

FITTING A TAPER FUNCTION TO MINIMIZE THE SUM OF ABSOLUTE DEVIATIONS

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ABSTRACT: Multiple product inventories of forests require accurate estimates of the diameter, length and volume of each product. Taper functions have been used to precisely describe tree form, once they provide estimates for the diameter at any height or the height at any diameter. This study applied a goal programming technique to estimate the parameters of two taper functions to describe individual tree forms. The goal programming formulation generates parameters that minimize total absolute deviations (*MOTAD*). These parameters generated by the *MOTAD* method were compared to those of ordinary least squares (*OLS*) method. The analysis used a set of 178 trees cut from cloned eucalyptus plantations in the Southern part of the state of Bahia, Brazil. The values of the estimated parameters for the two taper functions resulted very similar when the two methods were compared. There was no significant difference between the two fitting methods according to the statistics used to evaluate the quality of the generated estimates. *OLS* and *MOTAD* resulted equally precise in the estimation of diameters and volumes outside and inside bark.

Key words: MOTAD, ordinary least squares, goal programming, linear regression

AJUSTE DE UMA FUNÇÃO DE AFILAMENTO VIA MINIMIZAÇÃO DA SOMA DOS DESVIOS ABSOLUTOS

RESUMO: Os inventários florestais para múltiplos produtos requerem estimativas exatas do diâmetro, comprimento e volume de cada produto. As equações de afilamento têm sido usadas para descrever precisamente a forma da árvore uma vez que estas funções fornecem estimativas de diâmetro a qualquer altura ou de altura em qualquer diâmetro. Este trabalho aplica um modelo de programação por metas para estimar os parâmetros de duas equações de afilamento para descrever a forma do tronco de árvores individuais. O modelo de programação por metas gera parâmetros que minimizam a soma dos desvios absolutos (*MOTAD*). Esses parâmetros gerados pelo método *MOTAD* foram comparados aos parâmetros gerados pelo método dos mínimos quadrados ordinários (*OLS*). A análise se baseou em dados de cubagem de 178 árvores obtidas em plantios clonais de eucaliptos conduzidos na região sul da Bahia. Os valores dos parâmetros estimados por ambos os métodos de ajuste para as duas funções de afilamento mostraram-se muito semelhantes. Não houve diferença significativa entre os indicadores usados para avaliar a qualidade dos parâmetros estimados pelos dois métodos de ajuste. Os métodos *OLS* e *MOTAD* mostraram-se igualmente precisos na estimação de diâmetros e volumes com casca e sem casca.

Palavras-chave: minimização do desvio absoluto total, mínimos quadrados ordinários, programação por metas, regressão linear

INTRODUCTION

The quality of a good forest management plan relies on the precision level of the biometric system used to estimate future tree volumes. The biometric system contains independent variables which are measurable tree characteristics such as diameter, height,

form and basal area usually measured to monitor forest growth (Scolforo, 1993). The term “taper” is applied to the rate of decrease in diameter along the trunk. Taper functions provide estimates for the diameter at any height or the height at any diameter.

Tree form and size determines different outputs and taper-functions have been used to precisely

describe these tree characteristics (Ahrens & Holbert, 1981; Husch et al., 1972; Lima, 1986; Assis, 2000). The vertical integration of production activities in forest companies, where outputs from one production stage become input to the next stage, turns precise tree volume estimation even more relevant (Ahrens & Holbert, 1981; Assis, 2000). Therefore, taper functions become the primary tool for estimating the volume at any part of the trunk, by means of the mathematical integration of the section area along the tree axle.

Pioneer studies using mathematical functions to describe the trunk form date from 1903 and were developed by Höjer. With the advance of computing systems, more complex models, including polynomials, segmented and not segmented, considering sigmoid shaped forms were developed. Among them we can cite those developed by Kozak et al. (1969), Demaerschalk (1973), Ormerod (1973), Max & Burkhart (1976), Clutter (1980), Lima (1986), Guimarães & Leite (1992), and Andrade & Leite (1998).

In Brazil, the use of taper functions is more recent. Among the published studies, Ahrens & Holbert (1981), Lima (1986), McTague et al. (1989), Guimarães & Leite (1992), Leite et al. (1995), Schneider et al. (1996), Fischer (1997), Andrade & Leite (1998), and Assis (2000), the main question is how well different taper functions fit the data, how well they represent the diameters at different heights and vice-versa.

The parameters of a model representing the form of a tree are determined by specific fitting techniques, among which ordinary least squares has been the most frequently used fitting method. The objective of this work is to apply Goal Programming (GP) techniques as a fitting method to estimate the coefficients of two polynomial models used as taper functions and to compare the results of this fitting method with the estimates produced with the least square fitting method. This study contributes to the development of taper function fitting methods and to the analysis of such processes.

