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Assessing Allometric Models to Predict Vegetative Growth of Yams in Different Environments

Denis Cornet,* Jorge Sierra, and Régis Tournebize

ABSTRACT

Yams are a neglected crop, grown mostly in West Africa by resource-poor farmers. Little is known about the physiology of the crop, and researchers lack practical and efficient tools to conduct growth analysis. The objective of this study was to develop allometric models able to predict yam leaf area and leaf and stem dry mass with acceptable accuracy. The models were calibrated using a data set comprising 10 cultivars belonging to the two main species (Dioscorea alata L. and D. rotundata Poir.) grown at two locations in Guadeloupe (Lesser Antilles) and two locations in Benin (West Africa). The best models were selected based on Akaike’s information criteria and validated against independent data sets. A power regression was best for predicting leaf area from leaf measurements while linear relationships were sufficient to predict the relationship between crop leaf area and leaf and stem mass. The use of species-specific models for the estimation of leaf and stem mass significantly improved the models’ performance. Models predicting yam leaf area and leaf mass proved to be reliable and accurate (no significant deviation and adjusted $R^2 > 0.95$). For stem mass, overestimation always occurred during validation (9%). To overcome this discrepancy, a methodology was proposed that allows the user to calibrate the model by tailoring the sampling size to obtain the required precision. The use of the selected models provides a nondestructive and reliable alternative to estimate leaf area and leaf and stem biomass for different cultivars and sites.

Yams are grown throughout the tropical parts of the world and are important staple crops in at least 10 countries (ranging from Nigeria to Jamaica and the Solomon Islands), accounting for 155 million consumers (FAO, 2012). Despite their economic value, research on yam is still scarce (Varshney et al., 2012), and there is little literature available on the crop’s physiology (Onwueme and Haverkort, 1991; Marcos et al., 2011). To address this lack of knowledge, there is a clear need for practical methods allowing basic growth analysis. An essential aspect of crop growth analysis lies in the accurate and nondestructive estimation of key variables such as leaf area or crop biomass (Normand and Lauri, 2012). To perform such estimations, modern equipment (e.g., hand scanners or laser optic apparatus) may be expensive and too complex for basic and simple studies. Moreover, 85% of world yam production is in rural areas of West Africa, where modern facilities and instruments may be unavailable, yet simple practical tools and methods may suffice for researchers. The use of predictive allometric models allows this difficulty to be circumvented by quantifying statistical relationships between key variables and some easily measured predictors (Niklas, 2004).

Estimation of leaf area is a key part of classical growth analysis and is necessary in many plant physiological studies (Sestak et al., 1971). Models for the nondestructive estimation of leaf area are useful tools for researchers in agronomic experiments. The linear measurement method allows the leaf area to be estimated without disturbing the growth of the plant (Sestak et al., 1971). This simple, nondestructive, and accurate method is based on linear regressions of leaf area on various leaf measurements. For heart-shaped leaves, the most frequently used measurements are leaf maximum length, leaf maximum width, midrib length, or their combination (Sinoquet and Andrieu, 1993). Although simple methods exist for estimating the leaf area of some common crops (Kandiannan et al., 2002; Lu et al., 2004; Pompelli et al., 2012), only fragmentary studies are reported for yam (Gumbs and Ferguson, 1976; Ravi and Chowdhury, 1989; George, 1992). Unfortunately, these published reports on yam fail to show that any one specific approach for deriving leaf area is adequate across cultivars and environmental conditions. Moreover, sometimes the adequacy of the model’s assumptions for estimating leaf area has not been carefully examined (e.g., there may be no validation step).

Predictive models for estimating key variables at the crop level have been developed for agronomic or ecological purposes (Müller et al., 2000; Shipley and Meziane, 2002; McCarthy and Enquist, 2007). Some relate the leaf area to the leaf biomass through the specific leaf area, while others investigate...
allometric biomass partitioning. These classical allometric models differ from the models at the level of the leaf because they integrate all the growth processes of the plant (Normand and Lauri, 2012) and are thus often less accurate.

The objective of this study was to develop predictive models to estimate leaf area and leaf and stem mass of 10 cultivars belonging to the two main yam species (D. alata and D. rotundata). This study focused on two levels: the individual leaf and the whole crop. At the level of the individual leaf, we evaluated the currently used models and propose a reliable and accurate model using nondestructive linear measurements for the estimation of the leaf area of yam. At the crop level, we link the stem and leaf mass to the leaf area of the crop.

