

Characterizing vertical forest stands structure using data mining methods

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Vertical forest stands structure

DEFINITION:

“... the bottom to top configuration of above ground vegetation within a forest stand”

“distribution of tree heights within a forest stand”

Changes during natural development of stand:

- the number, height and biomass of trees,
- arrangement of branches
- leaf area on trees,
- understory composition and structure, etc.

Vertical forest stands structure

Influences working on stand:

- **natural forest development processes** (cyclical stand development, successions)
- **silviculture treatments** (part of forest management).

Factors that **affect** the development of vertical structure:

- physiological and morphological properties of individual trees,
- size of the trees
- spatial position of their neighbors,
- stand density,
- disturbance history,
- site conditions etc.

Vertical forest stands structure

Changes in vertical forest structure affect:

- both **microclimatic** factors and **processes** in the system,
- tree **growth**,
- **understory** community structure,
- suitability of the stand for **wildlife**,
- **hydrologic** response,
- fire **hazard**,
- susceptibility to **pest and disease**,
- **aesthetic** value etc.

Vertical forest stands structure

Stand structure is the OUTER reflection of the processes.

The study of vertical structure may make a significant contribution to the knowledge about growth and developmental processes of forest ecosystems.

Methods for vertical structures quantification

- **arbitrarily** defined and do not represent natural stratification patterns of forests
- are **too time consuming** for landscape analyses
- can be **too expensive**
- **inappropriate** at dense stands

To improve the methodological shortages we conducted a very detailed study of the vertical stand structure.

Goals

To identify properties of the vertical structure in both managed and virgin forest.

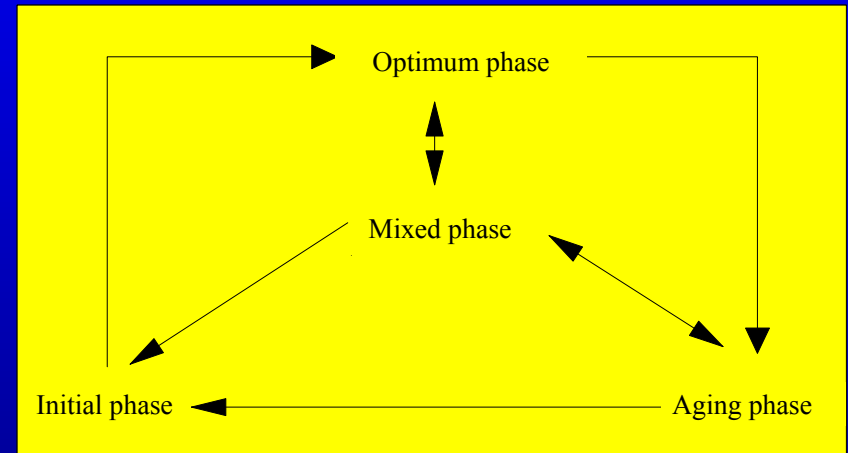
Case study

- study plots :
 - the **virgin forest remnant Rajhenavski Rog**,
 - **lightly managed forest in its vicinity**.
- the selection of **study plots** was restricted to the most dominant forest plant community *Omphalodo-Fagetum omphalodetosum*
=> high karst **Dinaric forests with silver fir** (*Abies alba* Mill.) and **beech** (*Fagus sylvatica* L.) as the most dominant tree species.

Case study

Stand dynamics was **described** by **four** indicative forest **cyclical developmental** phases, which have distinctive vertical and horizontal stand structures:

- a **juvenile** phase,
- an **optimal** phase,
- a **mixed** phase,
- a **regeneration** phase.



Sampling and data

Four research plots (35m by 35m) were **randomly** selected within optimal, mixed and regeneration developmental phase in **both** managed and virgin forest.

Stand structure was described by the following **attributes**:

- **tree species,**
- **dimager at breast high (DBH),**
- **tree height,**
- **layer,**
- **depth of the crown,**
- **width of the crown,**
- **social position,**
- **vitality.**

Data analysis

Structural patterns of vertical stand structure were **studied by automated data analysis** using **machine learning** techniques:

- **classification trees**
- **regression trees**

Classification trees

They predict the value of a **discrete dependent variable** with a **finite** set of values (called **class**)
from
the values of a set of **independent variables** (called **attributes**),
which may be either **continuous or discrete**.

