# Height-diameter relationship for *Pinus koraiensis* in Mengjiagang Forest Farm of Northeast China using nonlinear regressions and artificial neural network models

Nguyen Thanh Tuan<sup>1</sup>\*, Tai Tien Dinh<sup>2</sup>, Shen Hai Long<sup>3</sup>

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Abstract: Korean pine (*Pinus koraiensis* Sieb. et Zucc.) is one of the highly commercial woody species in Northeast China. In this study, six nonlinear equations and artificial neural network (ANN) models were employed to model and validate height-diameter (H-DBH) relationship in three different stand densities of one Korean pine plantation. Data were collected in 12 plots in a 43-year-old even-aged stand of *P. koraiensis* in Mengjiagang Forest Farm, China. The data were randomly split into two datasets for model development (9 plots) and for model validation (3 plots). All candidate models showed a good perfomance in explaining H-DBH relationship with error estimation of tree height ranging from 0.61 to 1.52 m. Especially, ANN models could reduce the root mean square error (RMSE) by the highest 40%, compared with Power function for the density level of 600 trees. In general, our results showed that ANN models were superior to other six nonlinear models. The H-DBH relationship appeared to differ between stand density levels, thus it is necessary to establish H-DBH models for specific stand densities to provide more accurate estimation of tree height.

Keywords: forest measurement; nonlinear growth functions; artificial intelligence technology; Korean pine plantation

Tree diameter at breast height (DBH) and tree height (H) are the most important variables used in forest inventory and management (Lei et al. 2009), particularly for estimating total tree and stand volumes (Sharma et al. 2004). In addition, tree sizes at the individual tree and stand levels are underlying variables for determining site index, timber volume, and forest structures and dynamics (Parresol, Bernard 1992; Lauri et al. 2015; Ahmadi et al. 2016). In forest inventory field, the measurement of tree height is time-consuming and costly because of the low visibility of tree tops in dense stands while diameter at breast height of a tree can

be measured quickly, easily, and accurately by basic instruments such as tree calipers and diameter tape (Staudhammer, Lemay 2000; Colbert et al. 2002; Sharma, Parton 2007; Mengesha et al. 2018). To save time and effort in forest inventory, foresters often predict tree height using height-diameter (H-DBH) models instead of direct measurement of height (Ahmadi et al. 2013).

Previous studies have used linear and nonlinear growth equations to develop H-DBH relationships for several tree species (Colbert et al. 2002; EMANUEL et al. 2018; MENGESHA et al. 2018). The nonlinear growth functions have been widely used

<sup>&</sup>lt;sup>1</sup> Department of Forestry, Vietnam National University of Forestry, Dong Nai, Vietnam

<sup>&</sup>lt;sup>2</sup>Institute of Resources and Environment, Hue University, Hue, Vietnam

<sup>&</sup>lt;sup>3</sup>School of Forestry, Northeast Forestry University, Harbin, P.R. China

<sup>\*</sup>Corresponding author: nttuan@vnuf2.edu.vn

to fit tree height-diameter models, however they may produce large extrapolation errors when applying beyond the range of model development data, thus the predictive capability of model should be carefully validated before application (ZHANG 1997). Artificial neural networks (ANNs) are computing systems in artificial intelligence technology which is based on the structures and functions of biological neural networks (ASHRAF et al. 2015). Recently, ANN models have been applied in many aspects of forest management, especially in forest modelling such as the estimation of tree volume (DIAMANTOPOULOU, MILIOS 2010), prediction of tree diameter and height (SHAO, REYNOLDS 2006; VIEIRA et al. 2018), and forest cover types (BLACK-ARD, DEAN 1999) because of its strong capacity of nonlinear mapping, high accuracy and robustness (Sheela, Deepa 2013). However, few studies have developed height-diameter (H-DBH) models using artificial neural networks (Özçelik et al. 2013). The H-DBH relationship of one tree species could vary with environmental gradients (SHARMA et al. 2004; ÖZÇELIK et al. 2014), thus the development of H-DBH model for specific site conditions is needed to provide more accurate estimation of tree height. Korean pine (Pinus koraiensis Sieb. et Zucc.) is widely distributed in Northeast China and it is one of the most valuable and commercially important timber and nut production species (JIN et al. 2017; NGUYEN et al. 2018). However, the information on H-DBH relationship in this species is limited (ZANG et al. 2016). Therefore, this

study aimed to develop H-DBH models using basic nonlinear growth equations and artificial neural networks (ANN) models to describe the height-diameter relationship in *P. koraiensis* in Mengjiagang Forest Farm, China.

