

Contextualizing Pan-Tropical Allometric Models for Biomass Estimation

Eustache Diemert, Anaëlle Dambreville

PUR



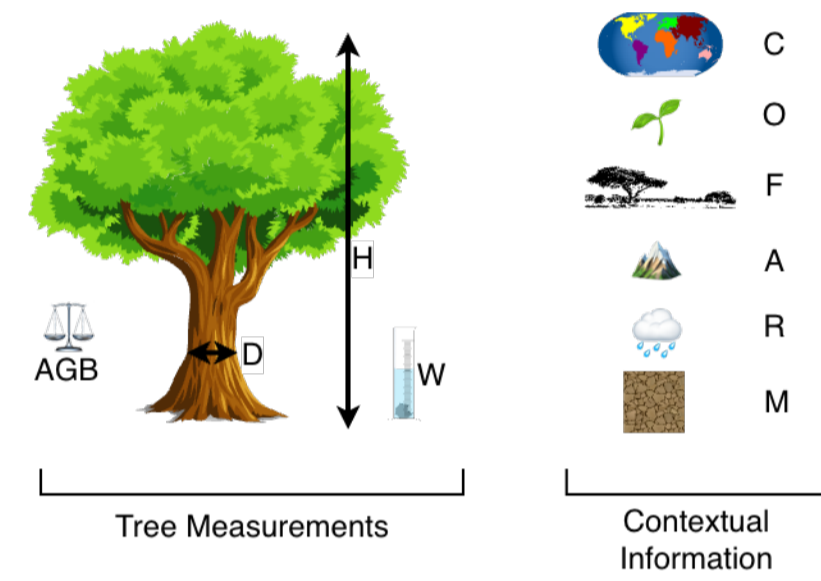
Introduction

Allometric Models (AMs) play a central role in monitoring and mitigating climate change as they provide accurate estimation of biomass and carbon sequestered by trees from non-destructive, easy to obtain physical measurements.

Unfortunately, practitioners spend considerable effort in researching, qualifying and choosing AMs for specific growth conditions.

Classical Pan-Tropical Allometric Models

The dataset from (author?) [1] contains measurements of trunk diameter at breast height D (cm), total tree height H (m), wood density W ($g.cm^{-3}$), and total oven-dry above ground dry matter AGB (kg) for 4004 trees across 58 sites worldwide.



Tree Measurements	
D	diameter at breast height (cm)
H	total height (m)
W	wood density ($g.cm^{-3}$)
Contextual Information	
C	continent (categorical)
O	old growth (binary)
F	forest type (categorical)
A	altitude (m)
R	annual rainfall (mm)
M	dry months (0 – 12)
Tree Biomass (prediction target)	
AGB	above ground dry matter (kg)

Figure 1. Tree measurements and contextual variables

The state of the art pan-tropical allometric model from [1]: $AGB = \alpha(D^2HW)^\beta$

Frontiers in Allometric Models Development

- Can allometric models be improved by incorporating richer contextual information?
- Can transfer error be predicted when using an allometric model in a new site?