Data mining analysis was performed by the **Weka** machine learning package.

We used **J4.8 algorithm**, which is Weka's implementation of **C4.5** decision tree algorithm - one of the most widely used decision tree system.

Regression trees

They predict the value of a **continuous dependent variable** (called **class**) or **linear function of some attributes** from the values of a set of **independent variables** (called **attributes**), which may be either **continuous or discrete**.

Data mining analysis was performed by the **Weka** machine learning package. We used **M5 algorithm** as one of the most widely used regression tree system.

Results – Classification trees

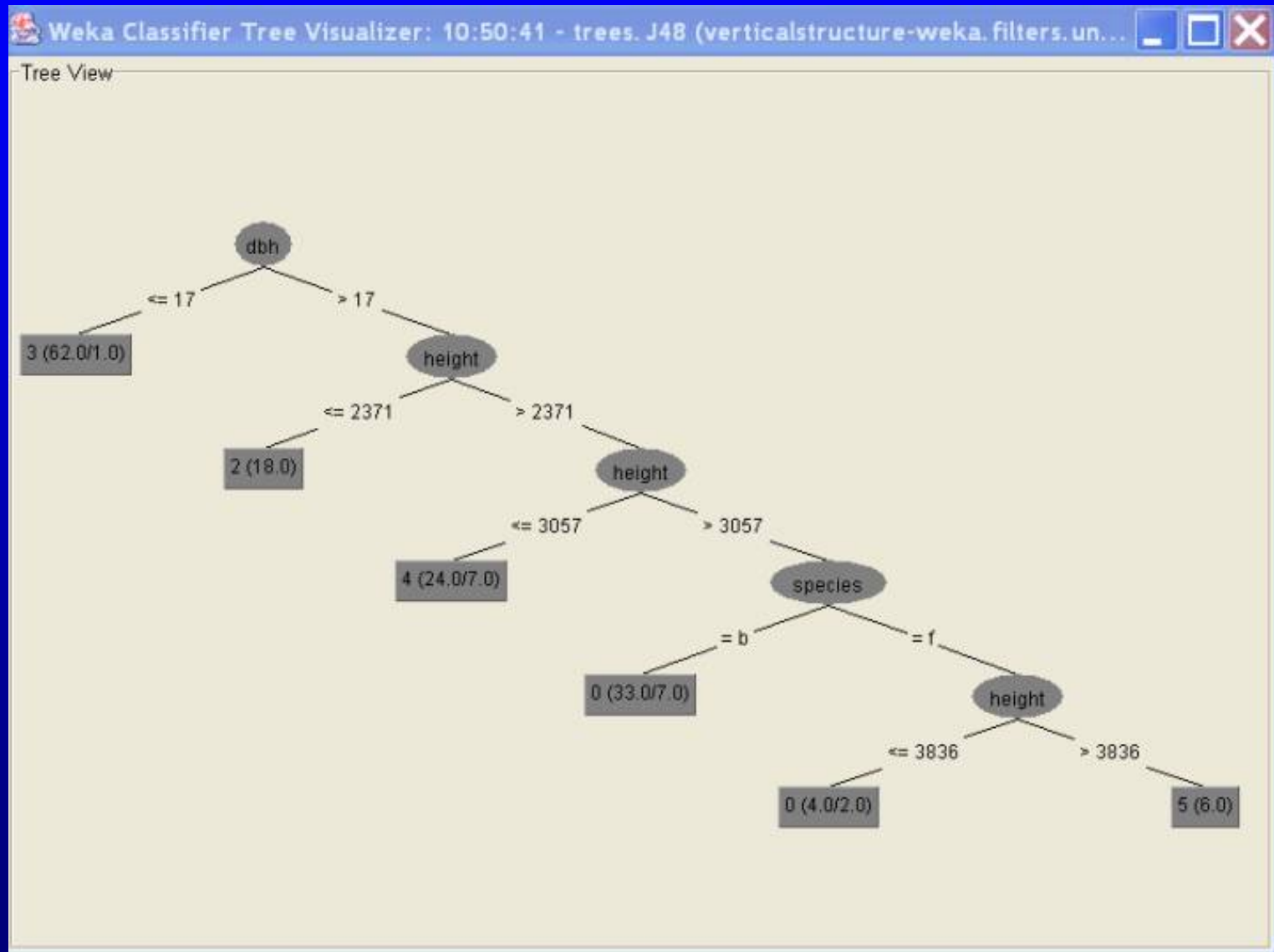
Induced models 24 different models:

| | Optimal phase | Mixed phase | Regeneration phase |
|---------------|--|--|--|
| VIRGIN forest | layer crown depth crown width 195 instances | layer crown depth crown width 303 instances | layer crown depth crown width 147 instances |
| MANGED forest | layer crown depth crown width 336 instances | layer crown depth crown width 325 instances | layer crown depth crown width 218 instances |

| | |
|---------------|--|
| VIRGIN forest | layer crown depth crown width 645 instances |
| MANGED forest | layer crown depth crown width 879 instances |

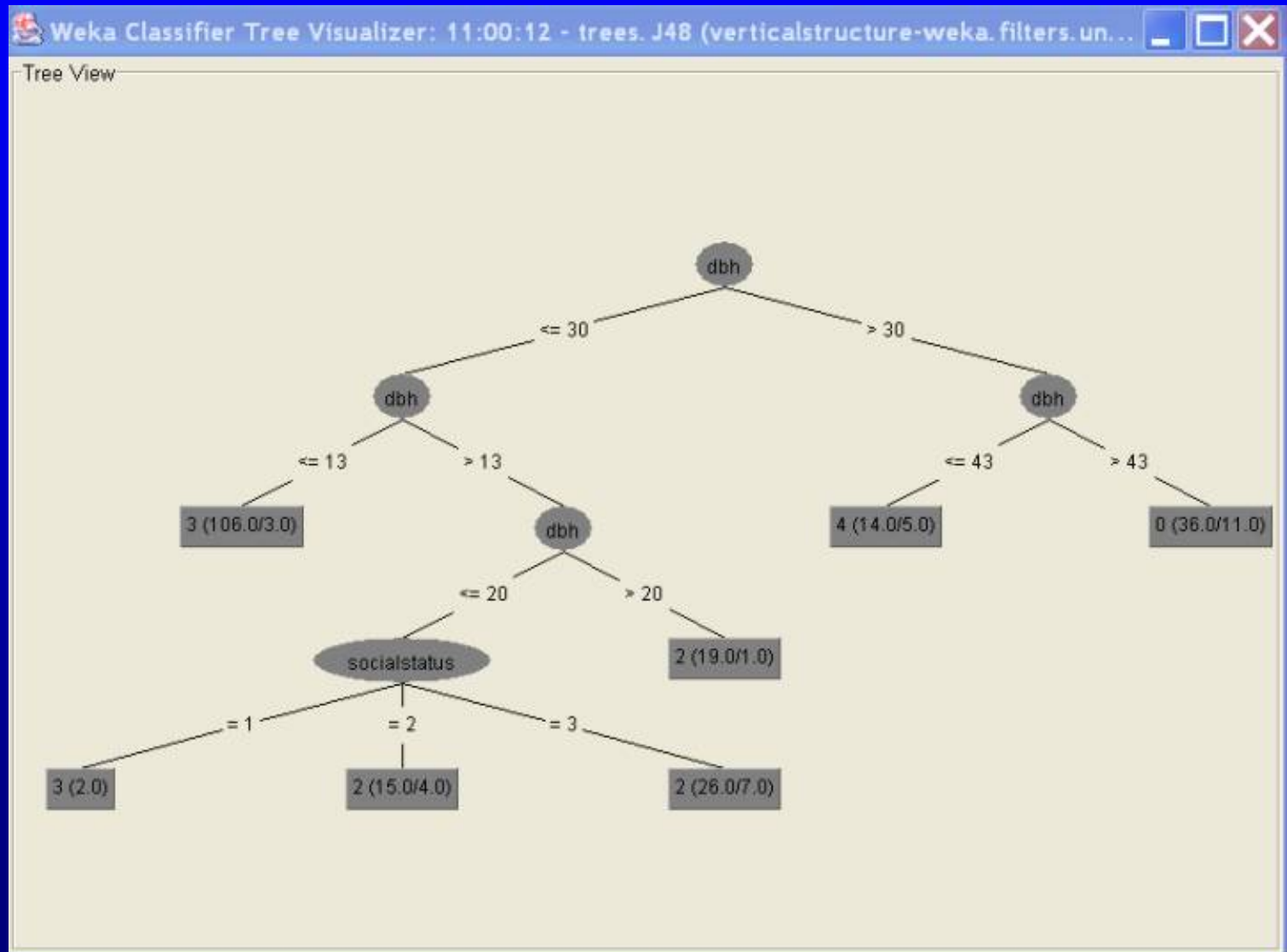
Results – Classification trees

Layer: virgin f.- regeneration ph.



