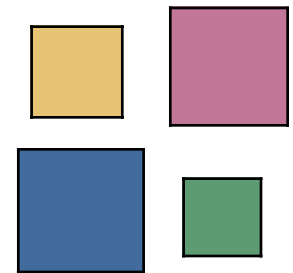


TWIX

Trees With eXtra splits

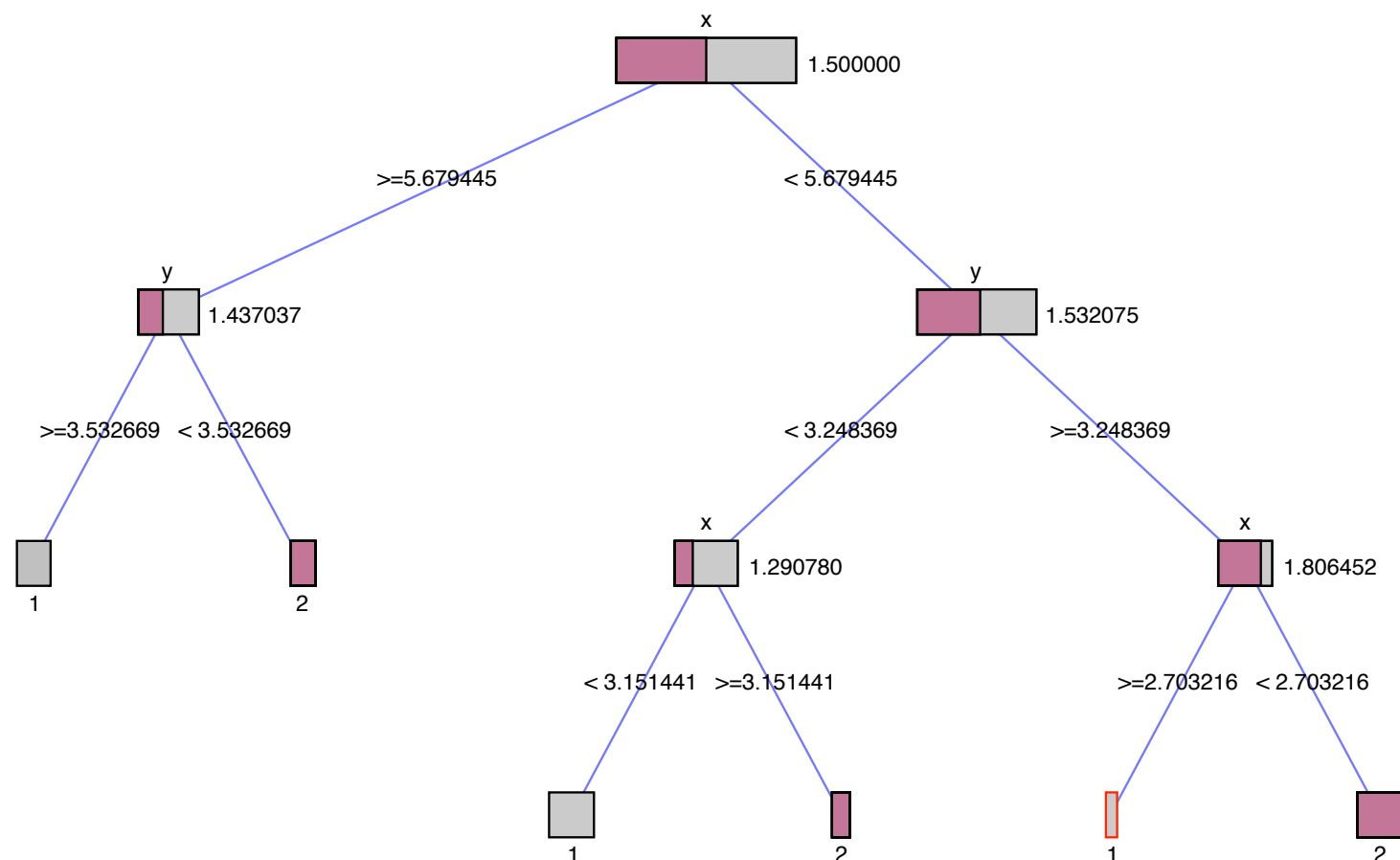