## MATERIAL AND METHODS

## Study area and data description

The study was carried out in Mengjiagang Forest Farm (46°32'16"N, 129°10'36"E) in Heilongjiang Province, China. The climate is East Asian continental monsoon with long winter and dry seasons. In the study site, mean annual temperature is 2.7°C and mean annual precipitation is 550 mm. Average annual maximum and minimum temperatures are 35.6°C in July and -34.7°C in January, respectively.

The plots were established in 1973 and they were planted in a random block arrangement with 3 density levels:  $400 \text{ trees} \cdot \text{ha}^{-1}$  (N1);  $600 \text{ trees} \cdot \text{ha}^{-1}$  (N2) and  $800 \text{ trees} \cdot \text{ha}^{-1}$  (N3). In summer of 2016, 4 plots ( $40 \times 25 \text{ m}$  each) were selected to develop H-DBH models for each density level. Diameter at breast height (DBH), tree height (H) of 758 pine trees within 12 plots (4 plots  $\times$  3 density levels) were measured. For each tree, two perpendicular diameters (outside-bark 1.3 m above ground level) were measured to the nearest 0.1 cm and were then averaged to obtain DBH (cm). Total H (m) was

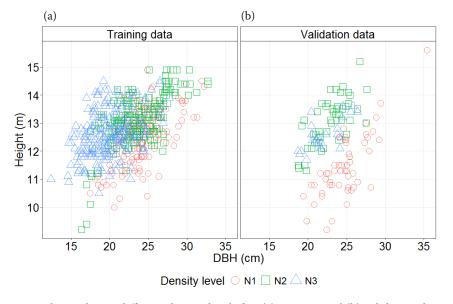


Fig. 1. Height diameter relationship in different density levels for: (a) training and (b) validation datasets

Table 1. Summary statistics for the training and the evaluation data sets

| A 44:14            | Training data set (n = | 630 trees in 9 plots) | Validation data set ( $n = 128$ trees in 3 plots) |       |
|--------------------|------------------------|-----------------------|---|-------|
| Attributes –       | DBH (cm)               | H (m)                 | DBH (cm)  | H (m) |
| Min                | 12.4                   | 9.2                   | 18.7  | 9.2   |
| Max                | 32.7                   | 14.9                  | 35.4  | 15.6  |
| Mean               | 22.29                  | 12.77                 | 23.51   | 12.22 |
| Standard deviation | 3.44                   | 0.91                  | 2.70  | 1.18  |

DBH - diameter at breast height, H - tree height

measured to the nearest 0.5 m, using a Vertex IV<sup>®</sup> (Haglof, Sweden). In order to evaluate the performance of developed H-DBH models, the data were randomly divided into a training dataset (9 plots) and a validation dataset (3 plots). Two datasets are described in Fig. 1 and summary statistics also are shown in Table 1.

## Model development and comparison

Nonlinear growth functions. A set of 6 nonlinear growth functions was selected from previously studies on H-DBH models (Table 2) because of their appropriate features such as sigmoid shape and flexibility. Besides, the parameters of these functions can be biologically interpreted (e.g., upper asymptote and change rate) as documented in the literature (Huang 1999). In this study, we employed only

functions that do not transform tree height variable to avoid problem of back transformation bias. We used *nls* function in R version 3.5.1 (R Core Team 2018) to fit 6 candidate models. For estimating the parameters, the initial values were obtained by using the package *lmfor* (Mehtatalo 2015).

