583997	5724	3.302992	0.000962	1.210579	3.073574
σ	3.110839	0.029338	5724	106.033502	<0.00001	3.054379	3.168342
$\alpha_{0,1}$	-3.371745	0.106547	407	-31.645570	<0.00001	-3.580578	-3.162913
$\alpha_{0,2}$	-2.840826	0.089770	320	-31.645570	<0.00001	-3.016775	-2.664877
$\alpha_{0,3}$	-0.601879	0.019019	631	-31.645570	<0.00001	-0.639157	-0.564601
$\alpha_{0,4}$	3.572580	0.112894	133	31.645570	<0.00001	3.351309	3.793851
$\alpha_{0,5}$	0.773802	0.024452	925	31.645570	<0.00001	0.725876	0.821729
$\alpha_{0,6}$	0.478565	0.015123	1109	31.645570	<0.00001	0.448925	0.508206

$\alpha_{0,7}$	0.945549	0.029879	364	31.645570	<0.00001	0.886985	1.004113
$\alpha_{0,8}$	-0.308794	0.009758	876	-31.645570	<0.00001	-0.327919	-0.289668
$\alpha_{0,9}$	0.012327	0.000390	654	31.645570	<0.00001	0.011564	0.013091
$\alpha_{0,10}$	1.340420	0.042357	317	31.645570	<0.00001	1.257400	1.423440

SE = asymptotic standard error; DF = freedom degrees; sd = standard deviation for random effect between cluster groups; σ = standard error within-cluster-group.

The training phase's fitting statistics are presented in [Table 4](#), which includes the overall training dataset as well as the cluster-groups individually. For both overall training dataset, and cluster-groups, the fitting statistics were accurate and showed the potential to offer the *NLMEM* approach for the *h-dbh* relationship. The RMSE value for the overall training dataset was 3.1085 m, the best value was 2.4735 for cluster-group 1 (C1) and the worst value for cluster-group 4 (C4). Additionally, the overall training dataset exhibited an *E* value of -0.0005. The highest value was observed in C3, indicating the poorer performance, while the lowest value was found in C4, indicating the best performance. In terms of AIC, the C4 demonstrated the best performance, whereas C6 exhibited relatively poorer performance.

Table 4. Fitting statistics for *h-dbh* relationship in Durango pine by *NLMEM*.

Dataset	n	RMSE	SEE	RSEE	FI	E	RE	AIC	BIC	logLik
All-dataset	5736	3.1085	3.1123	25.1193	0.6588	-0.0005	-0.0042	13039.75	13139.57	-13009.75
C1	631	2.4735	2.4889	6.4322	0.6182	-0.0324	-0.3161	8834.14	772.23	-736.18
C2	407	2.5289	2.5489	6.1485	0.5378	0.0050	0.0515	7113.33	627.39	-592.78
C3	925	2.9631	2.9749	8.4267	0.6525	-0.1111	-0.9526	16437.91	1408.51	-1369.83
C4	1109	3.8688	3.9442	2.9876	0.5610	0.3416	1.7378	4306.53	388.22	-358.88
C5	320	3.1341	3.1426	10.6971	0.6312	-0.0072	-0.0610	25348.12	2153.32	-2112.34
C6	654	2.8727	2.8792	10.3014	0.6134	0.0555	0.4528	28074.42	2381.60	-2339.53
C7	364	3.4345	3.4584	6.5911	0.6329	-0.0640	-0.4879	10767.05	932.64	-897.25
C8	876	3.3842	3.3939	10.2592	0.6015	0.0210	0.1632	25618.61	2175.54	-2134.88
C9	133	3.2098	3.2221	8.8088	0.6138	-0.0229	-0.1862	18292.63	1563.28	-1524.39
C10	317	3.6168	3.6458	5.1452	0.6196	-0.0058	-0.0353	9768.73	848.61	-814.06

RMSE = root mean square error (m); SEE = standard error of estimate (m); RSEE = relative SEE (%); FI = fitting index; E = mean error (m); RE = relative (%); AIC = Akaike information criterion; BIC = Bayesian information criterion; logLik = log-likelihood value; C = cluster-group.

3.1.1. RBPANN

The results about performed ANNs for *RBPANN*-tanh, *RBPANN*-softplus, and *RBPANN*-logistic activation functions are shown in [Table 5](#). The statistics are in standardized variables and observed in 10 repetitions in the learning process. In this scenario, all three activation functions exhibit favorable outcomes. Specifically, both *RBPANN*-tanh and *RBPANN*-softplus deliver comparable performance. In contrast, *RBPANN*-logistic exhibited the lowest performance among the three activation functions. Additionally, the *RBPANN*-logistic achieved the minimum number of steps (88) required for convergence.

Table 5. Main fitting statistics for *RBPANN*-tanh, *RBPANN*-softplus, and *RBPANN*-logistic activation functions tested for *h-dbh* relationship.

ANN	Error	Reached Threshold	Steps	AIC	BIC
<i>RBPANN</i> -tanh	27.8455	0.0775	301	577.69	2314.52
<i>RBPANN</i> -softplus	27.3939	0.0838	1885	576.79	2313.62
<i>RBPANN</i> -logistic	28.4113	0.0994	88	578.82	2315.65

AIC = Akaike information criterion; BIC = Bayesian information criterion.