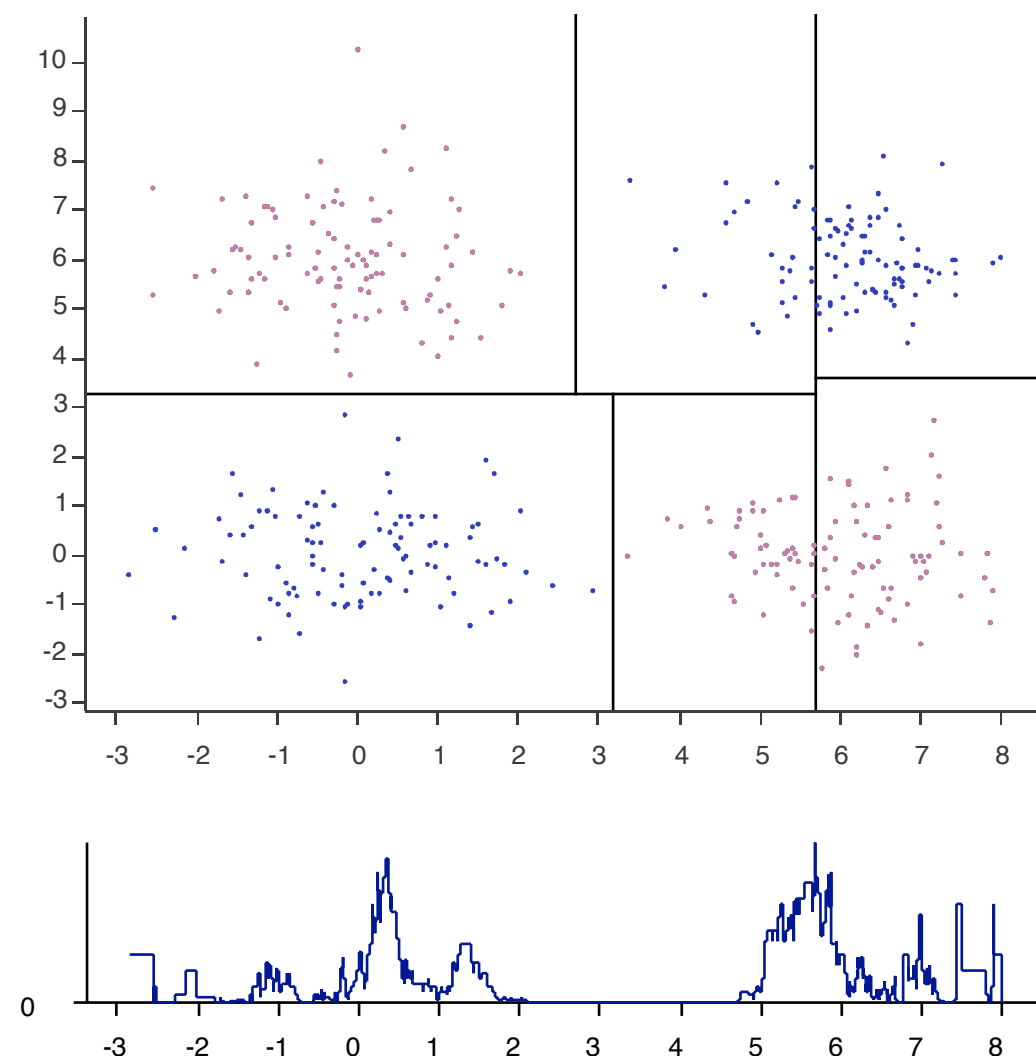
Sergej Potapov
Martin Theus
Simon Urbanek