Artificial neural networks. In this study, we used multilayer neural networks to develop H-DBH model in *P. koraiensis*. A proposed ANN model was designed using MATLAB Neural Network Toolbox (MathWorks, Natick, USA) and using multilayer perceptrons (MLP). In our study, the proposed neural network had one input layer, one hidden layer and one output layer. In each density level, the data of the input layer were DBH values and the data of output layer were height values in three plots used for ANN model training (Fig. 2). One remaining plot of each density level was used to validate the ANN models. The type and the

Table 2. A list of candidate H-DBH models

| Model   | References                              |
|---|---|
| Chapman-Richards: $H = 1.3 + a(1 - e^{-bDBH})^{c}$  | Richards (1959); Huang et al. (1992)    |
| Prodan: $H = 1.3 + \frac{DBH^2}{a + bDBH + cDBH^2}$ | Strand (1959)                           |
| Ratskowky: $H = 1.3 + ae^{\frac{-b}{DBH+c}}$        | Ratkowsky (1990); Huang et al. (1992)   |
| Logistic: $H = 1.3 + \frac{a}{1 + be^{-cDBH}}$      | Pearl, Reed (1920); Huang et al. (1992) |
| Gompertz: $H = 1.3 + ae^{-be^{-cDBH}}$              | Gompertz (1832); Huang et al. (1992)    |
| Power: $H = 1.3 + aDBH^b$                           | Stoffels; Van Soest (1953)              |

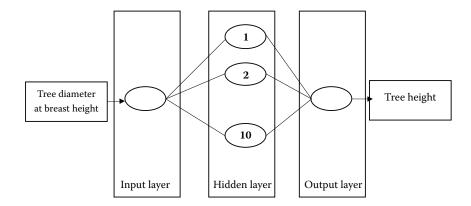


Fig. 2. Artificial neural network architectures (the model has three layers, for each density level, ten models are constructed by correspondingly varying the number of neurons in the hidden layer from 1 to 10)

complexity of the ANN model can be determined by the number of the neurons in the hidden layer. Several studies have revealed that the number of neurons in the hidden layer can not be determined directly. Generally, it should meet the following Eq. 1 (Yu, Jia 2012):

$$i = \sqrt{j + m} + R \tag{1}$$

where:

i – number of neurons in the hidden layer;

i – number of neurons in the output layer;

m – number of neurons in the input layer;

R – any number ranging from 0 to 10.

As the input layer in our study had one neuron and the output layer had one neuron, the number of neurons of the hidden layer meets the condition:  $1 \le i < 11$ . Thus, in our study, the number of neurons in the hidden layer was set from 1 to 10 to search for the best ANN models. The activation function used in the hidden layer was hyperbolic tangent sigmoid transfer function and training algorithm was the Levenberg-Marquardt Backpropagation. We set the target error as 0.01 and learning rate as 0.01 and the networks were trained for 1,000 epochs for all models as there was very negligible reduction of the root mean square error (RMSE) values after 1,000 epochs.

Model performance criteria. In the present study, three good-ness of fit statistics obtained from the residuals were examined to compare the performance of ANN models and six nonlinear models. These criteria were root mean square error (RMSE), which indicates the precision of the estimates; the coefficient of determination ( $R^2$ ), which is used to determine the relative correlation between the estimated and the measured data; and Akaike's information criterion (AIC), which is an index used to select the best model from a group of candidate models (AKAIKE 1974). The RMSE, R<sup>2</sup> and AIC were defined as in the following Eqs (2-4):

$$R^{2} = 1 - \frac{\sum (y_{i} - \widehat{y}_{i})^{2}}{\sum (y_{i} - \overline{y})^{2}}$$
 (2)

$$RMSE = \sqrt{\frac{\sum_{i=1}^{n} \left(y_i - \widehat{y_i}\right)^2}{n}}$$
 (3)

$$AIC = nln(RMSE) + 2p$$
 (4)

where:

– observed height value of the tree  $i^{th}$ ;

 $y_i$   $\hat{y}_i$ – estimated height value of the tree  $i^{th}$ ;

mean value of observed height;

- sample size,

- number of model parameters to be estimated;

RMSE - root mean square error.