The fitting statistics for three ANNs applied to examine the h - dbh relationship in Durango pine are presented for both the overall dataset and each cluster-group in [Table 6](#) for training phase. The nine fitting statistics illustrate the accuracy of *RBPANN*-tanh, *RBPANN*-softplus, and *RBPANN*-logistic activation functions in the ANNs models. The topology of each ANN, as depicted in [Figure 4](#), exhibited satisfactory results in predicting h based on dbh and an unsupervised clustering analysis for ten groups. Overall, the estimations demonstrate similar characteristics across the three activation models. However, *RBPANN*-tanh exhibits certain advantages that are comparable to the other activation functions in the training phase. All ANNs were trained using the resilience backpropagation learning algorithm, and the likelihood function was employed.

Table 6. Fitting statistics for h - dbh relationship in Durango pine by ANNs and different backpropagations activation functions.

Dataset	RMSE	SEE	RSEE	FI	E	RE	AIC	BIC	logLik
<i>RBPANN</i> -tanh									
All-dataset	2.8122	2.8134	22.7071	0.7208	0.0001	-0.0003	11872.59	11912.52	-11860.59
C1	2.6322	2.6427	8.0184	0.7258	0.0002	0.0018	14644.89	1259.09	-1220.41
C2	2.2127	2.2264	6.1634	0.6945	0.0073	0.0717	7745.69	681.53	-645.47
C3	2.8170	2.8246	10.2988	0.7020	-0.0087	-0.0741	22979.70	1955.95	-1914.98
C4	2.5795	2.5853	9.9082	0.6883	0.0636	0.5191	25209.16	2142.83	-2100.76
C5	2.4178	2.4370	6.2966	0.5776	-0.6268	-6.4583	6768.31	598.64	-564.03
C6	2.8907	2.9019	8.4977	0.6867	0.2559	2.0805	16649.44	1426.35	-1387.45
C7	3.1340	3.1558	6.4423	0.6943	0.3677	2.8037	9967.09	865.97	-830.59
C8	3.0652	3.0740	9.9535	0.6731	-0.3691	-2.8636	23537.45	2002.11	-1961.45
C9	3.7470	3.8200	3.0995	0.5882	1.5107	7.6864	4204.43	379.71	-350.37
C10	3.2490	3.2751	4.9509	0.6930	-0.1391	-0.8426	8952.96	780.63	-746.08
<i>RBPANN</i> -softplus									
All-dataset	2.8431	2.8443	22.9565	0.7143	0.7146	-0.0013	-0.0107	11997.88	12037.81
C1	2.6516	2.6621	8.0772	0.7218	0.0489	0.4190	14755.60	1268.32	-1229.63
C2	2.4591	2.4744	6.8498	0.6226	-0.9459	-9.2408	8777.19	767.49	-731.43
C3	2.8529	2.8606	10.4301	0.6944	0.4200	3.5697	23260.85	1979.38	-1938.40
C4	2.6003	2.6062	9.9882	0.6833	0.3109	2.5351	25423.15	2160.66	-2118.60
C5	2.4956	2.5154	6.4993	0.5499	-0.9140	-9.4178	7011.62	618.91	-584.30
C6	2.8841	2.8952	8.4781	0.6882	-0.1042	-0.8472	16613.23	1423.33	-1384.44
C7	3.1030	3.1246	6.3787	0.7003	0.1339	1.0213	9880.37	858.75	-823.36
C8	3.0748	3.0837	9.9847	0.6711	-0.3712	-2.8799	23603.31	2007.59	-1966.94
C9	3.8187	3.8931	3.1588	0.5723	1.6461	8.3754	4264.92	384.75	-355.41
C10	3.2550	3.2811	4.9600	0.6919	0.0992	0.6009	8966.90	781.80	-747.24
<i>RBPANN</i> -logistic									
All-dataset	2.8486	2.8498	23.0012	0.7135	-0.0052	-0.0420	12020.19	12060.12	-12008.19
C1	2.6570	2.6676	8.0938	0.7206	0.0785	0.6729	14786.65	1270.90	-1232.22
C2	2.4675	2.4828	6.8732	0.6201	-0.9254	-9.0403	8810.45	770.26	-734.20
C3	2.8586	2.8664	10.4510	0.6932	0.4146	3.5237	23305.34	1983.09	-1942.11
C4	2.6024	2.6083	9.9962	0.6827	0.2868	2.3386	25444.48	2162.44	-2120.37
C5	2.5183	2.5382	6.5583	0.5417	-0.9468	-9.7561	7081.01	624.69	-590.08
C6	2.9055	2.9167	8.5411	0.6835	-0.1415	-1.1503	16729.32	1433.01	-1394.11
C7	3.0957	3.1172	6.3636	0.7017	0.1066	0.8129	9859.77	857.03	-821.65
C8	3.0740	3.0828	9.9818	0.6712	-0.3751	-2.9095	23597.18	2007.08	-1966.43
C9	3.8192	3.8936	3.1592	0.5721	1.6756	8.5253	4265.34	384.79	-355.45
C10	3.2584	3.2845	4.9652	0.6912	0.1835	1.1115	8974.85	782.46	-747.90

RMSE = root mean square error (m); SEE = standard error of estimate (m); RSEE = relative SEE (%); FI = fitting index; E = mean error (m); RE = relative (%); AIC = Akaike information criterion; BIC = Bayesian information criterion; logLik = log-likelihood value; C = cluster-group.

The residual and predicted values are presented in [Figure 5](#) for each cluster-group by *RBPANN*-tanh, which was the best approach in training phase to model the *h-dbh* relationship. In general, the residuals ranged between -6 m and 6 m. Enhancing the training phase could involve increasing the number of repetitions or epochs, however, the computational process ought to be significantly time-consuming.