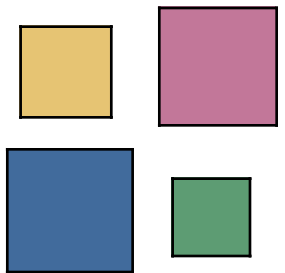


Motivation

- Where the classical CART algorithm fails
 - Greedy algorithms never go for a (locally) second best solution, which would result in a better overall (global) solution.

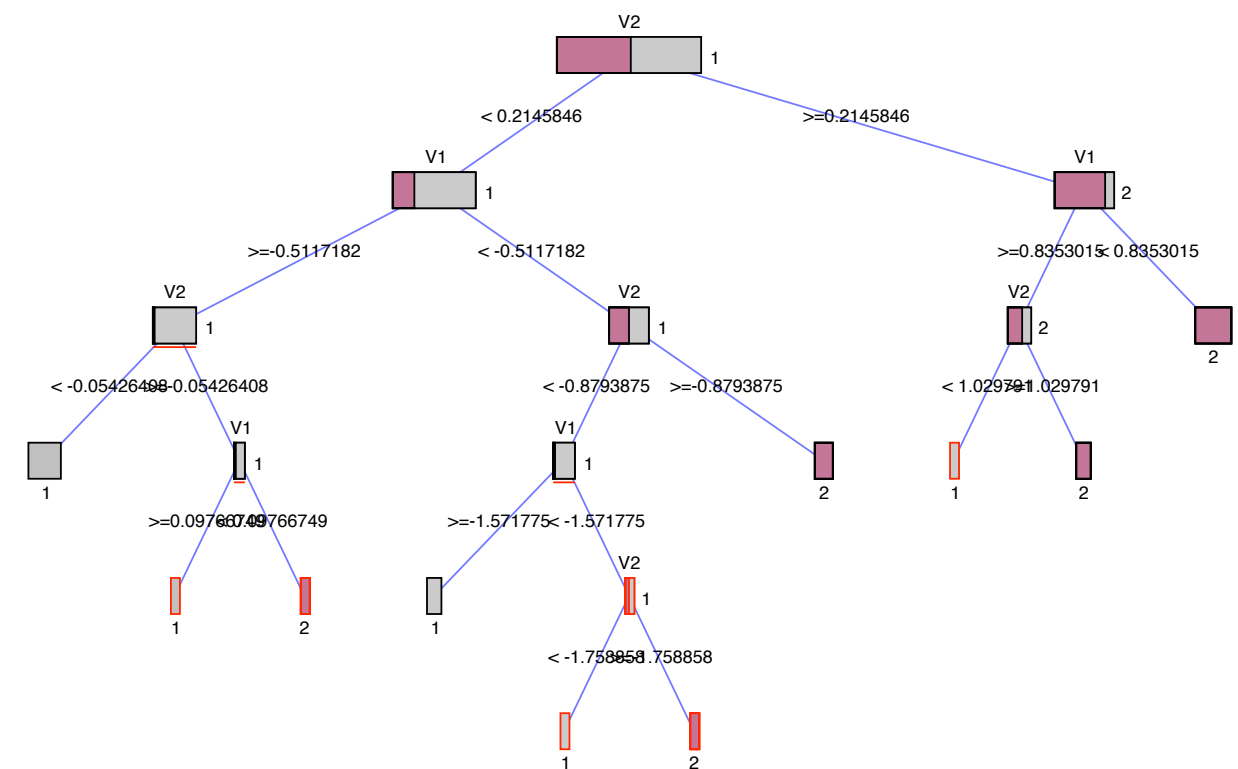
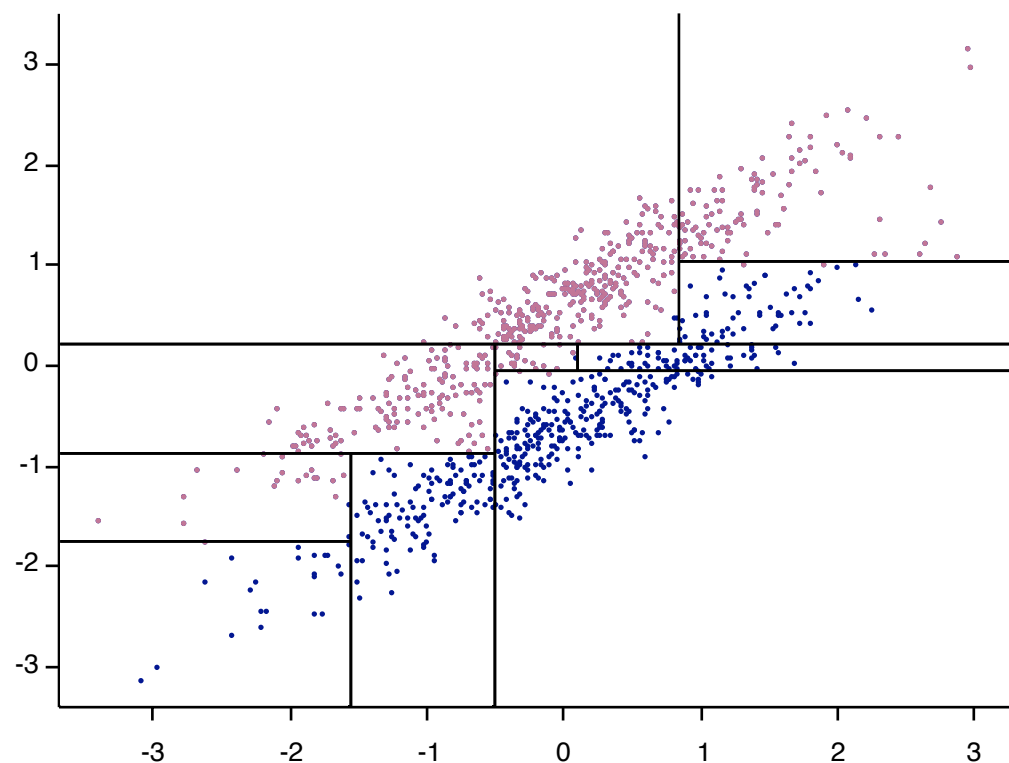
Example: XOR-data



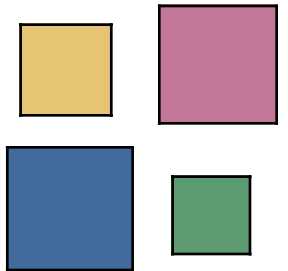


Trees: More Problems

- Non-orthogonal splitting directions ...