For each ANN model, the AIC value can be calculated by using the Eqs (5, 6) of PANCHAL et al. (2010) and D'EMILIO et al. (2018).

$$AIC = n \ln \left( \frac{RSS}{n} \right) + 2K \quad \text{if } n/K \ge 40 \tag{5}$$

$$AIC = n \ln \left( \frac{RSS}{n} \right) + 2K + \frac{2K(K+1)}{n-K-1} \text{ if } n/K < 40$$
 (6)

where:

n – number of observations,

K – number of free parameters,

*RSS* – residual sum of square.

#### RESULTS

# **Evaluation of nonlinear models**

For six nonlinear regression models (Table 3), we found that the coefficient estimates varied among density levels. The asymptotically maximum estimated height varied approximatly from 12 to 14 m. Among nonlinear models, the Power function showed a statistical significance in all coefficient estimates. Except for the parameter *c*, all coefficient estimates of Prodan model were not statistically significant. The results of goodness of fit statistics for the training and validation datasets are shown in Fig. 3. In all six nonlinear models, there was a small difference in RMSE values between training and validation data. Power model exhibited a smaller RMSE values in the validation dataset, compared with other nonlinear models. In general, Power model had the highest value of  $\mathbb{R}^2$ , except for N3 density level where Ratskowky model showed the highest  $R^2$  for the both training (0.742) and validation data (0.612). The AIC values of Power model were slighly smaller than those of other nonlinear models for both training and validation data, and for all stand density levels.

# ANN models

In this study, the number of neurons in the hidden layer of ANN models used for training process ranged from 1 to 10. In training dataset of N1 and N3 density levels, the AIC values tended to increase with the number of neurons in the hidden layer. In validation dataset, however, the AIC values increased with the neuron number for N1 and N2 density levels (Fig. 3). We observed that the coefficient of determination ( $R^2$ ) of all models varied from 0.61 to 0.77, and the differences in  $R^2$  values were found between the six nonlinear models and ANN models in each stand density level (Fig. 3).

For N1 and N2 density levels, the best model performance was achieved by using 10 neurons in the hidden layer, showing the smallest RMSE values for both training dataset (0.745 and 0.564, respectively) and validation dataset (1.277 and 0.610, respectively). Meanwhile, in N3 density level, the best model performance was observed in the ANN model using 7 neurons in hidden layer for both training dataset (RMSE = 1.279) and validation dataset (RMSE = 0.743).

In N1 density level, the RMSE values of validation data ranged from 0.743 to 0.745 and from 0.750 to 0.763 in ANN models and nonlinear models, respectively. In N2 density level, we found that the RMSE values of the ANN model using 10 neurons in the hidden layer were reduced by approximately 17 and 40% for validation and training data, respectively, compared to Power model. In N3 density level, the RMSE of the ANN model with 7 neurons in the hidden layer was slightly lower than those of nonlinear models for both training and validation data.

## **DISCUSSION**

Tree height is fundamental variable in forest management for developing growth model, biomass estimation, and for determining stand structure and dynamics (Diamantopoulou, Özçelik 2012; NAVROODI et al. 2016). In our study, six non-linear growth functions selected for estimating total tree height performed quite well with small RMSE values (0.651 to 1.517) and fairly well of  $R^2$  values (0.592-0.744) for all stand density levels. Overall, our results supported the findings of previous studies, indicating the appropriateness of nonlinear models in describing H-DBH relationship (DIA-MANTOPOULOU, ÖZÇELIK 2012; ÖZÇELIK et al. 2014; Mengesha et al. 2018; Emanuel et al. 2018). In general, H-DBH relationship has been considered to approach a sigmoidal shape (ÖZÇELIK et al. 2014; Costa et al. 2016; Ferraz et al. 2018), however the best fit model appeared to vary with tree species and site conditions. For example, the best model was the logistic growth function for Pinus kesiya in Benguet province, Philippines (Lumbres et al. 2013); was the Gompertz function for three economically important tree species of southern Turkey (Özçelik et al. 2014); was the Weibull function for Cupressus lusitanica in Gergeda Forest, Ethiopia (MENGESHA et al. 2018). In previous study on H-DBH model of Korean pine, Richards equation with four parameters appeared to explain well H-DBH relationship in Nature forest of Jilin province, China (DENG et al. 2001). ZANG et al. (2016) used Chapman-Richards function to describe the H-DBH relationship for Pinus koraiensis in mixed Mongolian oak-deciduous stands and mixed spruce-fir-deciduous stands in Wangqing Forestry Bureau, Jilin Province of Northeast China. On the contrary, our study demonstrated that Power model was the best fit model among