Results – Classification trees

Layer: managed f.- regeneration ph.



Results – Classification trees

Layer m.: virgin f.: regeneration

```

dbh <= 17: 3 (62.0/1.0)
dbh > 17
| height <= 2371: 2 (18.0)
| height > 2371
| | height <= 3057: 4 (24.0/7.0)
| | height > 3057
| | | species = b: 0 (33.0/7.0)
| | | species = f
| | | | height <= 3836: 0 (4.0/2.0)
| | | | height > 3836: 5 (6.0)

```

| | | |
|----------------------------------|-----------|-----------|
| Correctly Classified Instances | 124 | 84.3537 % |
| Incorrectly Classified Instances | 23 | 15.6463 % |
| Kappa statistic | 0.7853 | |
| Mean absolute error | 0.0866 | |
| Root mean squared error | 0.2299 | |
| Relative absolute error | 29.4685 % | |
| Root relative squared error | 60.065 % | |
| Total Number of Instances | 147 | |

Layer m.: managed f. regeneration

```

dbh <= 30
| dbh <= 13: 3 (106.0/3.0)
| dbh > 13
| | dbh <= 20
| | | socialstatus = 1: 3 (2.0)
| | | socialstatus = 2: 2 (15.0/4.0)
| | | socialstatus = 3: 2 (26.0/7.0)
| | dbh > 20: 2 (19.0/1.0)
dbh > 30
| dbh <= 43: 4 (14.0/5.0)
| dbh > 43: 0 (36.0/11.0)

```

| | | |
|----------------------------------|-----------|-----------|
| Correctly Classified Instances | 177 | 81.1927 % |
| Incorrectly Classified Instances | 41 | 18.8073 % |
| Kappa statistic | 0.7105 | |
| Mean absolute error | 0.0951 | |
| Root mean squared error | 0.233 | |
| Relative absolute error | 37.1597 % | |
| Root relative squared error | 65.3369 % | |
| Total Number of Instances | 218 | |

Results – Classification trees - conclusions

24 models : the **independent attributes** that **correlates** the most with the **selected (dependent) class-attribute**

Layer models: DBH, crown depth, social status

Crown depths models: social status, layer, tree height

Crown width models: vitality, crown depth, social status

Results – Classification trees - conclusions

Patterns of relationships among attributes:

lower DBH => lower layer => the thresholds between layers are lower in managed than in virgin forest

narrow crowns => lower vitality or/and trees are groups

low crown depths => trees are in groups

Results – Classification trees - conclusions

**VIRGINE forest: high diversity in vertical structures
between development phases**

**MANAGED forest: homogeneous vertical structure =>
no differences between development phases**

Results - regression trees

Virgin forest height model:

Instances: 645

Managed forest height model:

Instances: 879

Results - regression trees

Virgin forest: model of tree's height

| | | |
|------------------------------|-----------------------------|----------|
| dbh ≤ 24.5 : | Correlation coefficient | 0.9962 |
| dbh ≤ 7.5 : LM1 (209/6.253%) | Mean absolute error | 0.9469 |
| dbh > 7.5 : LM2 (198/3.491%) | Root mean squared error | 2.1788 |
| dbh > 24.5 : | Relative absolute error | 4.5358 % |
| dbh ≤ 44.5 : LM3 (88/6.504%) | Root relative squared error | 8.7022 % |
| dbh > 44.5 : | Total Number of Instances | 645 |
| dbh ≤ 61.5 : LM4 (68/4.559%) | | |
| dbh > 61.5 : LM5 (82/6.126%) | | |

LM num: 1

height = 4.9316 * phase=3,2 - 53.6045 * species=f + 118.5593 * dbh - 1.555 * crowndepth=2 - 5.7753 * socialstatus=1,2 + 6.7971 * socialstatus=2 + 4.3573 * vitality=1,2 + 52.7343