METODOLOGY

Area Characterization

Data used in this study proceed from plantations of cloned *Eucalyptus grandis* × *Eucalyptus urophylla* located in the South of Bahia, municipality of Eunápolis, Brazil (16°17'59"S; 39°28'42"N; altitude 168 m). The regional climate (Köppen) is of the *Af* type, hot and humid tropical, without dry seasons, with annual average temperature of 23.1°C and average rainfall of 1250 mm year⁻¹.

Tree Volume Definition

One hundred seventy eight *E. grandis* × *E. urophylla* trees, felled at age 5, with heights (H) varying from 20 to 30 m and diameter at breast height (DBH) varying from 9.23 to 23.00 cm had their volumes rigorously determined. Data, collected for each tree, included circumference at breast height (CBH), total height (H), log length (h) and circumference of the log's largest base inside bark (CIB_i) and outside bark (COB_i) for every log *i*.

The measurements of CIB and COB along tree trunks were made every meter, starting at 0.3 m from the soil (stump), resulting in a total of 4,333 observations. The volume of the section corresponding to the top of the tree was calculated taking the cone formula as a guide and the volumes of the other sections were calculated based on the Smalian formula. The total volume (outside and inside bark) was obtained by the sum of the volumes of the different sections of the tree.

Model fitting approach

Two different polynomial models were used in this study to relate all diameters taken along the trunk and respective heights with DBH and DAB (diameter at the base of the tree) and H. A detailed description of these models follows.

i) Model 1

The polynomial model 1 (M1) can be represented, mathematically, as follows:

$$\left(\frac{d}{DAB}\right)^2 = \beta_0\left(\frac{L}{H}\right)^2 + \beta_1\left(\frac{L}{H}\right)^3 + \beta_2\left(\frac{L}{H}\right)^4 + \varepsilon \quad (1)$$

where: d = diameter at height h from the soil (cm); $L = H - h$; β_i = parameters to be estimated; ε = estimation error.

Isolating d , a taper function is obtained to estimate the correspondent diameter at any height on the tree, if DAB, H and L are given.

$$d^2 = DAB^2 \left[\beta_0\left(\frac{L}{H}\right)^2 + \beta_1\left(\frac{L}{H}\right)^3 + \beta_2\left(\frac{L}{H}\right)^4 \right] \quad (2)$$

Considering the sectional area (A) of a tree with diameter d (m; at height h) equal to $(\pi/40000)d^2$ and the integration of this section along length L , we obtain the compatible volume equation:

$$V = \int_{L_2}^{L_1} \left(\frac{\pi}{40000} \right) d^2 dL \quad (3)$$

Substituting (1) into (2) and integrating we obtain:

$$V = \left(\frac{\pi}{40000} \right) DAB^2 \left\{ \beta_0 \left[\frac{(L_1^3 - L_2^3)}{3H^2} \right] + \beta_1 \left[\frac{(L_1^4 - L_2^4)}{4H^3} \right] + \beta_2 \left[\frac{(L_1^5 - L_2^5)}{5H^4} \right] \right\} \quad (4)$$

Setting $L_2 = 0$ (top of the tree) and $L_1 = H$ (base of the tree), the volume equation for the whole tree becomes:

$$V = \frac{\pi}{40000} DAB^2 H \left[\beta_0 \frac{1}{3} + \beta_1 \frac{1}{4} + \beta_2 \frac{1}{5} \right] \quad (5)$$

and the equation used to estimate volumes at heights $h_i = 3, 6$ and 12 m is:

$$V = \frac{\pi}{40000} DAB^2 \left\{ \beta_0 \left[\frac{H^3 - (H - h_i)^3}{3H^2} \right] + \beta_1 \left[\frac{H^4 - (H - h_i)^4}{4H^3} \right] + \beta_2 \left[\frac{H^5 - (H - h_i)^5}{5H^4} \right] \right\} \quad (6)$$

ii) Model 2

Polynomial model 2 (M2) can be represented, mathematically, as follows:

$$\begin{aligned} \left(\frac{d}{DBH} \right)^2 &= \beta_0 \left(\frac{H-h}{H-1.3} \right)^2 + \beta_1 \left(\frac{H-h}{H-1.3} \right)^3 + \\ &+ \beta_2 \left(\frac{H-h}{H-1.3} \right)^4 + \varepsilon \end{aligned} \quad (7)$$

Isolating d , we obtain the taper function to estimate the correspondent diameter at any height in the tree, if DBH , H and h are given.