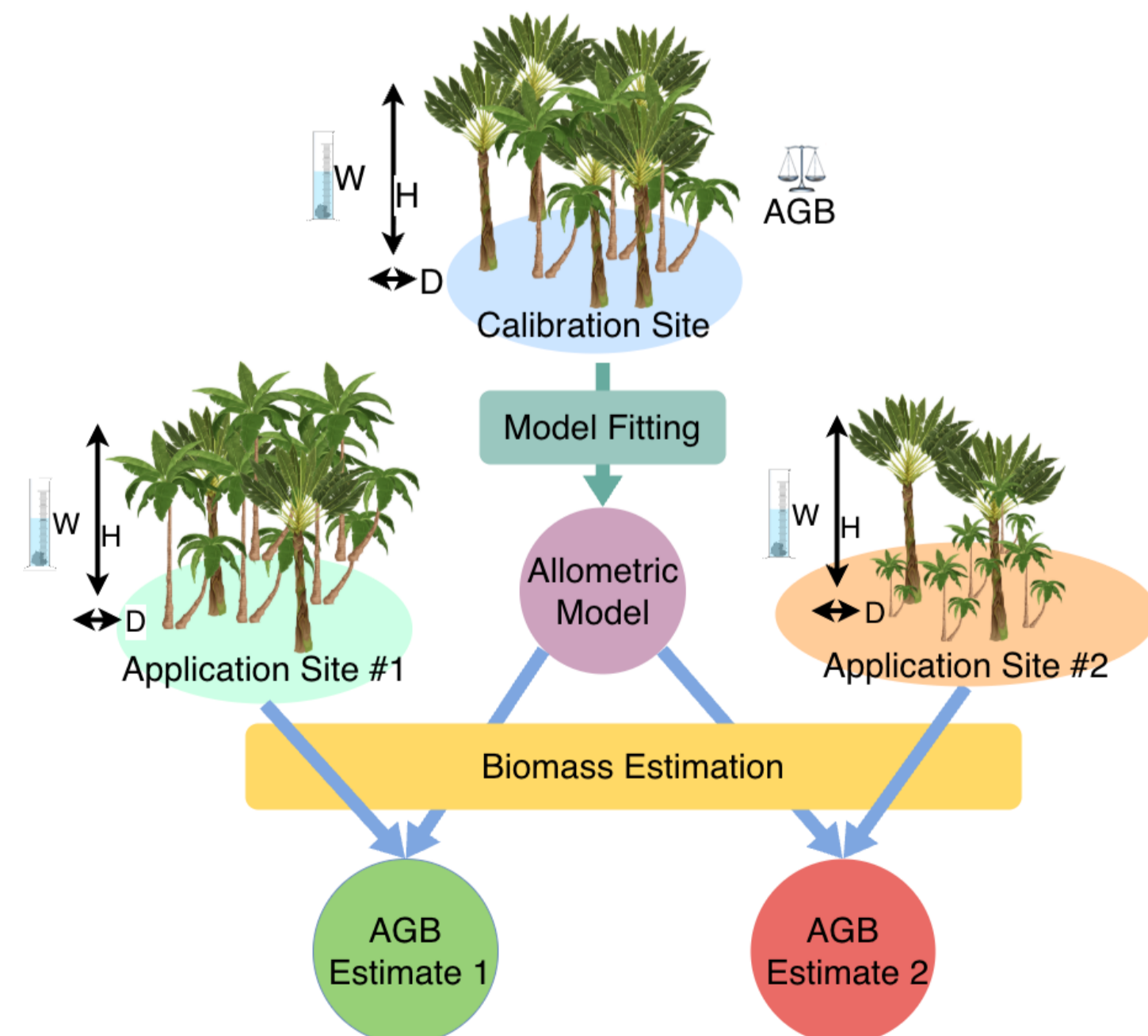


Figure 2. Illustration of how data distribution shift between Calibration site and two Application sites may affect estimated Biomass: a severe shift at site 2 jeopardizes estimated AGB Accuracy while a mild shift doesn't affect site 1 accuracy.

Contribution #1: Deep Allometric Model

First, [1] can be trained as $\log AGB = \alpha \log D + \beta \log H + \gamma \log W$ with L^2 regularization. Then, assuming contextual information can provide valuable insights to the model, we add *ContextualChave*:

$$AGB = \alpha(D^2HW)^{\beta_0} \tilde{C}^{\beta_1} \tilde{O}^{\beta_2} \tilde{F}^{\beta_3} A^{\beta_4} R^{\beta_5} M^{\beta_6} \quad (1)$$

where $\tilde{\cdot}$ designates Target Encoding, a popular and efficient method to encode categorical variables.

Finally, we propose deep learning models:

- COFARM** a model that embeds categorical covariates C, O, F in 2 dimensions before pooling them with D, H, W, A, R, M in a log-log linear regressor
- COFARM-NN**, inspired by wide and deep architectures pools embedded categorical covariates C, O, F with A, R, M numerical features and passes resulting information through 3 layers of 3 fully connected neurons and ReLU blocks before the final regressor, as depicted on Figure 3.