LM num: 2

height = 4.9316 * phase=3,2 - 723.1937 * species=f + 91.5362 * dbh - 1.555 * crowndepth=2 - 5.7753 * socialstatus=1,2 + 6.7971 * socialstatus=2 + 4.4243 * vitality=1,2 + 338.7362

LM num: 3

height = 8.2259 * phase=3,2 - 391.9149 * species=f + 48.393 * dbh - 2.5937 * crowndepth=2 - 9.6331 * socialstatus=1,2 + 11.3375 * socialstatus=2 + 5.1068 * vitality=1,2 + 1220.2757

LM num: 4

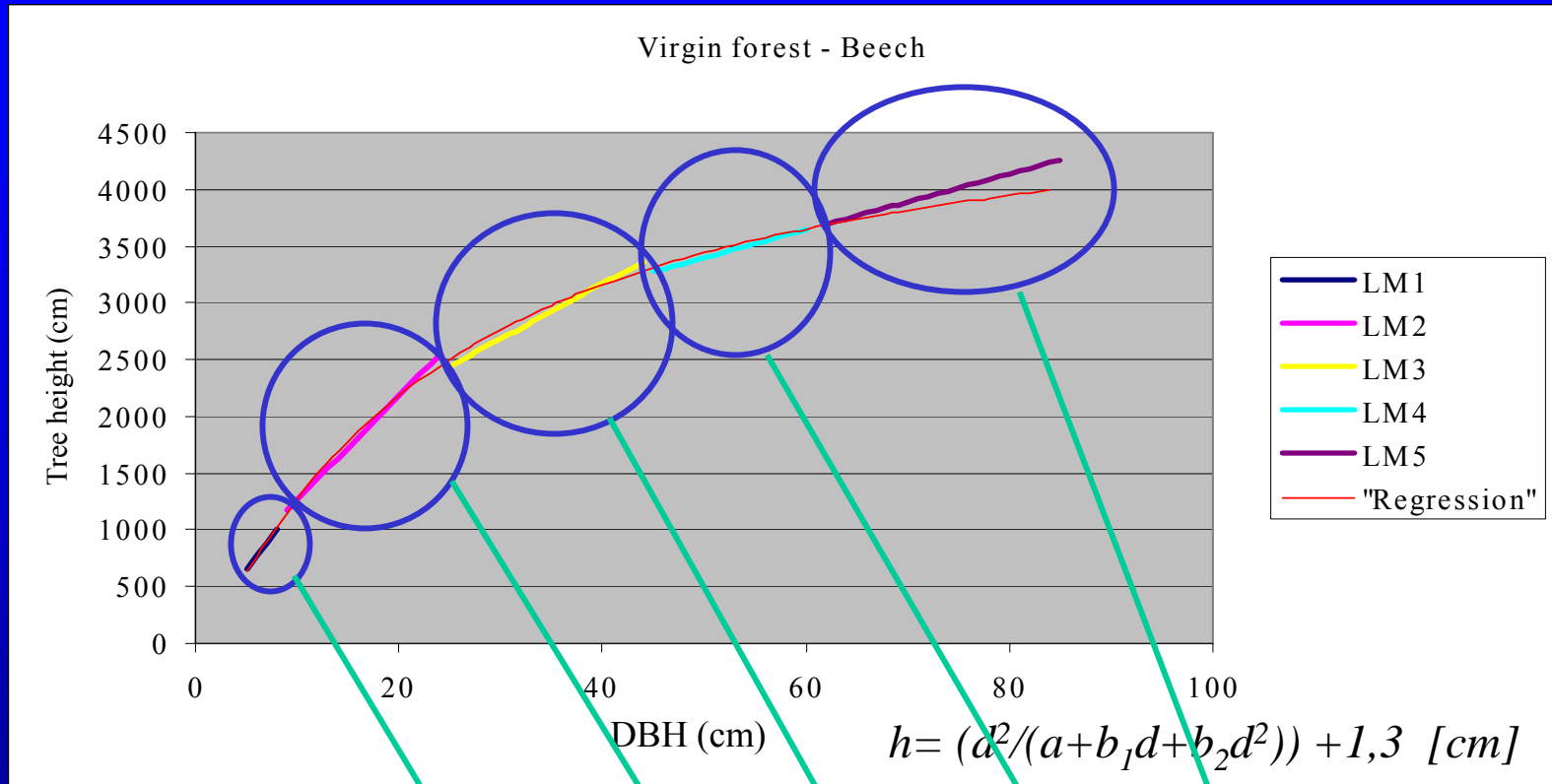
height = 8.2259 * phase=3,2 - 28.015 * species=f + 24.6605 * dbh - 2.5937 * crowndepth=2 - 9.6331 * socialstatus=1,2 + 11.3375 * socialstatus=2 + 5.1068 * vitality=1,2 + 2150.9432