$$\begin{aligned} d^2 &= DBH^2 \left[\beta_0 \left(\frac{H-h}{H-1.3} \right)^2 + \beta_1 \left(\frac{H-h}{H-1.3} \right)^3 + \right. \\ &\left. + \beta_2 \left(\frac{H-h}{H-1.3} \right)^4 \right] \end{aligned} \quad (8)$$

Setting $\frac{H-h}{H-1.3} = z$, the model becomes:

$$d^2 = DBH^2 [\beta_0 z^2 + \beta_1 z^3 + \beta_2 z^4] \quad (9)$$

As in M1, integration of section $(\pi/40000)d^2$ along length L (equation 3) results in the following compatible volume equation:

$$\begin{aligned} V &= \frac{\pi}{40000} DBH^2 \left\{ \beta_0 \left[\frac{(H-h_1)^3 - (H-h_2)^3}{3(H-1.3)^2} \right] + \right. \\ &\left. + \beta_1 \left[\frac{(H-h_1)^4 - (H-h_2)^4}{4(H-1.3)^3} \right] + \beta_2 \left[\frac{(H-h_1)^5 - (H-h_2)^5}{5(H-1.3)^4} \right] \right\} \end{aligned} \quad (10)$$

Again, setting $h_2 = H$ (top) and $h_1 = 0$ (tree base), the volume equation for the whole tree becomes:

$$\begin{aligned} V &= \frac{\pi}{40000} DBH^2 \left\{ \beta_0 \left[\frac{H^3}{3(H-1.3)^2} \right] + \beta_1 \left[\frac{H^4}{4(H-1.3)^3} \right] + \right. \\ &\left. + \beta_2 \left[\frac{H^5}{5(H-1.3)^4} \right] \right\} \end{aligned} \quad (11)$$

Equations used to estimate volumes at heights $h_i = 3, 6$ and 12 m are:

$$\begin{aligned} V &= \frac{\pi}{40000} DBH^2 \left\{ \beta_0 \left[\frac{H^3 - (H - h_i)^3}{3(H-1.3)^2} \right] + \right. \\ &\left. + \beta_1 \left[\frac{H^4 - (H - h_i)^4}{4(H-1.3)^3} \right] + \beta_2 \left[\frac{H^5 - (H - h_i)^5}{5(H-1.3)^4} \right] \right\} \end{aligned} \quad (12)$$

Data stored formed the “base” file, which was processed by a SAS[®] routine to generate linear regression estimates and by LINDO[®] to process the goal programming model.

β_i Parameter Estimation

Estimating β_i parameters involves the generation of the minimum possible error (ε). The minimum loss function can be defined as the differences between the observed data and the data estimated by the model.

In the ordinary least squares (*OLS*) method, the loss function to be minimized is set as the *sum of squared residuals*. Squaring the residues avoids residue canceling but weights more heavily large residues and emphasizes their importance (Batista, 1998).

In goal programming, the loss function used to estimate the β_i parameters is the *sum of absolute residuals* and the method is referred to as the *MOTAD* (minimization of total absolute deviations) method. The *MOTAD* method also avoids residue canceling, but differently to the *OLS* method large residues have the same importance as small residues (Batista, 1998).

Ignizio & Cavalier (1994) discuss the use goal programming as an alternative tool for developing predictive function. According to these authors, the *OLS* method is more frequently employed simply because it is easy to be applied and because it generates confi-

dence intervals based on the assumption that errors are normally distributed with equal variances, an assumption that sometimes may not hold. Alcântara et. al (2003) points out that the *MOTAD* method overcomes a deficiency in the *OLS* method when outliers are present in the data set due to *MOTAD*'s lower sensitivity to extreme values.

Model fitting with the OLS method

The models adopted in the present study can be written as multiple linear regression models with three predictor variables, as follows:

$$Y = \beta_0 X_1 + \beta_1 X_2 + \beta_2 X_3 + \epsilon \quad (13)$$

where: $Y = (d/DAB)^2$ or $(d/DBH)^2$; $X_1 = (L/H)^2$ or $[(H-h)/(H-1.3)]^2$; $X_2 = (L/H)^3$ or $[(H-h)/(H-1.3)]^3$; $X_3 = (L/H)^4$ or $[(H-h)/(H-1.3)]^4$

The condition $\beta_0 + \beta_1 + \beta_2 = 1$ was imposed to enforce coherence. This is needed because $(L/H)^2$, $(L/H)^3$ and $(L/H)^4$ are equal to 1 when L is equal to H , and also d equals to DAB , resulting consequently $(d/DAB)^2$ equal to 1. The same reasoning can be used for M2.