**MATERIALS AND METHODS**

**Data Collection**

Ten data sets from field experiments with a wide range of cultivars and cultural practices (Table 1) were selected. The experiments were performed in Guadeloupe (Lesser Antilles) at the INRA experimental stations of Duclos (16°12′ N, 61°39′ W, 250 m asl) and Godet (16°24′ N, 61°29′ W, 30 m asl) and in Benin (West Africa) at the IITA experimental stations of Cotonou (6°25′ N, 2°19′ E, 23 m asl) and Glazoué (7°56′ N, 2°15′ E, 200 m asl). Each field experiment consisted of 300 to 550 plants.

At the crop level, plants were sampled from an area of 1.5 to 3 m² every 2 to 3 wk from vine emergence to the beginning of tuber growth (approximately 110 d) to measure leaf and stem biomass and crop leaf area index. There were four replicates of one to four plants at each sampling date, depending on plant density. Crop leaf area was measured with a planimeter (Li-Cor Model 3100) and expressed as the area of leaves (m²) per square meter of ground. Leaf biomass (LM in g m⁻²) and stem biomass (SM in g m⁻²) were determined after drying the material at 70°C. To study the allometric relationships between crop leaf area and leaf or stem dry mass, the field experiments were divided into two groups: Exp. 1 to 6 (520 samples) were used to select models and estimate their parameters, while Exp. 7 to 10 were used to test the chosen models (259 samples). All cultivars belonged to the two main species (i.e., Florido, Belep, Oriental, Kinabay, Boutou, and Plimbite for D. alata and Kpouna, Morokorou, Gnidou, and Kokoro for D. rotundata).

At the individual leaf level, samples were collected in 2001, 2004, and 2007 (Table 1) to study the relationship between linear measurements of leaf blade and leaf area. All samples consisted of at least 40 leaves collected randomly from different positions on the vine. All cultivars had heart-shaped leaves on which maximum length (L in cm), maximum width (W in cm), and midrib length (M in cm) were measured. Afterward, individual leaf area (A in cm²) was measured with a planimeter. Samples from 2001 and 2007 were used for model calibration and investigation of the effects of site and cultivar (456 samples), while samples from 2004 were used for model validation (354 samples).

**Statistical Analysis**

Statistical analysis was performed in four steps. First, all the models were evaluated and the best equations for each of the three allometric relationships selected. Second, the effects of the covariates (cultivar, species, site, and year) on the allometric models were tested, and the selected models were validated. Third, the calibrated models were validated against an independent data set and compared with other published models. Finally, if a significant deviation in the model estimates was found during the validation step, the minimum sample size for the parameterization of such a model was then evaluated.

**Table 1. Experiment characteristics of each data set used in this study.**

<table>
<thead>
<tr>
<th>Data set</th>
<th>Year</th>
<th>Site</th>
<th>Cultivars</th>
<th>Plant density</th>
<th>Irrigation</th>
<th>Mineral fertilization</th>
<th>Soil type†</th>
<th>Mean temp.</th>
<th>Cumulated rainfall</th>
<th>References</th>
</tr>
</thead>
<tbody>
<tr>
<td>Calibration data set</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>1††</td>
<td>2001</td>
<td>Duclos</td>
<td>Belep, Kinabay, Oriental, Boutou</td>
<td>1.56</td>
<td>yes</td>
<td>N 110 P 80 K 150</td>
<td>ferralsol</td>
<td>25.8</td>
<td>4329</td>
<td>Patetsos (2001)</td>
</tr>
<tr>
<td>3</td>
<td>2006</td>
<td>Cotonou</td>
<td>Florido, Gnidou</td>
<td>0.11</td>
<td>no</td>
<td>N 80 P 13 K 132</td>
<td>arenosols</td>
<td>27.0</td>
<td>1060</td>
<td>Cornet et al. (2014)</td>
</tr>
<tr>
<td>4</td>
<td>2007</td>
<td>Cotonou</td>
<td>Florido, Morokorou, Kokoro</td>
<td>0.67</td>
<td>yes</td>
<td>N 80 P 13 K 132</td>
<td>arenosols</td>
<td>26.6</td>
<td>1395</td>
<td>Cornet et al. (2014)</td>
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<tr>
<td>5</td>
<td>2007</td>
<td>Glazoué</td>
<td>Florido, Kokoro, Kpouna</td>
<td>0.67</td>
<td>no</td>
<td>N 80 P 13 K 132</td>
<td>ferralsol</td>
<td>28.5</td>
<td>1010</td>
<td>unpublished data (2014)</td>
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<tr>
<td>6</td>
<td>2008</td>
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<td>Florido, Morokorou</td>
<td>0.67</td>
<td>yes</td>
<td>N 60 P 6 K 66</td>
<td>arenosols</td>
<td>27.0</td>
<td>1562</td>
<td>Cornet et al. (2014)</td>
</tr>
<tr>
<td>Validation data set</td>
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<td></td>
<td></td>
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<td></td>
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</tr>
<tr>
<td>7</td>
<td>2009</td>
<td>Cotonou</td>
<td>Florido, Morokorou</td>
<td>0.67</td>
<td>yes</td>
<td>N 120 P 80 K 120</td>
<td>arenosols</td>
<td>27.3</td>
<td>1427</td>
<td>Cornet et al. (2014)</td>
</tr>
<tr>
<td>8</td>
<td>2009</td>
<td>Glazoué</td>
<td>Florido, Morokorou</td>
<td>0.67</td>
<td>no</td>
<td>N 80 P 4 K 44</td>
<td>ferralsol</td>
<td>28.4</td>
<td>995</td>
<td>unpublished data (2014)</td>
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<td>10</td>
<td>2004</td>
<td>Duclos</td>
<td>Belep, Kinabay, Oriental, Plimbite, Boutou</td>
<td>3.33</td>
<td>yes</td>
<td>N 200 P 100 K 200</td>
<td>ferralsol</td>
<td>25.4</td>
<td>3664</td>
<td>Cornet (2005)</td>
</tr>
</tbody>
</table>