... can not be handled by single trees, no matter how we split



Bagging Revisited

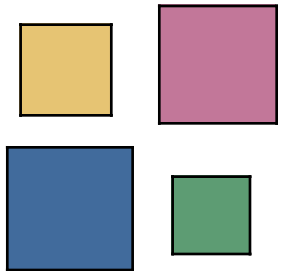
Bagging = **B**ootstrap **A**ggregation, tries to simulate an infinite sample by bootstrapping, i.e. sampling from the original sample with replacement.

Repeat N times:

1. Generate a bootstrap sample D_i of size n .
2. Fit model $\hat{f}_{D_i}$.

Depending on the problem the N results are aggregated:

- Classification: $g(x) = \operatorname{argmax}_{c \in C} \sum_{i=1}^N I(f_{D_i}(x) = c)$
- Regression: $g(x) = \frac{1}{N} \sum_{i=1}^N f_{D_i}(x)$

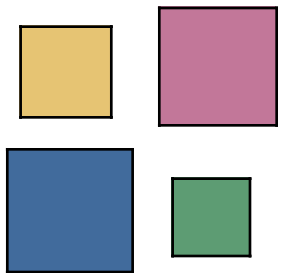


Ensembles

- **General Idea**
Use many “different” classifier and combine them to get more accurate results.
- Bagging: Instability of trees yields different models
- Random Forests: Restrict input space randomly to get wider range of models
- Boosting: Iterate to down-weight “bad” points

Question:

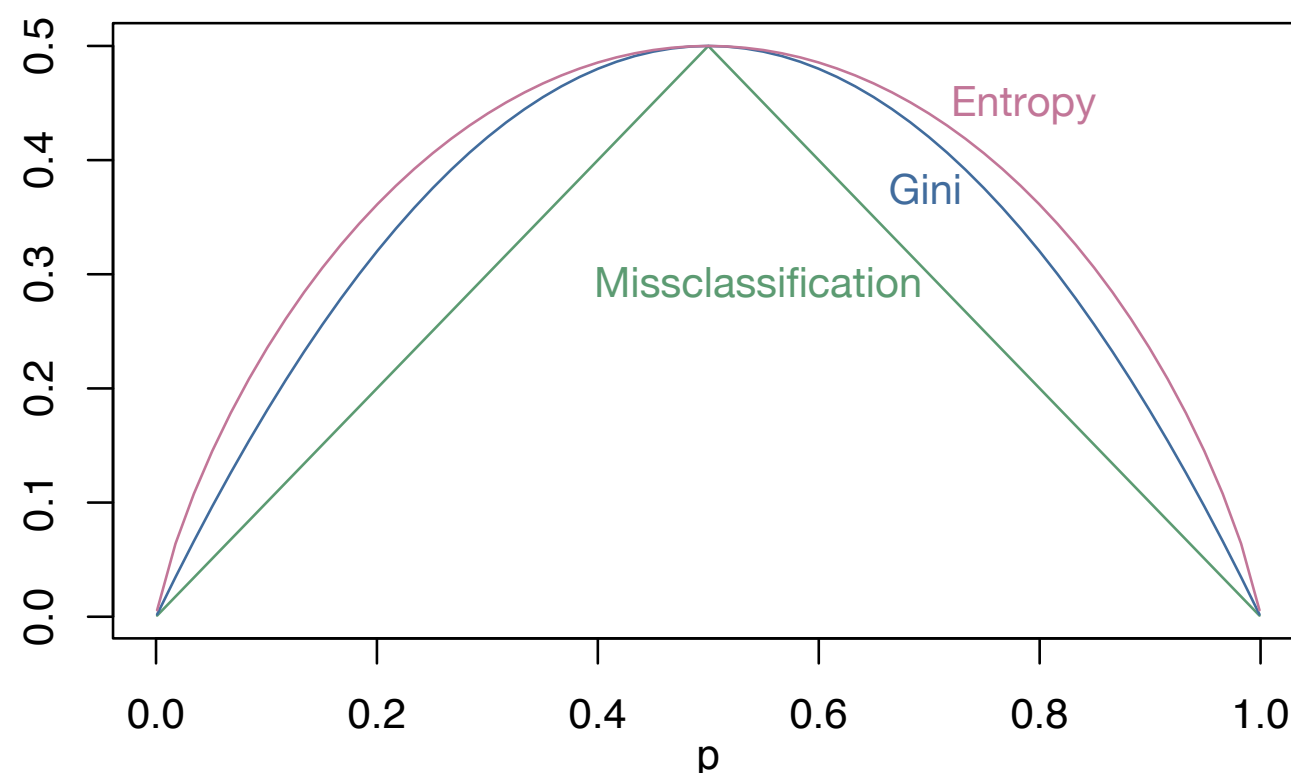
Why use randomly generated (sub-optimal) models?