LM num: 5

height = 8.2259 * phase=3,2 - 28.015 * species=f + 16.9728 * dbh - 2.5937 * crowndepth=2 - 9.6331 * socialstatus=1,2 + 11.3375 * socialstatus=2 + 5.1068 * vitality=1,2 + 2665.7544

Results - regression trees

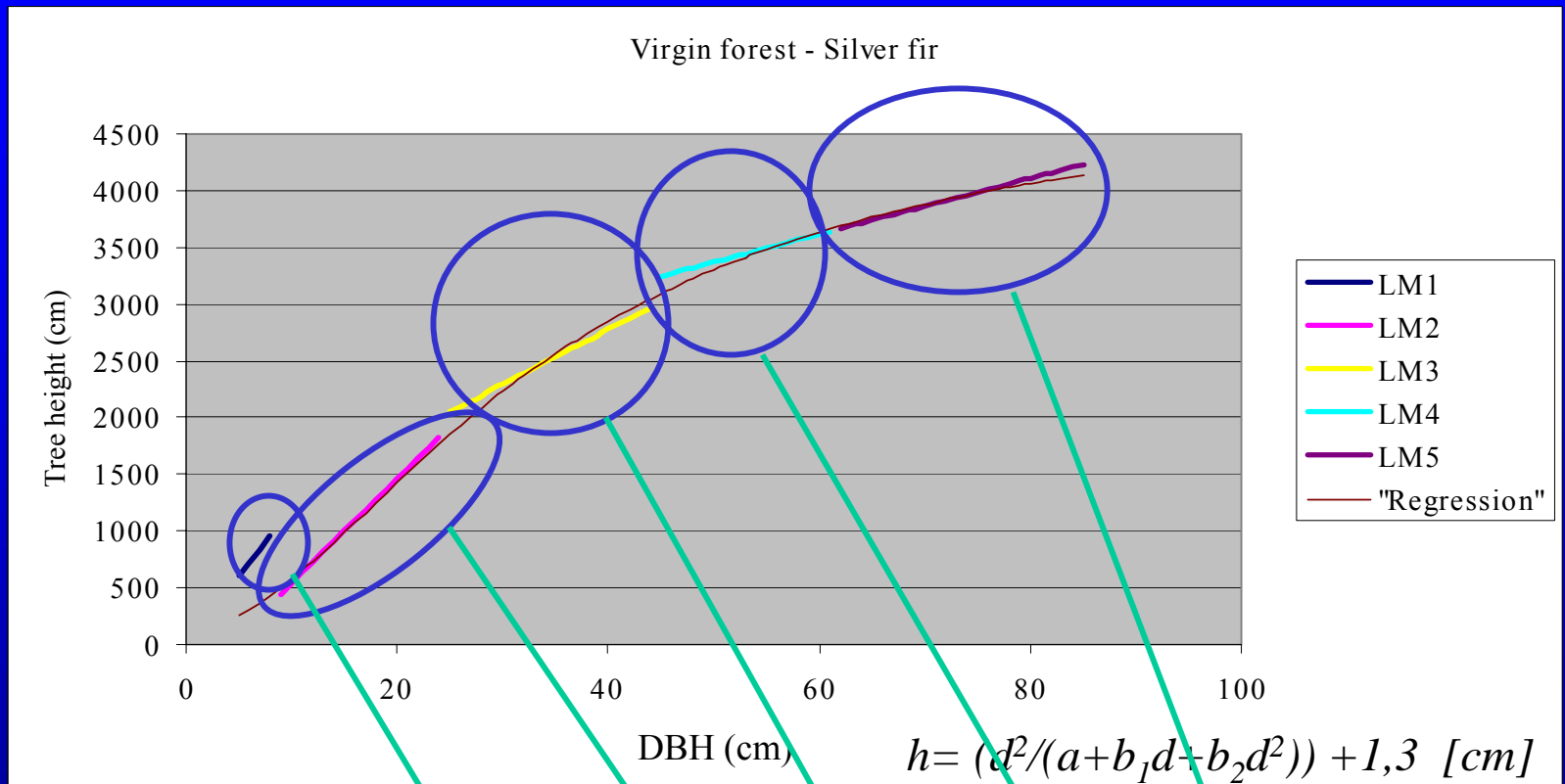
Virgin forest: model of tree's height



| Virgin f. | LM1 | LM2 | LM3 | LM4 | LM5 |
|------------------|---------|----------|----------|----------|----------|
| phase=3,2 | 4.932 | 4.932 | 8.226 | 8.226 | 8.226 |
| species=f | -53.605 | -723.194 | -321.915 | -28.015 | -28.015 |
| dbh | 118.559 | 91.536 | 48.393 | 24.661 | 16.973 |
| crowndepth=2 | -1.555 | -1.555 | -2.594 | -2.594 | -2.594 |
| socialstatus=1,2 | -5.775 | -5.775 | -9.633 | -9.633 | -9.633 |
| socialstatus=2 | 6.797 | 6.797 | 11.338 | 11.338 | 11.338 |
| vitality=1,2 | 4.357 | 4.424 | 5.107 | 5.107 | 5.107 |
| n | 52.734 | 338.736 | 1220.276 | 2150.943 | 2665.754 |

Results - regression trees

Virgin forest height model



| Virgin f. | LM1 | LM2 | LM3 | LM4 | LM5 |
|------------------|---------|----------|----------|----------|----------|
| phase=3,2 | 4.932 | 4.932 | 8.226 | 8.226 | 8.226 |