† FAO (1998).
†† Experiments in 2001 were used only for the calibration of leaf area estimation models.
All statistical analyses were performed using R software (R Core Team, 2013).

**Model Selection**

For each allometric relationship, several linear and nonlinear regression models were run using combinations of predictors and response variables. These were either chosen from the literature or derived for the present study based on their logical mathematical relation to the published models. Preliminary regression analysis showed that most of the relationships between crop leaf area and leaf and stem dry biomass followed linear patterns, while the relationship between linear measurements and leaf area was better modeled with power functions typical of allometric models.

Hence, in this study, only linear models are presented for the estimation of leaf and stem mass, while only power models are presented for leaf area. A logarithmic transformation was used to make the assumption of homogeneous variances reasonable in the power models (e.g., $y = ax^b$):

$$\ln y = b \ln x + \ln a$$

where $y$ is the response variable, $x$ the predictor, and $a$ and $b$ the model parameters.

Models were selected using Akaike’s information criterion (AIC). The AIC is a common model selection criterion that deals with both the goodness of fit and the complexity of the model (i.e., number of parameters) and enables several models to be compared simultaneously (Johnson and Omland, 2004). The lower the value of the AIC, the better the model. Because the AIC tells nothing about the quality of the model in an absolute sense, the adjusted coefficient of determination ($R^2$) and the mean standard error percentage ($\%$) were also calculated to illustrate the mean size of the error as a percentage of the estimated values:

$$\% = 100 \frac{\sum_{i=1}^{n} |\hat{y}_i - y_i|}{\sum_{i=1}^{n} \hat{y}_i}$$

where $y_i$ is the actual value, $\hat{y}_i$ is the estimated value, and $n$ is the number of observations (Schafer, 1980). Because the model estimates were computed on a logarithmic scale, they must then be back-transformed to their original, arithmetic scale. Because this transformation is nonlinear, and there is variability in the observed data around the fitted relationship, a simple “naive” exponential-based transformation will generate bias (Clifford et al., 2013). Consequently, correction factors are typically calculated to remove this bias when back-transforming. As proposed by Clifford et al. (2013), we calculated the uniform minimum variance unbiased estimations described by Bradu and Mundlak (1970). The R codes required to fit a log-log allometric model and to make predictions using the uniform minimum variance unbiased estimations were adapted from Clifford et al. (2013).

**Testing Covariates and Model Calibration**

Experimental factors (i.e., cultivars, sites, and year) can affect the slope and/or the intercept of the relationships and thus were studied as covariates. To test the influence of these covariates on the allometric relationships, a maximal model was fitted (i.e., a model including explanatory variables, covariates, and first-order interactions). Furthermore the maximal model was simplified according to Crawley’s (2007) procedure. The model summaries were inspected and the least significant term was removed. The full and simplified models were then compared using the F-test. If no significant difference was observed, the simplified version was kept and the least significant highest order term was again deleted. If the F-test revealed a significant difference ($P < 0.05$), the previous model was kept. Finally, a 1000 bootstrap resampling in conjunction with automated variable selection methods was applied to check the efficiency of the selected models (Austin and Tu, 2004). Variables that were identified as independent predictors of the response variable in at least 60% of the bootstrap samples were kept, and this last model was called the minimal adequate model. The relative importance of each term of the minimal adequate model was estimated using the Lindeman, Merenda, and Gold metric (Lindeman et al., 1980), allowing an estimation of the percentage of the variance explained by each term. The confidence intervals for relative importance were estimated using a bootstrap procedure on 1000 samples (Grömping, 2006). Differences in model performance (i.e., $\Delta R^2$, $\Delta AIC$, and $\Delta \%$) were calculated between the minimal adequate model and the classical pooled regression.