Model fitting with the MOTAD method

The *MOTAD* method formulates the minimization of the sum of absolute deviations as a goal programming problem (GP), a mathematical formulation for constrained multiple objectives. Deviations to these objectives are minimized, generating solutions close to certain aspiration levels. Aspiration levels are in fact goals associated with the multiple objectives and, therefore, the name goal programming.

The Simplex algorithm for solving linear programming problems is normally used in the solution of GP problems. These problems can also be formulated under the same hypothesis, limitations and conditions of linear programming: linearity, divisibility and deterministic characteristic (Lee et al., 1990).

The first application of GP to constrained regression was formulated by Charnes et al. (1955). Two deviations, DP_i and DN_i , are created to each pair of observations (X_i, Y_i) . DP_i represents a positive deviation and DN_i represents a negative deviation. So, the linear model $Y_i = f(X_i)$ becomes $0 = f(X_i) - DP_i + DN_i - Y_i$.

In the MOTAD version of the GP problem applied to the taper function fitting process, i identifies each observation in a set of N measurements, the coefficients of the linear model are the main decision variables and the GP formulation becomes:

$$\text{Min } Z = \sum_{i=1}^N (DP_i + DN_i) \quad (14)$$

subject to $0 = f(X_i) - DP_i + DN_i - Y_i$ ($i = 1, 2, \dots, N$)

For M1, the constraints can be represented as:

$$DN_i - DP_i + \left(\frac{L}{H} \right)_i^2 \beta_0 + \left(\frac{L}{H} \right)_i^3 \beta_1 + \left(\frac{L}{H} \right)_i^4 \beta_2 - \left(\frac{d}{DAB} \right)_i^2 = 0$$

and for M2, the expression becomes

$$DN_i - DP_i + \left[\frac{(H-h)}{(H-1.3)} \right]_i^2 \beta_0 + \left[\frac{(H-h)}{(H-1.3)} \right]_i^3 \beta_1 + \left[\frac{(H-h)}{(H-1.3)} \right]_i^4 \beta_2 - \left(\frac{d}{DBH} \right)_i^2 = 0$$

where: $i = i^{\text{th}}$ tree log and i^{th} line in the model;

DN_i = negative deviation of observation i in relation to

$$\left(\frac{d}{DAB} \right)_i^2 \text{ or } \left(\frac{d}{DBH} \right)_i^2;$$

DP_i = positive deviation of observation i in relation to

$$\left(\frac{d}{DAB} \right)_i^2 \text{ or } \left(\frac{d}{DBH} \right)_i^2;$$

$\left(\frac{d}{DAB} \right)_i^2$ and $\left(\frac{d}{DBH} \right)_i^2$ are goals;

$$\left(\frac{L}{H} \right)_i^2, \left(\frac{L}{H} \right)_i^3, \left(\frac{L}{H} \right)_i^4, \left[\frac{(H-h)}{(H-1.3)} \right]_i^2, \left[\frac{(H-h)}{(H-1.3)} \right]_i^3$$

and $\left[\frac{(H-h)}{(H-1.3)} \right]_i^4$ are coefficients of the parameters to be estimated; and

β_0 , β_1 and β_2 are the parameters to be estimated.

Comparing the OLS and MOTAD methods

The coefficient of determination (R^2), the root of the mean square error (Syx) and the dispersion of residuals were used to compare the results of the fitting process. Precision and accuracy analysis were also made based on the estimative generate by each fitting method and according to four statistics used by Parresol et al. (1987) and Assis et al. (2002): bias (\bar{D}), standard deviation of differences (SD), sum of squared

relative residuals (SSRR) and residuals percentage (RP), where:

$$\text{Bias (D): } \bar{D} = \frac{\sum(d_{\text{observed}} - d_{\text{estimated}})}{N} \quad (15)$$

Standard deviation of Differences (SD):

$$SD = \sqrt{\frac{\sum((d_{\text{observed}} - d_{\text{estimated}}) - \bar{D})^2}{(N-1)}} \quad (16)$$

Sum of Squared Relative Residuals (SSRR):

$$SSRR = \sum \left(\frac{(d_{\text{observed}} - d_{\text{estimated}})}{d_{\text{observed}}} \right)^2 \quad (17)$$

Residuals percentage (RP):

$$RP = \frac{\sum \left(\frac{(d_{\text{observed}} - d_{\text{estimated}})}{d_{\text{observed}}} \right) * 100}{N} \quad (18)$$

Both models and the fitting methods were compared and also ranked for the quality of the estimates obtained for volume outside and inside bark (V_{ob} and V_{ib}). The model with the worst value for each of the calculated characteristic scored 1, otherwise the score was 2. The sum of scores for each model determined its final performance score.

Table 1 - Statistics and estimated coefficients for models given by equations (1) and (7).