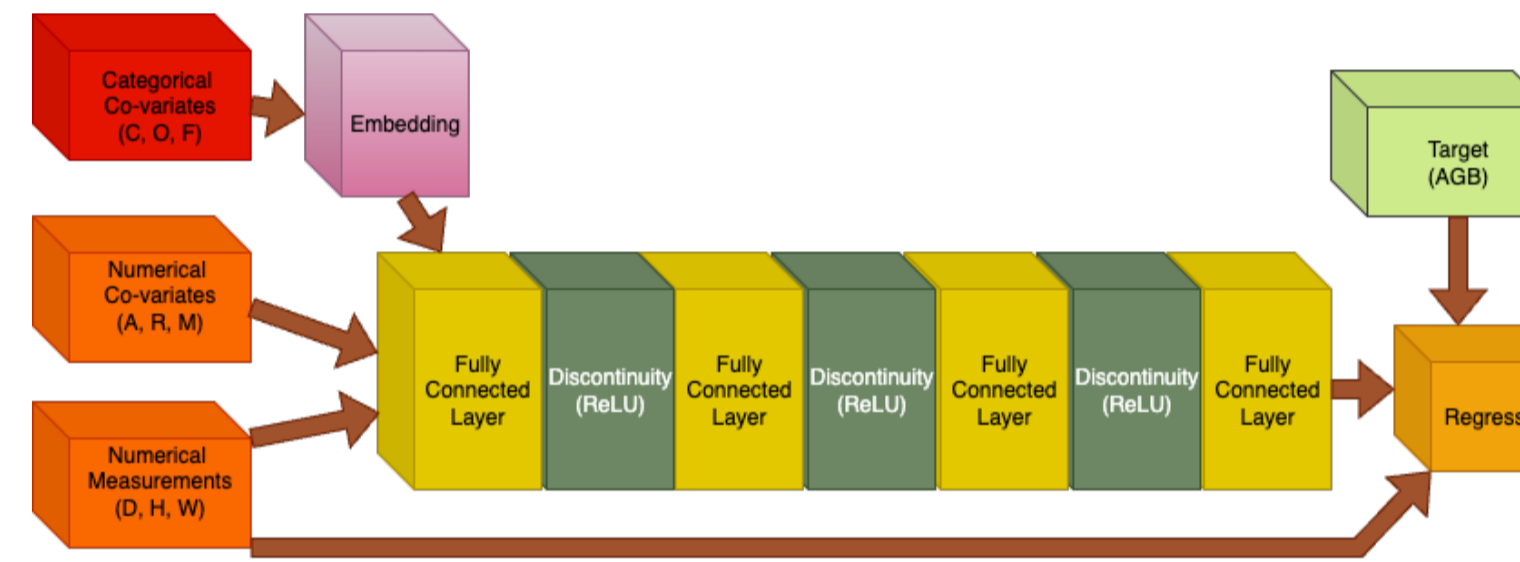


Figure 3. COFARM-NN model architecture for AGB prediction

Contribution #2: Transfer Error Model

We rely on [2] that establish a theoretical upper bound on the expected target error $\epsilon_t(f)$:

$$\epsilon_t(h) \leq \epsilon_s(h) + \eta_H(f_s, f_t) + \mathcal{H}Disc_{\mathcal{H},L}(p_t, p_s, h) \quad (2)$$

where $\eta_H(f_s, f_t)$ is the sum of the errors made by the ideal predictor on both source and target domains and $\mathcal{H}Disc_{\mathcal{H},L}(p_t, p_s, h)$ is the hypothesis-discrepancy associated with h and defined as

$$\mathcal{H}Disc_{\mathcal{H},L}(p_t, p_s, h) = \max_{h' \in \mathcal{H}} |\mathbb{E}_{x \sim p_t}[L(h'(x), h(x))] - \mathbb{E}_{x \sim p_s}[L(h'(x), h(x))]| \quad (3)$$

While η is purely related to the labeling discrepancy, $\mathcal{H}Disc$ measures the discrepancy between source and target distributions, relatively to the model h .

We propose a quadratic model of the additional prediction error:

$$\Delta_{s,t}MAE(h) = \alpha \times \mathcal{H}Disc_{\mathcal{H},L}(p_t, p_s, h) + \beta \times \text{sign}(\Delta_{s,t}Z) + \gamma \times \mathcal{H}Disc_{\mathcal{H},L}(p_t, p_s, h)^2 + \delta \times (\text{sign}(\Delta_{s,t}Z) \times \mathcal{H}Disc_{\mathcal{H},L}(p_t, p_s, h)) \quad (4)$$

where $\Delta_{s,t}Z = \mathbb{E}_{X \sim p_t}[D^2HW] - \mathbb{E}_{X \sim p_s}[D^2HW]$. The model is learned as a Ridge regression with L^2 penalty strength optimized by internal cross-validation.