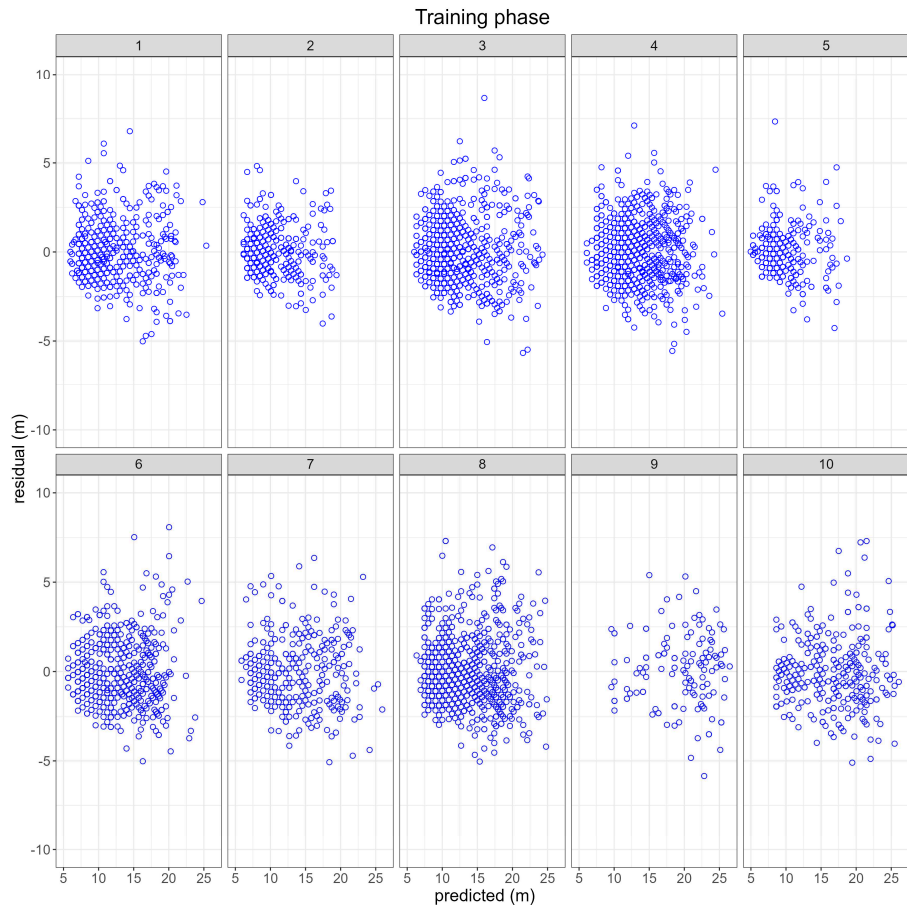


Figure 5. Residual versus predicted values for *RBPANN*-tanh model in training phase and each cluster-group.

Finally, The ranks and sums of ranks for hierarchy of *NLMEN* and *ANNs* are presented in [Table 7](#). The statistics for both the overall dataset and the cluster-group were ranked from 1 to 4. In terms of the overall dataset, the *RBPANN*-softplus exhibited the best performance during the training phase, which was comparable to the *NLMEN* model and other *ANNs*. The *RBPANN*-tanh activation function ranked second, while the *RBPANN*-logistic ranked third. On the other hand, the *NLMEN* approach had the lowest rank sum, indicating poorer performance compared to the other models. A similar pattern was observed within the cluster-groups, and the sum of ranks resulted in the following rankings: 176, 241, 269, and 304 for *RBPANN*-tanh, *RBPANN*-softplus, *NLMEN*, and *RBPANN*-logistic, respectively. It is worth noting that only the *RBPANN*-logistic demonstrated lower performance compared to the *NLMEN* approach. The number in parenthesis indicate the ranking for models in the overall dataset.

Table 7. Ranks and sum of ranks based on the fitting statistics for fitted models in training phase.

Model	Dataset	RMSE	SEE	RSEE	FI	E	RE	AIC	BIC	logLik	Rank
<i>NLMEN</i>	Overall	4	4	4	4	3	2	4	4	4	33 (4)
<i>RBPANN</i> -tanh	Overall	1	1	1	1	4	4	2	1	2	17 (2)
<i>RBPANN</i> -softplus	Overall	2	2	2	2	1	3	1	2	1	16 (1)