**Model Validation and Minimum Sample Size**

To validate the calibrated models, leaf area, leaf mass, and stem mass were estimated using the selected allometric models on independent data. The estimated values were compared with observed ones using $\%$, $R^2$, and the bias ($\bar{b}$%). The bias is defined as the scaled mean error (i.e., the mean difference between the estimated and the observed values expressed as a percentage of the mean observed value):

$$\bar{b} = \frac{100}{n} \sum_{i=1}^{n} \frac{\hat{y}_i - y_i}{y_i}$$

The bias indicates whether the model, on average, overestimates ($\bar{b} > 0$) or underestimates ($\bar{b} < 0$) data (Walther and Moore, 2005).

Paired t-tests were run to determine whether the differences between the estimated and the observed values were statistically significant at $P < 0.05$. When this difference was significant, we estimated the minimum sample size allowing user recalibration with minimum bias. In this case, parameter values and bias were calculated on randomly selected subsamples of increasing size. All sample sizes were submitted to a bootstrapping procedure with 100 resamples with replacement. The mean bias was then expressed as a function of the sample size.

**RESULTS AND DISCUSSION**

**Model Selection**

The domain of validity of the allometric models is defined by the experimental range for crop leaf area, leaf and stem mass, and individual leaf area used for model calibration. A summary of the data sets is given in Table 2. Table 3 summarizes the performance of candidate models during the calibration phase. Models were fitted to the calibration data set and ranked according to their performance. Individual leaf area was best explained by a power
function of the square of the maximum leaf length multiplied by the maximum leaf length ($S^2$% of 5.4% of mean leaf area and $R^2$ of 0.987; Table 3). Leaf and stem mass were better explained with a linear function of crop leaf area. The best allometric models for leaf and stem mass were not as good (i.e., 5% of 10.5 and 15.3% of 0.977 and 0.935, respectively; Table 3) as for leaf area similar to those reported in previous studies (Normand and Lauri, 2012; Pompelli et al., 2012; Kandianann et al., 2002). The higher $S^2$% and lower $R^2$% for leaf mass estimation may be due to the large variation in specific leaf area within a single stand, plant (Cooper and Qualls, 1967; Reddy et al., 1989), or even leaf (Sestak et al., 1971). Moreover the specific leaf area could have been influenced by the growth stage of the plant (Koller, 1972) and by the time of day (Chatterton et al., 1972) when samples were taken. Cornet et al. (2014) showed that yam emergence may occur over 1 mo, creating a very heterogeneous population of individuals of different ages with different specific leaf areas. This variability in specific leaf area may explain some of the scatter of leaf mass estimates with the evaluated model.

The leaf area/stem mass allometric model showed the highest $S^2$% and lowest $R^2$. This was also observed by other researchers and may be due to changes in both specific leaf area and leaf/stem ratio (Normand and Lauri, 2012). In fact, herbaceous self-supporting plants show an ontogenetic trend toward a decreasing leaf/stem allocation ratio of assimilates, with an increasingly larger investment in the support tissues needed by larger plants (Poorter et al., 2012). Because all yam species grown for food have a climbing growth habit, they are likely to behave differently. Unfortunately, literature on the functional anatomy of climbing species is scarce. Some studies have indicated that vines and lianas have a higher leaf/stem ratio but also higher plasticity (i.e., greater variation) in their anatomical characteristics in response to changing environmental conditions than their self-supporting relatives (Gianoli, 2003; den Dуббelden and Verburg, 1996; Werth et al., 2013). Thus, the increasing competition between plants with age can modify the leaf/stem ratio and therefore increase the variability of related allometric relationships.

### Year, Site, and Cultivar Effects on Allometric Models

Once the best model for each allometric relationship was selected, the relative importance of cultivar and environmental factors (year and site) was investigated. A summary of all terms of the minimally adequate model and their relative importance is presented in Table 4. Explanatory variables (i.e., linear measurements and crop leaf area for the estimation of leaf area or leaf and stem mass, respectively) always accounted for >83% of the variance explained by the model. Among covariates, cultivar was the most influential and accounted for 0, 4, and 3% of the explained variance for leaf area, leaf mass, and stem mass, respectively, while year accounted for 0, 4, and 3% of the explained variance for leaf area, leaf mass, and stem mass, respectively.
Despite the quite large fraction of the variance explained by the individual leaf area or stem biomass model terms, the effects of year, site, and their interaction had negligible impact on the performance of the models.