To estimate $\mathcal{H}Disc$ in practice we solve the following constrained optimization problem

$$\max_{\alpha', \beta'} |(\hat{\epsilon}_s(f_{\alpha, \beta}) - \hat{\epsilon}_s(f_{\alpha', \beta'})) - (\hat{\epsilon}_t(f_{\alpha, \beta}) - \hat{\epsilon}_t(f_{\alpha', \beta'}))| \text{ s.t. } |\alpha' - \alpha| < \delta_1 \text{ and } |\beta' - \beta| < \delta_2 \quad (5)$$

, where $\hat{\epsilon}$ is the empirical MAE.

Experiment: Deep Allometric Models Performance

We benchmark a wide range of models to understand what can lead to reduced prediction errors. Baseline: [1] and simple log-log regression (*LogReg*); More complex: MLP (*LogReg-NN*); More features: Perceptron + Embeddings (*COFARM*); More complex and more features: tree based (*HGBRT*), MLP + Embeddings (*COFARM-NN*).

Table 1. Predictive power of different AMs (bold = best average, * = stat. sig. at 90% vs [1])

Dataset	Metric	[1]	ContextualChave	HGBRT	LogReg	LogReg-NN	COFARM	COFARM-NN
train	RMSE	829.9 ± 23.3	822.8 ± 21.2	628.8 ± 24.8	762.3 ± 18.9	762.3 ± 18.9	692.2 ± 20.2	590.0 ± 35.2
train	MAE	175.0 ± 2.9	163.4 ± 2.3	83.2 ± 2.4	168.8 ± 2.6	168.8 ± 2.6	147.7 ± 2.3	130.6 ± 3.8
train	R^2	0.888 ± 0.007	0.890 ± 0.006	0.936 ± 0.004	0.905 ± 0.006	0.905 ± 0.006	0.922 ± 0.005	0.942 ± 0.007
test	RMSE	804.9 ± 80.7	797.6 ± 78.9	785.4 ± 111.1	747.3 ± 69.8	747.2 ± 69.8	694.6 ± 71.7	702.9 ± 91.7
test	MAE	180.4 ± 11.1	168.0 ± 9.3	170.5 ± 13.7	173.6 ± 10.2	173.6 ± 10.2	155.4 ± 9.2* [-14%]	148.9 ± 9.4* [-17%]
test	R^2	0.859 ± 0.025	0.866 ± 0.027	0.813 ± 0.062	0.865 ± 0.030	0.865 ± 0.030	0.880 ± 0.029	0.881 ± 0.026

Experiment: Predict Transfer Error

We experiment with synthetic covariate shifts by splitting the data according to explanatory variables. Then the model from [1] is learned on one split and tested on the other. Finally we compare the additional transfer error $\Delta_{s,t}MAE(h)$ as predicted by Eq. 4 vs as estimated from observed empirical error.