Model	Fitting Method	β_0	β_1	β_2	R^2	Sy.x
					----- % -----	
1	Outside bark	MOTAD	3.988983	-7.015488	4.026512	96.04
		OLS	4.071990	-7.235140	4.163150	98.83
	Inside bark	MOTAD	3.960312	-6.109268	3.148956	97.23
		OLS	3.887129	-5.949500	3.062370	99.20
2	Outside bark	MOTAD	4.075310	-5.809204	2.733894	96.34
		OLS	4.332605	-6.579273	3.246668	98.95
	Inside bark	MOTAD	3.795416	-6.106291	3.310874	95.76
		OLS	3.688512	-5.834353	3.145841	98.78

Table 2 - Sum of squared deviations (SSD) and sum of absolute deviations (SAD) for OLS and MOTAD fitting methods for models given by equations (1) and (7).

Model		SSD		SAD	
		OLS	MOTAD	OLS	MOTAD
1	Outside bark	11.3426	11.3636	164.1304	163.9313
	Inside bark	9.4168	9.4404	155.7328	155.5305
2	Outside bark	17.1846	17.6491	162.4276	158.2298
	Inside bark	14.4022	14.4327	169.0957	168.7593

RESULTS AND DISCUSSION

Model fitting

The tested models fitted adequately to data given that both models resulted in coefficients of determination above 96% (Table 1). As expected, models fitted with the MOTAD method produced coefficients of determination slightly lower than models fitted with the OLS method. The root of the mean square error ranged from 10.58% to 13.10% and resulted very similar when comparing the two fitting methods. The *t* and *F* tests were significant at 5%, showing a strong correlation between the dependent variable (d/DAB or d/DBH) and the independent variables (L/H or $[(H-h)/H-1.3]$).

For diameters close to DBH and above, M2 shows residuals more clustered around zero (graphs *c*, *d*, *g* and *h*, Figure 1). Graphs (*a*), (*b*), (*e*) and (*f*) in Figure 1 show that outside bark M1 taper function with the dependent variable $(d/DAB)^2$ represents better the form of the trunk at the base of the tree. At the base of the tree, the dependent variable $(d/DBH)^2$ is overestimated by M2 when fitting outside bark data (graphs *c* and *d*, Figure 1) and underestimated by M2 when fitting inside bark data (graphs *g* and *h*, Figure 1).

Loss Function Minimization

Table 2 shows very small differences between the two models and the two fitting methods. Models fitted for inside bark diameters produced smaller residuals sums.