<i>RBPANN-logistic</i>	Overall	3	3	3	3	2	1	3	3	3	24 (3)
<i>NLMEM</i>	<i>C1</i>	1	1	1	4	2	2	1	1	1	14
<i>RBPANN-tanh</i>	<i>C1</i>	2	2	2	1	1	1	2	2	2	15
<i>RBPANN-softplus</i>	<i>C1</i>	3	3	3	2	3	3	3	3	3	26
<i>RBPANN-logistic</i>	<i>C1</i>	4	4	4	3	4	4	4	4	4	35
<i>NLMEM</i>	<i>C2</i>	4	4	1	4	1	1	1	1	1	18
<i>RBPANN-tanh</i>	<i>C2</i>	1	1	2	1	2	2	2	2	2	15
<i>RBPANN-softplus</i>	<i>C2</i>	2	2	3	2	4	4	3	3	3	26
<i>RBPANN-logistic</i>	<i>C2</i>	3	3	4	3	3	3	4	4	4	31
<i>NLMEM</i>	<i>C3</i>	4	4	1	4	2	2	1	1	1	20
<i>RBPANN-tanh</i>	<i>C3</i>	1	1	2	1	1	1	2	2	2	13
<i>RBPANN-softplus</i>	<i>C3</i>	2	2	3	2	4	4	3	3	3	26
<i>RBPANN-logistic</i>	<i>C3</i>	3	3	4	3	3	3	4	4	4	31
<i>NLMEM</i>	<i>C4</i>	4	4	1	4	4	2	1	1	1	22
<i>RBPANN-tanh</i>	<i>C4</i>	1	1	2	1	1	1	2	2	2	13
<i>RBPANN-softplus</i>	<i>C4</i>	2	2	3	2	3	4	3	3	3	25
<i>RBPANN-logistic</i>	<i>C4</i>	3	3	4	3	2	3	4	4	4	30
<i>NLMEM</i>	<i>C5</i>	4	4	4	1	1	1	4	4	4	27
<i>RBPANN-tanh</i>	<i>C5</i>	1	1	1	2	2	2	1	1	1	12
<i>RBPANN-softplus</i>	<i>C5</i>	2	2	2	3	3	3	2	2	2	21
<i>RBPANN-logistic</i>	<i>C5</i>	3	3	3	4	4	4	3	3	3	30
<i>NLMEM</i>	<i>C6</i>	1	1	4	4	1	1	4	4	4	24
<i>RBPANN-tanh</i>	<i>C6</i>	3	3	2	2	4	4	2	2	2	24
<i>RBPANN-softplus</i>	<i>C6</i>	2	2	1	1	2	2	1	1	1	13
<i>RBPANN-logistic</i>	<i>C6</i>	4	4	3	3	3	3	3	3	3	29
<i>NLMEM</i>	<i>C7</i>	4	4	4	4	1	1	4	4	4	30
<i>RBPANN-tanh</i>	<i>C7</i>	3	3	3	3	4	4	3	3	3	29
<i>RBPANN-softplus</i>	<i>C7</i>	2	2	2	2	3	3	2	2	2	20
<i>RBPANN-logistic</i>	<i>C7</i>	1	1	1	1	2	2	1	1	1	11
<i>NLMEM</i>	<i>C8</i>	4	4	4	4	1	1	4	4	4	30
<i>RBPANN-tanh</i>	<i>C8</i>	1	1	1	1	2	2	1	1	1	11
<i>RBPANN-softplus</i>	<i>C8</i>	3	3	3	3	3	3	3	3	3	27
<i>RBPANN-logistic</i>	<i>C8</i>	2	2	2	2	4	4	2	2	2	22
<i>NLMEM</i>	<i>C9</i>	1	1	4	1	1	1	4	4	4	21
<i>RBPANN-tanh</i>	<i>C9</i>	2	2	1	2	2	2	1	1	1	14
<i>RBPANN-softplus</i>	<i>C9</i>	3	3	2	3	3	3	2	2	2	23
<i>RBPANN-logistic</i>	<i>C9</i>	4	4	3	4	4	4	3	3	3	32
<i>NLMEM</i>	<i>C10</i>	4	4	4	4	1	1	4	4	4	30
<i>RBPANN-tanh</i>	<i>C10</i>	1	1	1	1	3	3	1	1	1	13
<i>RBPANN-softplus</i>	<i>C10</i>	2	2	2	2	2	2	2	2	2	18
<i>RBPANN-logistic</i>	<i>C10</i>	3	3	3	3	4	4	3	3	3	29

RMSE = root mean square error; *SEE* = standard error of estimate; *RSEE* = relative SEE; *FI* = fitting index; *E* = mean error; *RE* = relative *E*; *AIC* = Akaike information criterion; *BIC* = Bayesian information criterion; *logLik* = log-likelihood value; *C* = cluster-group.

3.1. Testing or Validation phase

3.1.1. NLMEM

During the testing phase, 5,736 pairs of heights and diameters from 50% of the dataset were utilized. Height estimations were performed using fixed and random parameters for each cluster-group provided by *NLMEM* approach. The nine testing statistics were computed at two levels: for the overall dataset and for each cluster-group. The [Table 8](#) presents the statistics for testing the

advantages of NLMEN for overall dataset and for each cluster-group. All the statistics displayed satisfactory performance, which depended on the number of observations. Additionally, the cluster-groups with limited information exhibited the lowest values. Among the cluster-groups, C4 had the maximum number of observations, whereas C9 had the minimum number of observations.

Table 8. Testing statistics for *h-dbh* relationship in Durango pine by *NLMEM* in testing phase.

Dataset	n	RMSE	SEE	RSEE	FI	E	RE	AIC	BIC	logLik
All-dataset	5736	3.1438	3.1476	25.8549	0.6464-0.1611	-1.3229	13169.29	13269.10	-13139.29	
C1	631	2.7037	2.7141	24.1931	0.6893-0.0987	-0.8795	15671.32	1344.87	-1305.94	
C2	407	2.9045	2.9228	29.6973	0.4680	1.2223	12.4194	10249.76	890.11	-854.15
C3	925	3.0941	3.1029	27.5443	0.6136-0.7206	-6.3964	23870.14	2029.86	-1989.18	
C4	1109	3.0509	3.0578	25.0690	0.6102-0.0942	-0.7725	29971.18	2539.72	-2497.60	
C5	320	2.7563	2.7797	28.6992	0.5175	0.9194	9.4926	7263.59	639.50	-605.30
C6	654	3.1718	3.1834	25.4080	0.6085	0.1428	1.1398	19047.71	1626.51	-1587.31
C7	364	3.4512	3.4752	26.2781	0.6545-0.7996	-6.0462	10839.28	938.67	-903.27	
C8	876	3.0615	3.0704	24.6303	0.6290-0.1456	-1.1680	23216.40	1975.28	-1934.70	
C9	133	5.0061	5.1138	25.4219	0.2128-2.4332	-12.0959	4665.31	417.55	-388.78	
C10	317	3.8668	3.8957	24.6024	0.5555-0.7956	-5.0245	10991.37	950.90	-915.95	

RMSE = root mean square error (m); SEE = standard error of estimate (m); RSEE = relative SEE (m); FI = fitting index; E = mean error (m); RE = relative E (%); AIC = Akaike information criterion; BIC = Bayesian information criterion; logLik = log-likelihood value; C = cluster-group.