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Table 3. Parameter estimates for the basic H-DBH nonlinear models

|                  |          |        |         |                 |          | Parameter estimates | estimates |                 |          |       |                 |             |
|------------------|----------|--------|---------|-----------------|----------|---------------------|-----------|-----------------|----------|-------|-----------------|-------------|
| Model            |          |        | a       |                 |          | 9                   | ,         |                 |          | 0     |                 |             |
| I                | Estimate | SE     | t-value | <i>P</i> -value | Estimate | SE                  | t-value   | <i>P</i> -value | Estimate | SE    | <i>t</i> -value | P-value     |
| N1 density level |          |        |         |                 |          |                     |           |                 |          |       |                 |             |
| Chapman-Richards | 12.517   | 0.453  | 27.611  | < 2e-16***      | 0.182    | 0.061               | 2.983     | 0.003**         | 7.410    | 8.061 | 0.919           | 0.35929     |
| Prodan           | 25.994   | 14.548 | 1.787   | 0.075763        | -1.263   | 1.217               | -1.038    | 0.300           | 0.095    | 0.025 | 3.768           | 0.000227*** |
| Ratskowky        | 13.904   | 0.923  | 15.057  | < 2e-16***      | 2.426    | 1.383               | 1.754     | 0.081           | -11.946  | 2.879 | -4.150          | 5.26e-05*** |
| Logistic         | 12.432   | 0.393  | 31.615  | < 2e-16***      | 13.543   | 14.486              | 0.935     | 0.351           | 0.208    | 0.060 | 3.455           | ***26900000 |
| Gompertz         | 12.501   | 0.434  | 28.785  | < 2e-16***      | 8.060    | 8.230               | 0.979     | 0.328           | 0.185    | 0.058 | 3.195           | 0.001**     |
| Power            | 2.927    | 0.441  | 6.642   | 4.00e-10***     | 0.425    | 0.047               | 8.986     | 4.78e-16***     |          |       |                 |             |
| N2 density level |          |        |         |                 |          |                     |           |                 |          |       |                 |             |
| Chapman-Richards | 13.686   | 0.773  | 17.694  | < 2e-16***      | 0.108    | 0.047               | 2.306     | 0.0221*         | 1.676    | 1.227 | 1.367           | 0.173       |
| Prodan           | 8.565    | 7.500  | 1.142   | 0.255           | -0.009   | 0.655               | -0.014    | 0.989           | 0.069    | 0.014 | 4.857           | 2.3e-06***  |
| Ratskowky        | 13.236   | 0.256  | 51.673  | < 2e-16***      | 0.908    | 0.240               | 3.781     | 0.000203***     | -14.328  | 0.668 | -21.463         | < 2e-16***  |
| Logistic         | 13.559   | 0.623  | 21.778  | < 2e-16***      | 2.913    | 1.672               | 1.742     | 0.08292         | 0.129    | 0.040 | 3.195           | 0.001**     |
| Gompertz         | 13.502   | 0.587  | 23.008  | < 2e-16***      | 2.331    | 1.368               | 1.705     | 69680.0         | 0.124    | 0.040 | 3.115           | 0.002**     |
| Power            | 3.623    | 0.343  | 10.580  | < 2e-16***      | 0.375    | 0.030               | 12.580    | < 2e-16***      |          |       |                 |             |
| N3 density level |          |        |         |                 |          |                     |           |                 |          |       |                 |             |
| Chapman-Richards | 14.032   | 8.829  | 1.589   | 0.113           | 0.029    | 0.146               | 0.200     | 0.841           | 0.264    | 0.419 | 0.631           | 0.529       |
| Prodan           | 1.010    | 5.330  | 0.190   | 0.85            | 0.233    | 0.559               | 0.416     | 0.678           | 0.074    | 0.015 | 5.105           | 6.77e-07*** |
| Ratskowky        | 13.098   | 1.218  | 10.751  | < 2e-16***      | 2.283    | 2.681               | 0.851     | 0.395           | -4.226   | 8.344 | -0.507          | 0.613       |
| Logistic         | 12.913   | 2.086  | 6.191   | 2.58e-09***     | 0.686    | 0.348               | 1.973     | 0.049*          | 0.079    | 0.089 | 0.884           | 0.377       |
| Gompertz         | 12.862   | 1.956  | 6.574   | 3.04e-10***     | 0.587    | 0.351               | 1.670     | 960.0           | 920.0    | 0.089 | 0.858           | 0.392       |
| Power            | 6.301    | 0.590  | 10.685  | < 2e-16***      | 0.195    | 0.031               | 6.226     | 2.12e-09***     |          |       |                 |             |

