



Background

- Forest growth and yield tables/functions as a common practice to predict growth.
- No incorporation of genetic gain in conventional models → might not take into account outcomes from tree breeding programs (10 25 % gain in growth traits)?



Potential underestimations?

Objective → modifications of currently used growth functions for improved forest reproductive material (FRM) categories





Material and Methods

Based on NFI data, generalized algebraic difference approach (GADA) used for King-Prodan equation (Donis et al. 2018):

$$H_2 = 1.3 + \frac{A_2^{b_1}}{b_2 + 100 b_3 \frac{A_1^{b_1}}{H_1 - 1.3} - b_2} + \frac{A_1^{b_1}}{H_1 - 1.3} - b_2}{100b_3 + A_1^{b_1}} + \frac{A_1^{b_1}}{100b_3 + A_1^{b_1}} A_2^{b_1}$$

Species	b_1	b ₂	b_3
Scots pine	1.15697	-27.04027	16.4512
Norway spruce	1.28394	-47.34926	23.60081
Silver birch	1.257	-47.475	21.726

Donis et al. 2018



Material and Methods (2)

- Data from open pollinated progeny trials in Latvia
 - -10 % of best performing families «*Tested*»;
 - remaining 90 % «Qualified»
- FRM-cateogry-specific modifiers of GADA model coefficients estimated

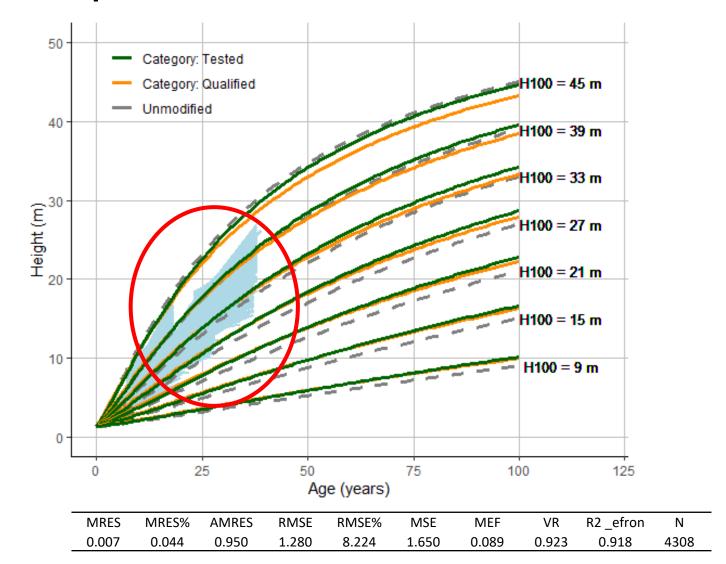
$$H_2 = 1.3 + \frac{A_2^{b_1}}{b_2 + 100 b_3 \frac{A_1^{b_1}}{H_1 - 1.3} - b_2} + \frac{A_1^{b_1}}{H_1 - 1.3} - b_2}{100b_3 + A_1^{b_1}} + \frac{A_1^{b_1}}{100b_3 + A_1^{b_1}} A_2^{b_1}$$



$$H_{2} = 1.3 + \frac{A_{2}^{g_{1} \cdot b_{1}}}{\frac{A_{1}^{b_{1}}}{H_{1} - 1.3} - b_{2}} + \frac{A_{1}^{b_{1}}}{\frac{H_{1} - 1.3}{100b_{3} + A_{1}^{b_{1}}}} + \frac{A_{1}^{b_{1}}}{100b_{3} + A_{1}^{b_{1}}} A_{2}^{g_{1} \cdot b_{1}}$$



Results: Scots pine

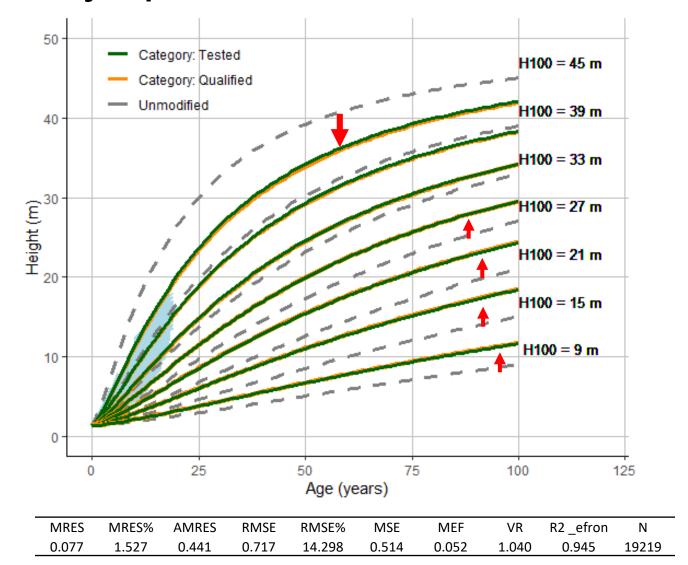






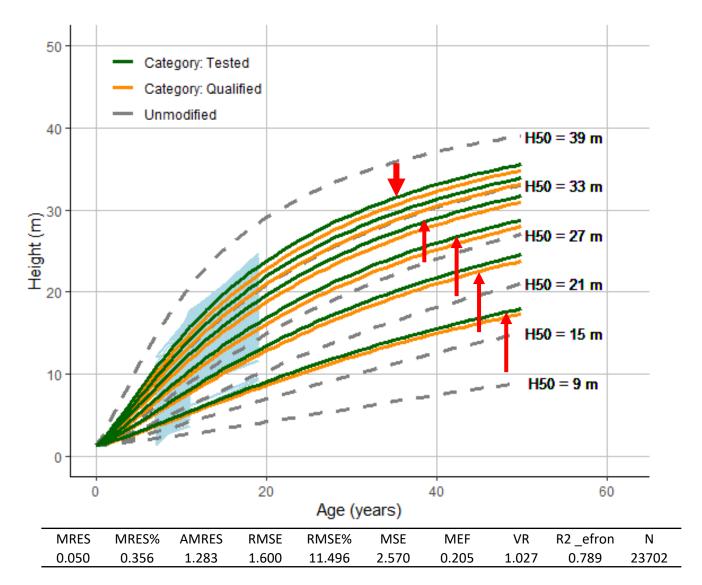


Results: Norway spruce





Results: Silver birch





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