Thus, plants bearing cataphylls produce significant stem mass before developing any leaf area and thus could modify the estimation of the stem mass of D. rotundata. Only in the estimation of the stem mass of D. alata was the intercept significantly different from zero. Differences between species may be due to the heteroblastic growth of D. rotundata. Indeed, leaves at the first few nodes of this species may be reduced to form modified shield-shaped cataphylls (Onwueme, 1978). These cataphylls are thick, lack a distinct leaf lamina, and are limited in aerial spread. Cornet (unpublished data, 2014) observed that a D. rotundata plant can bear up to 16 nodes carrying cataphylls before the appearance of the first true leaf, while cataphyll occurrence in D. alata was scarce (<2%). Thus, plants bearing cataphylls produce significant stem mass before developing any leaf area and thus could modify the intercept of the leaf area–stem mass relationship.

### Model Validation

Final models were validated against a completely independent data set (Table 4). The mean standard error percentage of the intercept of the leaf area–stem mass relationship was best described by a power model calibrated on pooled data. Conversely, leaf and stem mass estimations were best fitted with species-specific linear models. Only in the estimation of the stem mass of D. rotundata was the intercept significantly different from zero. Differences between species may be due to the heteroblastic growth of D. rotundata.
always above 0.93 and, except for stem mass ($r = 0.93\%$), the bias was negligible (–0.8 and 1.2 for leaf mass and leaf area estimation, respectively). Comparing the leaf area model obtained in this study with available models from the literature confirmed its greater accuracy (Fig. 1). Indeed, the model developed in this study always outperformed models from the literature. A paired Student’s t-test did not reveal any differences between the estimated and observed leaf areas using our model, but significant differences were observed when using the published models.

Models to estimate yam leaf and stem mass could not be found in the available literature. No differences between estimated and observed leaf mass were observed using our model. Moreover, the use of species-specific models decreased the model bias for both leaf and stem mass estimation (Fig. 1). However, the estimation of stem mass was very biased. The diagnostic plot shows clear differences between the stem mass model and the leaf area model (Fig. 2). Estimated and observed values were closely correlated for both leaf area and stem mass models (0.99 and 0.95, respectively), but the slope of the regression line was significantly different from 1 for stem mass but not for leaf area. This confirmed the formerly observed bias in stem mass estimation and showed that this bias was greater for D. rotundata than D. alata (slopes of 1.09 vs. 1.05, respectively). Residuals of the stem mass model present a biased and heteroscedastic pattern, with a clear overestimation for stem mass >30 g. The potential number of cataphylls per stem depends on the seed-tuber physiological age and weight (unpublished data, 2014). Because of differences in agricultural practices, the mean physiological age and weight of the seed tuber varied among trials. This could have led to the presence of varying numbers of cataphylls per stem, which may explain the large differences in the leaf/stem ratio among trials. To overcome this lack of reliability, we analyzed the relationship between the number of samples and the performance of the stem mass model. The variation in the estimate of the model parameter $b$ tended to decline as the sampling size increased (Fig. 3A). The inferred model bias decreased sharply with an increased number of samples as the sampling size increased (Fig. 3A). The inferred model bias decreased sharply with an increased number of samples as the sampling size increased (Fig. 3A).

![Fig. 1: Statistical analysis (Student's t-test for paired data) of the deviation between the estimated and observed leaf area, leaf mass, and stem mass. Estimates were based on the final best models (pooled or species-specific) of this study and models from the literature where available. Vertical bars denote the bias, and spreads denote 95% confidence intervals; ns = nonsignificant, *significant at $P < 0.05$, **significant at $P < 0.01$, ***significant at $P < 0.001$.](image)

![Fig. 2: Diagnostic plots of allometric models to predict (A) leaf area and (B) stem mass for the two main yam species (D. alata and D. rotundata). (A) Main plot presents the regression of estimated vs. actual values (black line and equation) of the pooled allometric model for leaf area estimation from linear measurement, while the inset presents the residuals of the model compared with the data estimated by the model. (B) Main plot presents the regression of estimated vs. actual values (lines and equations) of the species-specific allometric model for stem mass estimation from crop leaf area, while the inset presents the residuals of the model compared with the data estimated by the model.](image)
represent practical, cheap, and reliable alternatives for an accurate and nondestructive estimation of key variables of yam vegetative growth.

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