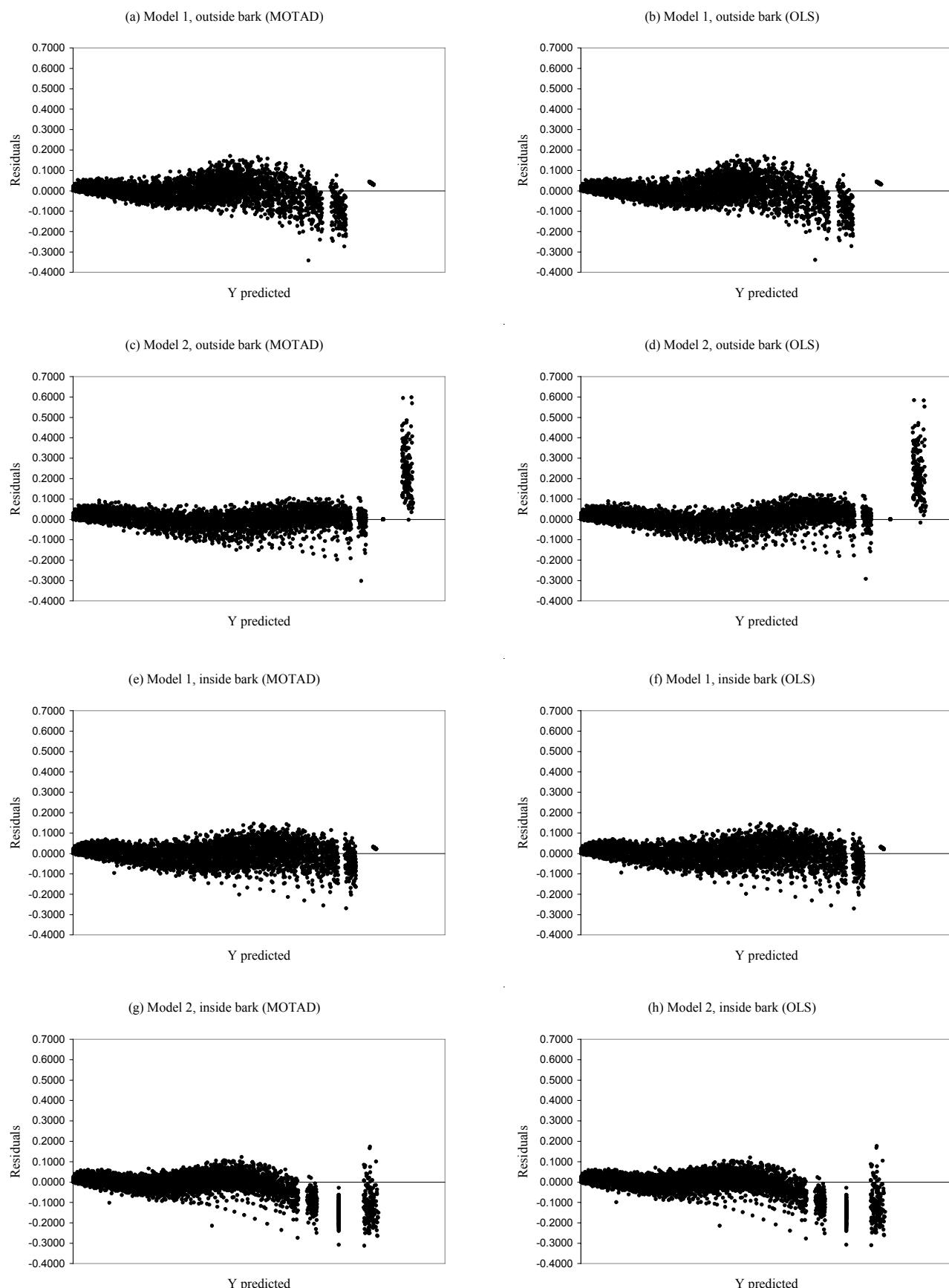


Figure 1 - Dispersion of residuals for the two models (1 and 2) and for the two fitting methods (MOTAD and OLS).

As expected, linear regression effectively resulted in the minimum sum of squared deviations while goal programming produced the minimum sum of absolute deviations.

Diameter, V_{ib} and V_{ob} estimation

Table 3 shows the values of the four selected statistics to evaluate the quality of estimates for diameter at different heights: bias (\bar{D}), standard deviation of differences (SD), sum of squared relative residuals ($SSRR$) and residuals percentage (RP).

The models showed to be equally precise in the estimation of diameters outside bark at relative heights equal to 3, 6 and 12 m. This could be observed for the *OLS* method, as well as for the *MOTAD* method. Considering the total height of the trees, M1 was slightly superior to M2.

Low values were observed for bias. Although low and considering the estimated diameters at 3, 6 and 12 m heights, this statistic shows that M1 tended to slightly overestimate the diameters outside and inside bark; M2 tended to slightly underestimate the diameters outside bark and overestimate the diameters

Table 3 - Bias (D), standard deviation of difference (SD), sum of squared relative residuals ($SSRR$), and residue percentage (RP) for diameter estimation at different heights.

Model	Section length	Fitting Method	Diameter			
			D	SD	$SSRR$	RP
1 (outside bark)	H = 3 m	MOTAD	-0.003882	0.009343	2.474496	-2.490603
		OLS	-0.003642	0.009149	2.400416	-2.331850
	H = 6 m	MOTAD	-0.001796	0.008566	4.030625	-1.045359
		OLS	-0.001503	0.008455	3.973617	-0.844549
	H = 12 m	MOTAD	-0.000591	0.007626	7.253995	-0.321753
		OLS	-0.000433	0.007573	7.234619	-0.224043
	H = Total	MOTAD	0.000235	0.007408	57.625167	2.178148
		OLS	0.000125	0.007387	56.411772	1.935972
	H = 3 m	MOTAD	-0.001345	0.005460	1.020240	-0.901078
		OLS	-0.001370	0.005569	1.024279	-0.917760
1 (inside bark)	H = 6 m	MOTAD	-0.000539	0.005571	2.095854	-0.316954
		OLS	-0.000525	0.005632	2.103390	-0.303103
	H = 12 m	MOTAD	-0.000559	0.005779	5.459162	-0.536793
		OLS	-0.000406	0.005795	5.375176	-0.391826
	H = Total	MOTAD	0.000384	0.006202	59.106371	2.240221
		OLS	0.000660	0.006186	60.227470	2.625113
	H = 3 m	MOTAD	0.004212	0.009030	1.949796	2.323320
		OLS	0.004470	0.008523	1.810318	2.525125
	H = 6 m	MOTAD	0.002778	0.007335	2.362544	1.609580
		OLS	0.003651	0.006943	2.381547	2.246184
2 (outside bark)	H = 12 m	MOTAD	0.001091	0.006457	4.004871	0.443287
		OLS	0.002026	0.006105	4.029127	1.144342
	H = Total	MOTAD	0.001924	0.006700	64.006751	3.893092
		OLS	0.002073	0.006444	58.779860	3.668973
	H = 3 m	MOTAD	-0.008451	0.009652	2.816026	-5.365199
		OLS	-0.008503	0.009826	3.371938	-5.839304
	H = 6 m	MOTAD	-0.005069	0.007896	3.374180	-3.160691
		OLS	-0.005233	0.008025	3.985348	-3.580717
	H = 12 m	MOTAD	-0.002526	0.006604	5.077676	-1.563215
		OLS	-0.002603	0.006660	5.734359	-1.841434
	H = Total	MOTAD	-0.000763	0.006375	51.228707	1.025899
		OLS	-0.000571	0.006432	54.654965	1.619614