\*significant at P < 0.05; \*\*significant at P < 0.005 level; \*\*significant at P < 0.005 level; SE – standard errors; a, b, c – parameters of the equation

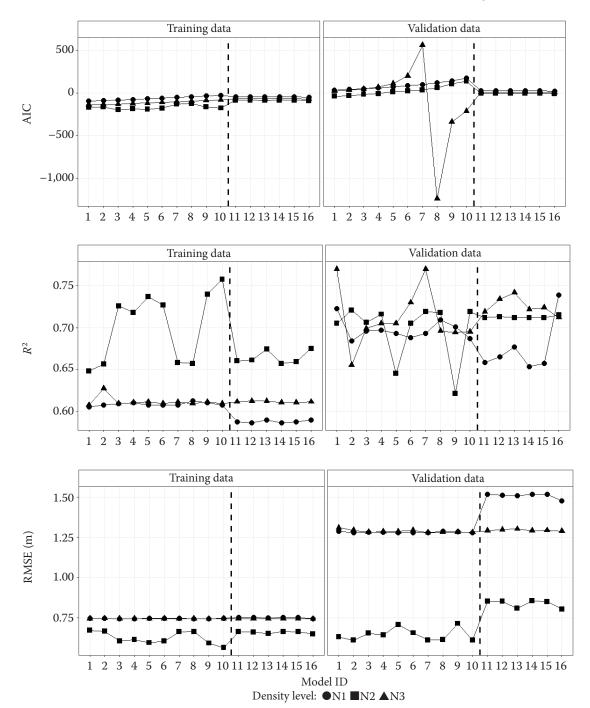


Fig. 3. Goodness of fit statistics for 10 ANN models and 6 nonlinear models. The statistic values are shown for both training and validation datasets. Model ID from 1 to 10 refers to ANN models with corresponding neuron number in the hidden layer. Model ID from 11 to 16 refers to the model of Chapman-Richards, Prodan, Ratskowky, Logistic, Gompertz and Power, respectively. RMSE,  $R^2$ , AIC are root mean square error, coefficient of determination and Akaike's Information Criterion, respectively.

6 candidate nonlinear equations evaluated in terms of RMSE and AIC values for Korean pine plantation in Mengjiagang Forest Farm, China. The inconsistency between our study and others could be explained by the differences in site condition and

in stand structure and age, which can affect the H-DBH relationship (ÖZÇELIK et al. 2014).