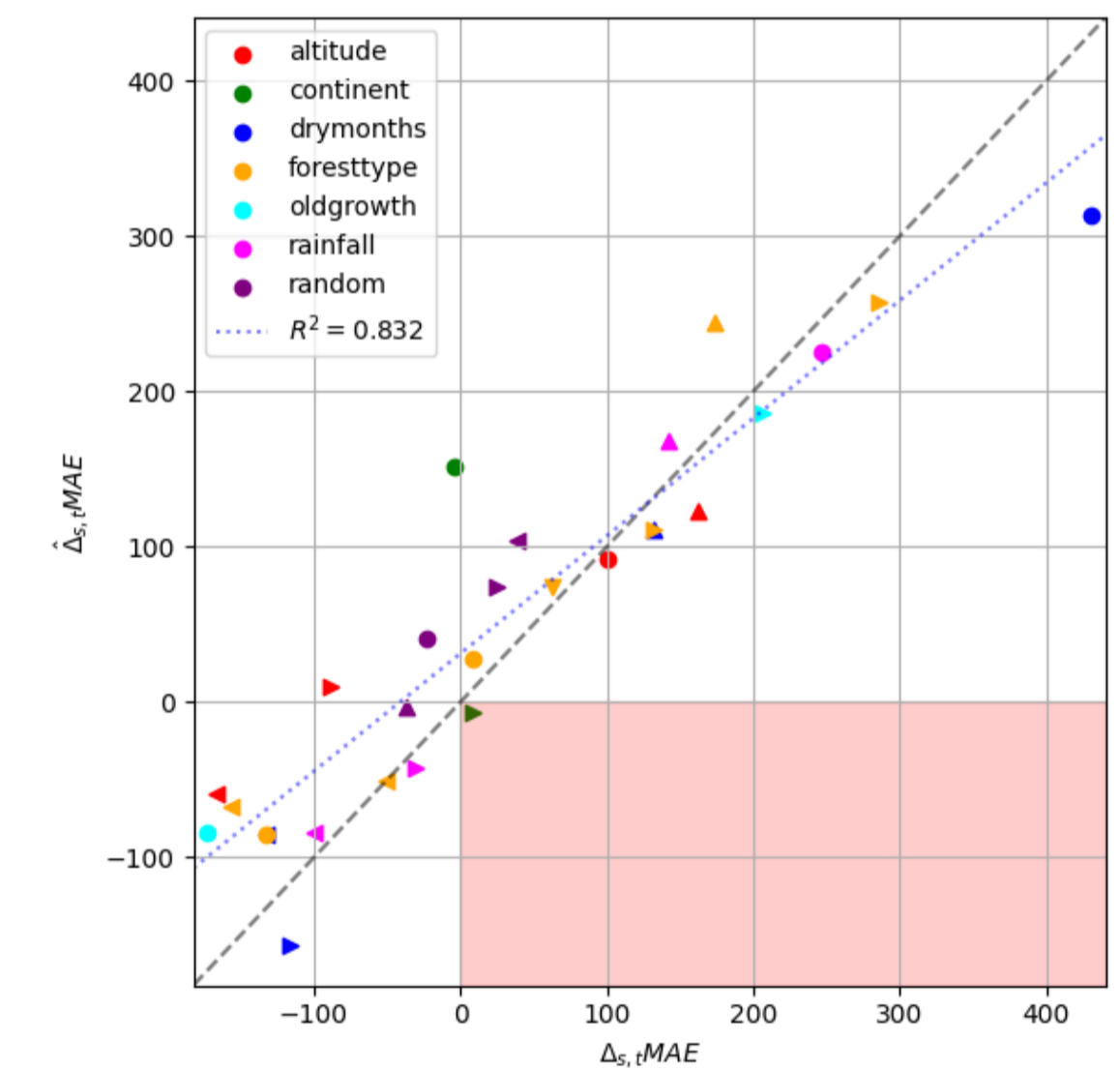


Figure 4. Predicted vs Actual Additional Error due to Covariate Shift. Color indicates the variable chosen to simulate the shift while marker shapes indicate different shifts magnitudes and directions.

Conclusion

- Embedding categorical contextual information leads to -14% MAE, while adding deep layers reduces further MAE by -3%
- Additional transfer error when applying a model to a new plot can be predicted ($R^2 = .832$) by quantifying the discrepancy between source and target features distribution

References

[1] J. Chave, M. Réjou-Méchain, A. Búrquez, E. Chidumayo, M. S. Colgan, W. B. Delitti, A. Duque, T. Eid, P. M. Fearnside, R. C. Goodman, M. Henry, A. Martínez-Yrizar, W. A. Mugasha, H. C. Muller-Landau, M. Mencuccini, B. W. Nelson, A. Ngomanda, E. M. Nogueira, E. Ortiz-Malavassi, R. Péliissier, P. Ploton, C. M. Ryan, J. G. Saldarriaga, and G. Vieilledent. Improved allometric models to estimate the aboveground biomass of tropical trees. *Global Change Biology*, 20(10):3177–3190, Oct. 2014.

[2] G. Richard, A. de Mathelin, G. Hébrail, M. Mougeot, and N. Vayatis. Unsupervised Multi-Source Domain Adaptation for Regression. In *Lecture Notes in Computer Science book series (LNCS, volume 12457)*, volume 12457 of *Lecture Notes in Computer Science book series*, pages 395–411, Ghent, Belgium, Sept. 2020. Springer International Publishing.