inside bark; and for the top height diameters: M2 slightly overestimates inside bark while underestimating outside bark, and M1 underestimates both outside and inside.

Table 4 presents the score values for each diameter estimation model at different heights (3, 6, 12 m, and total height), for each statistic separately and for the final total score.

The statistics calculated for the volume estimation at different heights are shown in Tables 5 and 6.

The *OLS* and *MOTAD* methods fitted models 1 and 2 with similar precision. M1 showed consistently better results than M2 for inside bark volume estimation at any height. For outside bark, M1 resulted better only for heights 3 and 6 m. For volumes up to 12 m, both fitting models resulted similar, and for total tree outside bark volume M2 showed more precise.

Both polynomial functions fitted well the data along the main part of the tree except for the base

Table 4 - Ranking attributes (2=best; 1=worst) for diameter estimation according to some performance statistics: bias (*D*), standard deviation of difference (*SD*), sum of squared relative residuals (*SSRR*) and residue percentage (*RP*)

Model	Section Length	Fitting Method	Scores				Total score
			<i>D</i>	<i>SD</i>	<i>SSRR</i>	<i>RP</i>	
1 (outside bark)	H = 3 m	MOTAD	2	1	1	1	5
		OLS	2	1	1	2	6
	H = 6 m	MOTAD	2	1	1	2	6
		OLS	2	1	1	2	6
	H = 12 m	MOTAD	2	1	1	2	6
		OLS	2	1	1	2	6
	H = Total	MOTAD	2	1	2	2	7
		OLS	2	1	2	2	7
	H = 3 m	MOTAD	2	2	2	2	8
		OLS	2	2	2	2	8
	H = 6 m	MOTAD	2	2	2	2	8
		OLS	2	2	2	2	8
1 (inside bark)	H = 12 m	MOTAD	2	2	1	2	7
		OLS	2	2	2	2	8
	H = Total	MOTAD	2	2	1	1	6
		OLS	1	2	1	1	5
	H = 3 m	MOTAD	1	2	2	2	7
		OLS	1	2	2	1	6
	H = 6 m	MOTAD	1	2	2	1	6
		OLS	1	2	2	1	6
	H = 12 m	MOTAD	1	2	2	1	6
		OLS	1	2	2	1	6
	H = Total	MOTAD	1	2	1	1	5
		OLS	1	2	1	1	5
2 (outside bark)	H = 3 m	MOTAD	1	1	1	1	4
		OLS	1	1	1	1	4
	H = 6 m	MOTAD	1	1	1	1	4
		OLS	1	1	1	1	4
	H = 12 m	MOTAD	1	1	2	1	5
		OLS	1	1	1	1	4
	H = Total	MOTAD	1	1	2	2	6
		OLS	2	1	2	2	7
	H = 3 m	MOTAD	1	1	1	1	4
		OLS	1	1	1	1	4
	H = 6 m	MOTAD	1	1	1	1	4
		OLS	1	1	1	1	4
2 (inside bark)	H = 12 m	MOTAD	1	1	2	1	5
		OLS	1	1	1	1	4
	H = Total	MOTAD	1	1	2	2	6
		OLS	2	1	2	2	7

part of the trunk. Differences between the two fitting methods, Ordinary Least Squares (*OLS*) and Minimization of Total Absolute Deviations (*MOTAD*), were practically insignificant in terms of generating good estimates for the taper function coefficients. Weights to reduce heteroscedasticity were not considered in this study, and are strongly recommended on further analysis of the two fitting methods in the future. Although rarely used on practical forest assessments, the diameter at the base of the tree (*DAB*) was here considered to cre-

ate the opportunity to evaluate the sensitivity of the *MOTAD* fitting method to situations where the variance highly increases when measurement values also increase.