| species=f | -53.605 | -723.194 | -321.915 | -28.015 | -28.015 |
| dbh | 118.559 | 91.536 | 48.393 | 24.661 | 16.973 |
| crowndepth=2 | -1.555 | -1.555 | -2.594 | -2.594 | -2.594 |
| socialstatus=1,2 | -5.775 | -5.775 | -9.633 | -9.633 | -9.633 |
| socialstatus=2 | 6.797 | 6.797 | 11.338 | 11.338 | 11.338 |
| vitality=1,2 | 4.357 | 4.424 | 5.107 | 5.107 | 5.107 |
| n | 52.734 | 338.736 | 1220.276 | 2150.943 | 2665.754 |

Results - regression trees

Managed forest height model

| | | |
|--------------------------------|-----------------------------|-----------|
| dbh ≤ 17.5 : LM1 (559/15.305%) | Correlation coefficient | 0.9923 |
| dbh > 17.5 : | Mean absolute error | 41.7166 |
| dbh ≤ 35.5 : LM2 (194/18.573%) | Root mean squared error | 87.8429 |
| dbh > 35.5 : | Relative absolute error | 7.3238 % |
| dbh ≤ 54.5 : LM3 (69/7.654%) | Root relative squared error | 12.3599 % |
| dbh > 54.5 : LM4 (57/4.948%) | Total Number of Instances | 879 |

LM num: 1

height = 64.7499 * dbh + 4.9658 * layer=4,0,5 + 0.7611 * crowndepth=2 + 2.6777 * vitality=2,1 + 525.534

LM num: 2

height = 39.7414 * dbh + 77.4803 * layer=4,0,5 + 1.304 * crowndepth=2 + 4.588 * vitality=2,1 + 954.3475