3.1.1. RBPANN

During the testing phase, the results for the ANNs were similar to the training phase. The ANN utilizing the tangent hyperbolicus activation function (*RBPANN-tanh*) exhibited the highest performance, followed by *RBPANN-logistic*, and finally *RBPANN-softplus*. The [Table 9](#) records the statistics for the testing dataset, both in the overall dataset and within each cluster-group. The *FIs* values for the overall dataset were higher than 0.7029, when *RBPANN-tanh* demonstrating the best performance and *RBPANN-logistic* exhibiting the poorest performance. A similar pattern was observed for other statistics such as AIC BIC and logLik. Furthermore, in this instance, the ANNs demonstrate superior performance compared to the *NLMEM* approach.

Table 9. Testing statistics for both overall dataset and each cluster-group in testing phase with ANNs approaches.

Dataset	RMSE	SEE	RSEE	FI	E	RE	AIC	BIC	logLik
<i>RBPANN-tanh</i>									
All-dataset	2.8693	2.8706	23.5793	0.7055	0.6603	5.4241	12103.39	12143.32	-12091.39
C1	2.5090	2.5186	8.1057	0.7324	0.6015	5.3617	14492.43	1246.63	-1207.70
C2	2.4793	2.4949	7.1292	0.6124	0.8309	8.4421	8726.27	763.15	-727.19
C3	2.7294	2.7372	10.1704	0.6994	0.4046	3.5917	21218.12	1808.86	-1768.18
C4	2.8842	2.8907	11.1931	0.6516	0.8969	7.3531	28460.68	2413.85	-2371.72
C5	2.3880	2.4083	6.0227	0.6378	0.0825	0.8517	6234.38	553.73	-519.53
C6	3.0888	3.1001	9.1436	0.6287	1.2080	9.6415	18609.70	1590.01	-1550.81
C7	3.1665	3.1885	6.4643	0.7091	0.8613	6.5128	10085.03	875.82	-840.42
C8	2.7710	2.7790	9.2458	0.6961	0.1519	1.2183	21146.63	1802.80	-1762.22
C9	4.2322	4.3232	3.2613	0.4374	1.8156	9.0259	4177.60	376.91	-348.13
C10	3.4732	3.4992	5.7063	0.6414	0.5225	3.2997	10118.01	878.12	-843.17
<i>RBPANN-softplus</i>									
All-dataset	2.8764	2.8776	23.6371	0.7040	0.6578	5.4029	12131.48	12171.41	-12119.48
C1	2.5181	2.5278	8.1353	0.7305	0.6756	6.0219	14550.00	1251.43	-1212.50
C2	2.3388	2.3536	6.7253	0.6551	-0.0814	-0.8266	8164.99	716.38	-680.42

C3	2.8312	2.8393	10.5500	0.6765	0.8131	7.2176	21992.85	1873.42	-1832.74
C4	2.9657	2.9724	11.5095	0.6317	1.1148	9.1400	29209.97	2476.29	-2434.16
C5	2.4000	2.4204	6.0529	0.6342	-0.2679	-2.7660	6270.23	556.72	-522.52
C6	2.9772	2.9881	8.8135	0.6550	0.7866	6.2785	18002.41	1539.40	-1500.20
C7	3.1029	3.1244	6.3344	0.7207	0.6147	4.6485	9907.23	861.00	-825.60
C8	2.7748	2.7828	9.2584	0.6952	0.1450	1.1632	21174.89	1805.15	-1764.57
C9	4.3045	4.3971	3.3171	0.4180	2.0025	9.9547	4226.81	381.01	-352.23
C10	3.5266	3.5530	5.7940	0.6303	0.8555	5.4026	10242.05	888.46	-853.50
<i>RBPANN-logistic</i>									
All-dataset	2.8820	2.8832	23.6832	0.7029	0.6484	5.3263	12153.85	12193.77	-12141.85
C1	2.5071	2.5168	8.0998	0.7328	0.6480	5.7760	14481.07	1245.68	-1206.76
C2	2.3476	2.3624	6.7505	0.6525	-0.0800	-0.8123	8200.94	719.38	-683.41
C3	2.8287	2.8368	10.5405	0.6771	0.8418	7.4726	21973.90	1871.84	-1831.16
C4	2.9718	2.9785	11.5331	0.6302	1.1385	9.3342	29264.97	2480.87	-2438.75
C5	2.3833	2.4035	6.0108	0.6392	-0.2199	-2.2702	6220.20	552.55	-518.35
C6	2.9674	2.9783	8.7844	0.6573	0.8301	6.6255	17947.97	1534.87	-1495.66
C7	3.1024	3.1240	6.3335	0.7208	0.6342	4.7956	9905.96	860.90	-825.50
C8	2.7763	2.7843	9.2634	0.6949	0.1438	1.1539	21186.18	1806.09	-1765.51
C9	4.2538	4.3453	3.2780	0.4316	1.9452	9.6698	4192.42	378.14	-349.37
C10	3.4913	3.5174	5.7360	0.6377	0.7841	4.9519	10160.28	881.65	-846.69

RMSE = root mean square error (m); *SEE* = standard error of estimate (m); *RSEE* = relative *SEE* (%); *FI* = fitting index; *E* = mean error (m); *RE* = relative *E* (%); *AIC* = Akaike information criterion; *BIC* = Bayesian information criterion; *logLik* = log-likelihood value; *C* = cluster-group.