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Table 5 - Bias (*D*), standard deviation of difference (*SD*), sum of squared relative residuals (*SSRR*) and residues percentage (*RP*) for volume estimation at different heights.

Model	Section length	Fitting Method	Volume			
			<i>D</i>	<i>SD</i>	<i>SSRR</i>	<i>RP</i>
1 (outside bark)	H = 3 m	MOTAD	0.011071	0.005700	4.681501	14.778097
		OLS	0.011241	0.005715	4.796915	15.001054
	H = 6 m	MOTAD	0.007332	0.009824	1.923516	6.495603
		OLS	0.007765	0.009793	1.999113	6.846832
	H = 12 m	MOTAD	0.002309	0.016823	1.407697	1.538785
		OLS	0.002850	0.016726	1.413479	1.806309
	H = Total	MOTAD	-0.008639	0.024495	1.764325	-3.041795
		OLS	-0.008695	0.024508	1.767484	-3.064399
1 (inside bark)	H = 3 m	MOTAD	0.012323	0.004814	6.984797	19.108403
		OLS	0.012304	0.004810	6.964637	19.078205
	H = 6 m	MOTAD	0.009527	0.007298	2.319237	9.246342
		OLS	0.009526	0.007295	2.320073	9.248438
	H = 12 m	MOTAD	0.003743	0.013398	1.239464	2.172295
		OLS	0.004040	0.013376	1.249489	2.363919
	H = Total	MOTAD	-0.005009	0.019804	1.567816	-2.502846
		OLS	-0.004120	0.019689	1.520680	-2.073619
2 (outside bark)	H = 3 m	MOTAD	0.018288	0.006418	10.046917	23.659334
		OLS	0.018332	0.006434	10.093778	23.715020
	H = 6 m	MOTAD	0.015478	0.006518	2.988615	12.601863
		OLS	0.016593	0.006758	3.407987	13.510578
	H = 12 m	MOTAD	0.008502	0.009874	0.648043	4.329850
		OLS	0.011107	0.010091	0.876826	5.656509
	H = Total	MOTAD	-0.000463	0.015459	0.501039	-0.195116
		OLS	0.001325	0.015306	0.497988	0.532694
2 (inside bark)	H = 3 m	MOTAD	0.018529	0.006490	10.311861	23.972620
		OLS	0.018523	0.006487	10.305559	23.965225
	H = 6m	MOTAD	0.020273	0.007536	5.024223	16.546106
		OLS	0.020064	0.007482	4.925819	16.377198
	H = 12m	MOTAD	0.024571	0.011958	3.151408	12.741846
		OLS	0.024289	0.011861	3.091489	12.607593
	H = Total	MOTAD	0.023703	0.016672	2.085584	9.714352
		OLS	0.024065	0.016733	2.136633	9.864727

Table 6 - Ranking attributes (2=best; 1=worst) for volume estimation according to some performance statistics: bias (*D*), standard deviation of difference (*SD*), sum of squared relative residuals (*SSRR*) and residue percentage (*RP*).

Model	Section length	Fitting Method	Scores				Total score	
			<i>D</i>	<i>SD</i>	<i>SSRR</i>	<i>RP</i>		
1 (outside bark)	H = 3 m	MOTAD	2	2	2	2	8	
		OLS	2	2	2	2	8	
	H = 6 m	MOTAD	2	1	2	2	7	
		OLS	2	1	2	2	7	
	H = 12 m	MOTAD	2	1	1	2	6	
		OLS	2	1	1	2	6	
	H = Total	MOTAD	1	1	1	1	4	
		OLS	1	1	1	1	4	
	1 (inside bark)	H = 3 m	MOTAD	2	2	2	2	8
			OLS	2	2	2	2	8
		H = 6 m	MOTAD	2	2	2	2	8
			OLS	2	2	2	2	8
		H = 12 m	MOTAD	2	1	2	2	7
			OLS	2	1	2	2	7
		H = Total	MOTAD	2	1	2	2	7
			OLS	2	1	2	2	7
2 (outside bark)	H = 3 m	MOTAD	1	1	1	1	4	
		OLS	1	1	1	1	4	
		H = 6 m	MOTAD	1	2	1	1	5
			OLS	1	2	1	1	5
		H = 12 m	MOTAD	1	2	2	1	6
			OLS	1	2	2	1	6
		H = Total	MOTAD	2	2	2	2	8
			OLS	2	2	2	2	8
	H = 6 m	MOTAD	1	1	1	1	4	
		OLS	1	1	1	1	4	
		H = 12 m	MOTAD	1	1	1	1	4
			OLS	1	1	1	1	4
		H = Total	MOTAD	1	2	1	1	5
			OLS	1	2	1	1	5

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