LM num: 3

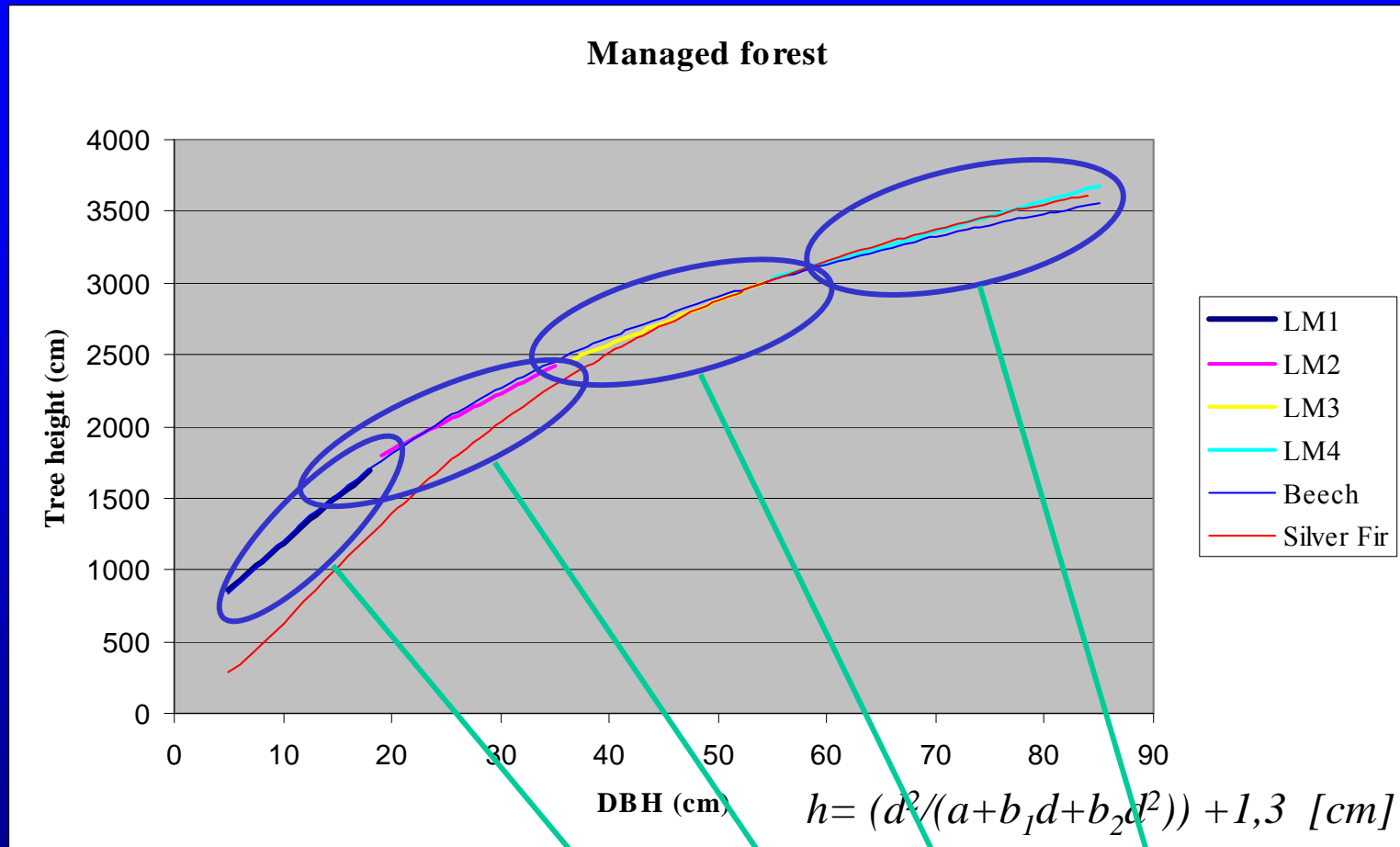
height = 30.2793 * dbh + 24.2647 * layer=4,0,5 + 1.304 * crowndepth=2 + 4.588 * vitality=2,1 + 1329.8797

LM num: 4

height = 21.4912 * dbh + 24.2647 * layer=4,0,5 + 1.304 * crowndepth=2 + 4.588 * vitality=2,1 + 1823.0698

Results - regression trees

Managed forest height model



| Managed f. | LM1 | LM2 | LM3 | LM4 |
|--------------|---------|----------|---------|---------|
| dbh | 64.7499 | 39.7414 | 30.2793 | 21.4912 |
| layer=4,0,5 | 4.9658 | 77.4803 | 24.2647 | 24.2647 |
| crowndepth=2 | 0.7611 | 1.304 | 1.304 | 1.304 |
| vitality=2,1 | 2.6777 | 4.588 | 4.588 | 4.588 |
| n | 525.534 | 954.3475 | 1329.88 | 1823.07 |

Results – Regression trees - Conclusions

Explanations of the shape of the Prodan's tree height curve.

M5 didn't distinguish between beech and silver fir tree height curves.

General conclusions

Classification trees: identification of the most descriptive attributes of vertical structure and their hierarchic relationship.

Regression trees: identification of the attributes which have the strongest effect on the Prodan's tree-height regression curve.

Managed forest has unified vertical structure while virgin forest show distinguished differences among development phases.

Thank you!