The [Figure 6](#) displays the representation of residual and predicted values for each cluster-group obtained through *RBPANN-tanh*. In this scenario, the residual dispersion appears to be larger compared to the training phase. However, this can be attributed to the utilization of a new dataset, where the predictions are made under different training conditions. Cluster-groups 4 and 10 exhibited higher levels of dispersion compared to the remaining cluster-groups.

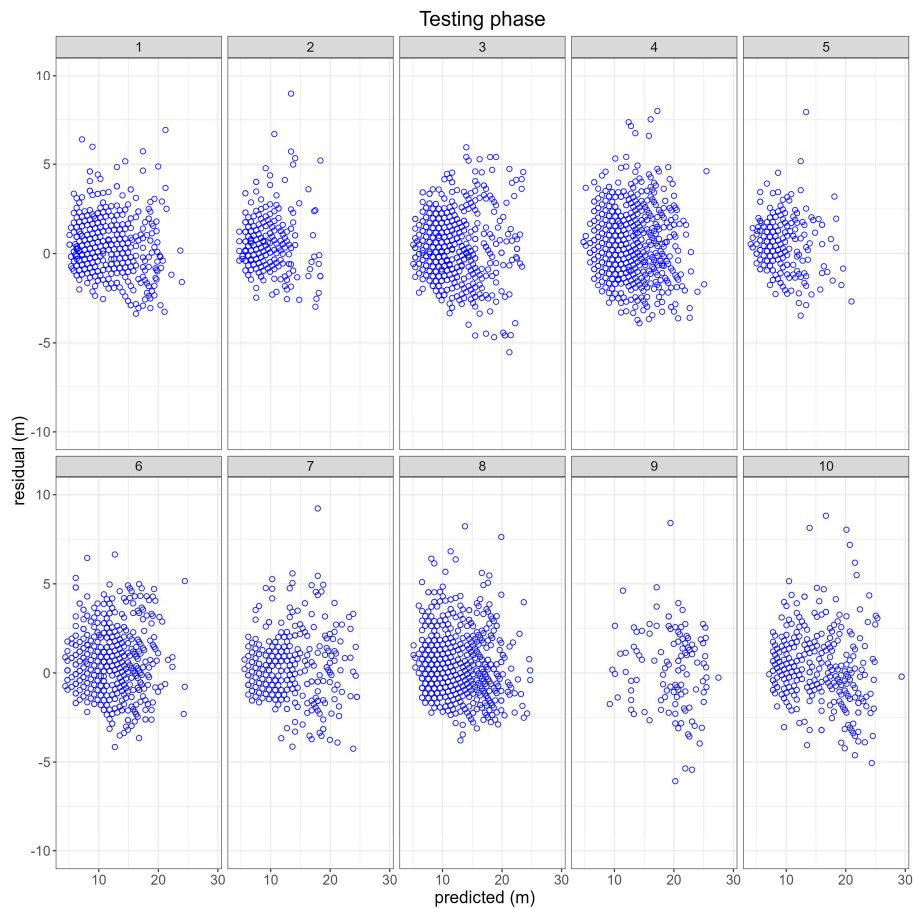


Figure 6. Residual versus predicted values for *RBPANN-tanh* model in testing phase for each cluster-group.

Finally, the ranks and sums of ranks for hierarchy of *NLMEN* and *ANNs* are presented in [Table 10](#). During the testing phase, the *RBPANN-tanh* achieved the highest rank of 1 with a sum of ranks of 9, while the *NLMEM* approach performed the poorest, ranking 4 with sum of ranks of 36. The *RBPANN-softplus* and *RBPANN-logistic* exhibited relatively similar performance conditions. Among the proven models of *ANNs*, the *RBPANN-logistic* exhibited the poorest performance. In terms of the sum of ranks for the combined overall dataset and cluster-groups, the *RBPANN-tanh* demonstrated the best performance with a sum of ranks of 181. Following that, the *RBPANN-logistic*, *RBPANN-softplus*, and *NLMEM* were ranked 2nd, 3rd, and 4th, with sum of ranks of 201, 240, and 368, respectively.

Table 10. Ranks and sum of ranks based on the fitting statistics for fitted models in testing phase.

Model	Dataset	RMSE	SEE	RSEE	FI	E	RE	AIC	BIC	logLik	Rank
<i>NLMEM</i>	Overall	4	4	4	4	4	4	4	4	4	36 (4)
<i>RBPANN-tanh</i>	Overall	1	1	1	1	1	1	1	1	1	9 (1)
<i>RBPANN-softplus</i>	Overall	2	2	2	2	2	2	2	2	2	18 (2)
<i>RBPANN-logistic</i>	Overall	3	3	3	3	3	3	3	3	3	27 (3)
<i>NLMEM</i>	<i>C1</i>	4	4	4	4	1	1	4	4	4	30
<i>RBPANN-tanh</i>	<i>C1</i>	2	2	2	2	2	2	2	2	2	18
<i>RBPANN-softplus</i>	<i>C1</i>	3	3	3	3	4	4	3	3	3	29
<i>RBPANN-logistic</i>	<i>C1</i>	1	1	1	1	3	3	1	1	1	13
<i>NLMEM</i>	<i>C2</i>	4	4	4	4	4	4	4	4	4	36
<i>RBPANN-tanh</i>	<i>C2</i>	3	3	3	3	3	3	3	3	3	27
<i>RBPANN-softplus</i>	<i>C2</i>	1	1	1	1	2	2	1	1	1	11