Our data showed that coefficient estimates and model performance criteria varied among three stand density levels. The performance of H-DBH models

tended to differ between density levels, and this could be explained by the variation in local environment and stand conditions (Sharma et al. 2004). Stand density is an important factor affecting tree growth in coniferous species (Mäkinen, Isomäki 2004). Low stand density may reduce the competition among individual trees, which results in advantageous conditions for tree growth (Woodruff et al. 2000). Moreover, low tree density might increase soil moisture availability and generate favorable physiological responses, whereby affecting tree growth (Nguyen et al. 2018). In accordance with previous studies (Krieger 1998; Ritchie et al. 2012), our data indicated that H-DBH relationship of *P. koraiensis* could be conditioned on stand density.

Compared with stand density of 600 trees·ha<sup>-1</sup> and 800 trees·ha-1, tree height in the stand density of 400 trees⋅ha<sup>-1</sup> tended to be smaller. Similary, Zeide, VANDERSCHAAF (2002) documented that Loblolly pine trees in higher density stand were taller than those in lower density stand. It is likely that the competition for light between a tree and its neighbors may promote the height growth (DORN et al. 2000). According to Lauri et al. (2015), this phenomenon can be explained by the biological growth rhythm of trees in dense stands that they firstly focus on height growth to compete for light, and then use their resources to increase diameter. This strategy may result in a higher height variation in small-diameter stand compared with those of large-diameter stand. The influence of stand density on the H-DBH relationship was also found in many studies (e.g. Krieger 1998; STAUDHAMMER, LEMAY 2000; RITCHIE et al. 2012). For instance, Krieger (1998) constructed H-DBH models for 6 stand density levels (from 100 to 1,100 trees·ha<sup>-1</sup>) for white spruce species. In our study, the difference between AIC and RMSE values for model performance were observed (Fig. 3). The contradiction of RMSE and AIC has been also reported in previous work (KINGSTON et al. 2008; Luo et al. 2018). Compared with other model performance criteria, RMSE is commonly implemented to select the best model (ÖZÇELIK et al. 2013; COSTA et al. 2016, ZANG et al. 2016; FERRAZ et al. 2018), thus in this study, we determined RMSE as the main goodness of fit statistic for mode evaluation. Based on the RMSE values, we suggest using the ANN model with 10 neurons in the hidden layer for stand density of 400 trees ha<sup>-1</sup> and 600 trees ha<sup>-1</sup>, and the ANN model with 7 neurons in the hidden layer for stand density of 800 trees· $ha^{-1}$  for H-DBH relationship of Korean pine in our study site. In the present study, ANN models performed superior to non-linear growth functions in explaining H-DBH relationship in P. koraiensis, as indicated by small RMSE for all stand density levels, This finding is consistent with previous studies indicating that ANN models can be an effective alternative to regression analysis, providing an accurate tree height estimation for forest inventory (ÖZÇELIK et al. 2013; Zhi, Gan 2017; Vieira et al. 2018). Besides, the model developed in this study showed the better overall performance in terms of RMSE compared with other approaches in previous studies such as using nonlinear regression for estimating tree height of Crimean juniper with RMSE values ranging from 1.28 to 1.89 (Özçelik et al. 2013), and using nonlinear mixed effects models for Pinus koraiensis with RMSE values ranging from 1.40 to 2.65 and from 0.95 to 2.67 in training data and validaton data, respectively (ZANG et al. 2016). Although our study referred only to the specific three stand density levels in Mengjiagang Forest Farm, the obtained results could be a basis for further research by highlighting the applicability of ANN models in tree height prediction and this would provide forest managers and researchers with useful guidelines for predicting tree height of Pinus koraiensis in the research site.

# **CONCLUSIONS**

Nonlinear growth functions have been commonly used for modeling tree height-diameter relationship. This study used 6 non-linear functions and ANN models to describe H-DBH relationship of *P. koraiensis* in three different stand densities in Mengjiagang Forest Farm, China. Based on RMSE and *R*<sup>2</sup> statistics, we showed that the ANN models were superior to all examined nonlinear models. Our data suggest that it is necessary to establish H-DBH models for different stand densities of *P. koraiensis* to provide more accurate estimates of tree height. The ANN models provide a new approach for tree height estimation and could be implemented at individual tree and stand levels for *P. koraiensis* in this studied region.

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