<i>RBPANN-logistic</i>	C2	2	2	2	2	1	1	2	2	2	16
<i>NLMEM</i>	C3	4	4	4	4	2	2	4	4	4	32
<i>RBPANN-tanh</i>	C3	1	1	1	1	1	1	1	1	1	9
<i>RBPANN-softplus</i>	C3	3	3	3	3	3	3	3	3	3	27
<i>RBPANN-logistic</i>	C3	2	2	2	2	4	4	2	2	2	22
<i>NLMEM</i>	C4	4	4	4	4	1	1	4	4	4	30
<i>RBPANN-tanh</i>	C4	1	1	1	1	2	2	1	1	1	11
<i>RBPANN-softplus</i>	C4	2	2	2	2	3	3	2	2	2	20
<i>RBPANN-logistic</i>	C4	3	3	3	3	4	4	3	3	3	29
<i>NLMEM</i>	C5	4	4	4	4	4	4	4	4	4	36
<i>RBPANN-tanh</i>	C5	2	2	2	2	1	1	2	2	2	16
<i>RBPANN-softplus</i>	C5	3	3	3	3	3	3	3	3	3	27
<i>RBPANN-logistic</i>	C5	1	1	1	1	2	2	1	1	1	11
<i>NLMEM</i>	C6	4	4	4	4	1	1	4	4	4	30
<i>RBPANN-tanh</i>	C6	3	3	3	3	4	4	3	3	3	29
<i>RBPANN-softplus</i>	C6	2	2	2	2	2	2	2	2	2	18
<i>RBPANN-logistic</i>	C6	1	1	1	1	3	3	1	1	1	13
<i>NLMEM</i>	C7	4	4	4	4	3	3	4	4	4	34
<i>RBPANN-tanh</i>	C7	3	3	3	3	4	4	3	3	3	29
<i>RBPANN-softplus</i>	C7	2	2	2	2	1	1	2	2	2	16
<i>RBPANN-logistic</i>	C7	1	1	1	1	2	2	1	1	1	11
<i>NLMEM</i>	C8	4	4	4	4	3	3	4	4	4	34
<i>RBPANN-tanh</i>	C8	1	1	1	1	4	4	1	1	1	15
<i>RBPANN-softplus</i>	C8	2	2	2	2	2	2	2	2	2	18
<i>RBPANN-logistic</i>	C8	3	3	3	3	1	1	3	3	3	23
<i>NLMEM</i>	C9	4	4	4	4	4	4	4	4	4	36
<i>RBPANN-tanh</i>	C9	1	1	1	1	1	1	1	1	1	9
<i>RBPANN-softplus</i>	C9	3	3	3	3	3	3	3	3	3	27
<i>RBPANN-logistic</i>	C9	2	2	2	2	2	2	2	2	2	18
<i>NLMEM</i>	C10	4	4	4	4	3	3	4	4	4	34
<i>RBPANN-tanh</i>	C10	1	1	1	1	1	1	1	1	1	9
<i>RBPANN-softplus</i>	C10	3	3	3	3	4	4	3	3	3	29
<i>RBPANN-logistic</i>	C10	3	3	3	3	4	4	3	3	3	29

RMSE = root mean square error (m); SEE = standard error of estimate; RSEE = relative SEE; FI = fitting index; E = mean error; RE = relative E; AIC = Akaike information criterion; BIC = Bayesian information criterion; *logLik* = log-likelihood value; C = cluster-group.

4. Discussion

Having knowledge about the total tree height and diameter at breast height is essential for both the development and application of many growth and yield models. Models focusing on the *h-dbh* relationship serve as valuable tools for accurately predicting tree height based on *dbh* measurements. Because of the *dbh* can be conducted quickly, easily, and accurately, but the measurement of total tree height is comparatively complex, time consuming, and expensive [11]. The *NLME* had been a capable approach to generate models in *h-dbh* relationship for different species and assumed fixed and random parameters for specific-groups or covariables to study the variability inter-and intra-plots, ecological regions or cluster-groups [10,16,39]. Also, these models have been studied for local and generalized formulations with *NLMEM* approach [12,13,16,28]. In this case of study the *NLMEM* performance was accurately strong to model the *h-dbh* relationship for Durango pine and the inclusion of unsupervised clustering analysis improve the estimated parameters and its statistics properties [36,47], which involve fixed parameter for the overall dataset in training phase and random parameter for each cluster-group, also parameter to give information about general variability and variability within cluster-group.

The *NLMEM* demonstrated outstanding performance during the training phase, with the fitting process converging quickly and effortlessly. Additionally, the maximum likelihood approach yielded favorable and suitable results particularly when expressing the asymptote parameter with mixed effects ([Table 3](#) and [Table 4](#)). All parameters in fitting process were significantly different to zero at 5% of significance level and the random parameters allow suitable estimations in training phase and those were used for cluster-groups in testing phase. The application of the *NLMEM* approach on the testing dataset resulted in successful outcomes that aligned with the expected results (as shown in [Table 8](#)), accompanied by the utilization of appropriate statistical measures. As an illustration, the root mean square error (RMSE) for the overall dataset during the testing phase was determined to be 3.1438 m, with an average value of 3.3773 m observed within the cluster-groups (refer to [Table 8](#)). By employing a mixed-effect model and incorporating cluster-group inclusion, the Chapman-Richards growth equation [[30](#)] (Equation 2) proves to be a highly effective model for predicting the height of Durango pine trees. Similar results have been conducted for several species an different conditions [[11,16,28](#)]. Even though the *NLMEM* method is accurate for height prediction based on diameter measurements, it is worth considering that *ANNs* could be a suitable alternative for modeling the *h-dbh* relationship under several dataset conditions and the incorporation of grouped strategies [[7,14,48](#)]. In recent times, there has been a growing application of AI and ML techniques in the fields of biology and forestry. These advanced approaches have proven valuable in addressing challenges that require substantial computational resources and unsupervised learning methods [[1,41](#)]. Several of these approaches have been employed in studying the height-diameter at breast height (*h-dbh*) relationship, leading to notable outcomes and reported successes for various species and under diverse forest management conditions, demonstrating their versatility and effectiveness [[7,14,15,17,48](#)]. In this context the *ANNs* model outperformed the *NLMEM* approach.

In this study, the *ANNs* were evaluated and compared with the traditional *NLMEM* method. The *ANNs* utilized the RBP learning algorithm along with three activation functions. In most cases, the *ANNs* employing *RBPANN-tanh*, *RBPANN*, and *RBPANN-logistic* ([Equations 7, 8, an 9](#), respectively) exhibited superior performance compared to the results obtained by *NLMEM*, both during the training and testing phases. The training statistics for three *ANNs*, as presented in [Table 6](#), exhibited enhanced fitting performance compared to the statistics obtained by *NLMEM* (see [Table 4](#)). This improvement was observed in both the overall dataset and cluster-group analyses. These findings provide evidence that the clustering analysis using the k-means algorithm effectively grouped the dataset utilized in this study [[36,47](#)]. The *RBPANN-tanh* model, employing a tangent hyperbolic activation function, demonstrated the highest performance in predicting height measurements during both the training and testing phases (it can see in [Table 6](#) and [9](#)). Furthermore, the ranks and sum of ranks, based on the ranking system proposed by [Kozak and Smith \[46\]](#), provided evidence of the advantages of the *ANNs* models over the *NLMEM* approach. Models such as *RBPANN-logistic* were reported by [Özcelik, Diamantopoulou, Crecente-Campo and Eler \[7\]](#) revealed that models such as *RBPANN-logistic* exhibited advantages over *NLMEM* models when predicting the growth of Crimean juniper in the southern and southwestern regions of Turkey. Conversely [Shen, Hu, Sharma, Wang, Meng, Wang, Wang and Fu \[17\]](#) developed *ANNs* models utilizing *RBPANN-logistic* and *RBPANN-tanh* transfer or activation functions for *Populus spp.* L. in China, where the *RBPANN-logistic* model outperformed both the *NLMEN* and the *RBPANN-tanh* model. In our case, the best *ANN* was the *RBPANN-tanh* and this outperformed other tested *ANNs* and *NLMEM* approach. Similar results have been reported the advantages of *ANNs* or deep learning algorithms over the ordinary least square and *NLMEM* models in both training and testing or validation phases [[14,15,48,49](#)]. In all cases, the implementation of *ANNs* exhibited significant advantages over traditional approaches when modeling the *h-dbh* relationship.

In this study, based on the implemented ranking system, the *RBPANN-tanh* model emerged as the top performer (residual and predicted values are showed in [Figures 5](#) and [6](#)). It achieved a sum of ranks of 176 for the training phase and 81 for the testing phase. These sums of ranks account for both the overall dataset and cluster-groups, as illustrated in [Tables 7](#) and [10](#), respectively. In terms of training, the *RBPANN-softplus* model ranked second, whereas during the testing phase, the

RBPANN-logistic model exhibited the second-best performance. On the other hand, the *RBPANN*-logistic model performed least effectively in the training phase, while the *NLMEM* model demonstrated comparatively lower performance during the testing phase. The *ANNs* developed in this study, as depicted in [Figure 4](#), were trained using the RBP algorithm. The *ANNs* were then evaluated using three different activation functions: *RBPANN*-tanh, *RBPANN*-softplus, and *RBPANN*-logistic. These models comprised a total of five layers, including three hidden layers. The training process involved ten repetitions to ensure robustness and accuracy. Even though the *RBPANN*-logistic converging in 88 steps, it exhibited relatively poorer performance compared to the *RBPANN*-tanh, which achieved better results within 301 steps. Interestingly, the *RBPANN*-logistic required a longer convergence time of 1885 steps, indicating its comparatively poorer performance in this aspect. As a result, the developed *ANNs* model showcased a high capability for predicting total tree height measurements. This highlights the potential application of AI in modeling the *h-dbh* relationship, not only for Durango pine but also for general forest modeling purposes or other variables [[6,19,21,50](#)]. The *ANNs* could be used to improve the estimations in forest inventory and forest management and planning in mixed-species forest in Durango, Mexico.

5. Conclusions

The nonlinear mixed effect modeling (*NLMEM*) and Artificial Neural Networks (*ANN*) with resilience backpropagation (RBP) were employed to model the height-diameter at breast height (*h-dbh*) relationship for Durango pine species. Unsupervised clustering analysis was conducted to enhance the capability of the trained and tested models.. Three activation functions, namely tangent hyperbolicus (*RBPANN*-tanh), softplus (*RBPANN*-softplus), and logistic (*RBPANN*-logistic), were utilized in *RBPANN*. Those activation functions were trained and tested on both overall dataset and each cluster group. In general, the *ANNs* outperformed the *NLMEM* for predictions of heights in training and testing phases. The best model in both training and testing phases was the *RBPANN*-tanh, which assumed five layers in the *ANN* and three of them were hidden. The use of *ANNs* proves to be a suitable and effective approach for estimating the total tree height of Durango pine species. Additionally, incorporating unsupervised clustering analysis enhances the estimation accuracy in *ANN* models, highlighting the capabilities of artificial intelligence (AI) in this context. In conclusion, AI techniques such as *ANNs* prove to be suitable and modern statistical tools for forest modeling.

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