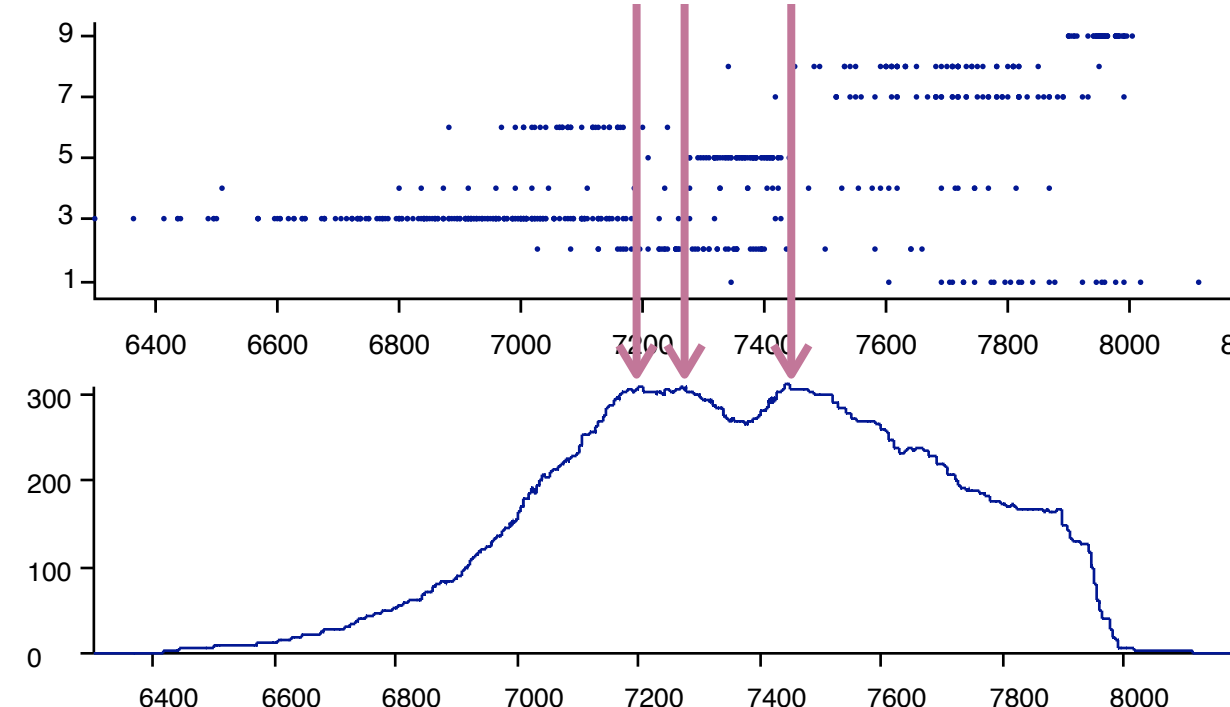
Tree Mechanics

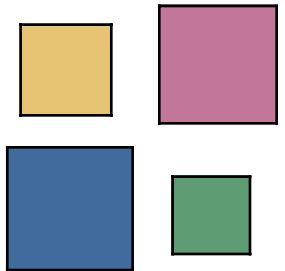
- CART is a recursive partitioning algorithm
- Each node is split according to the maximum gain in the loss function
- Mountain plots shows the loss function for a variable for all possible split points

Loss Functions



Mountain Plot



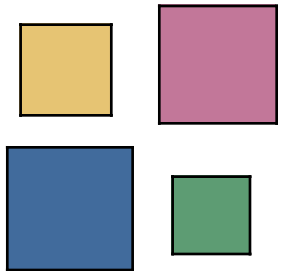


Idea behind TWIX

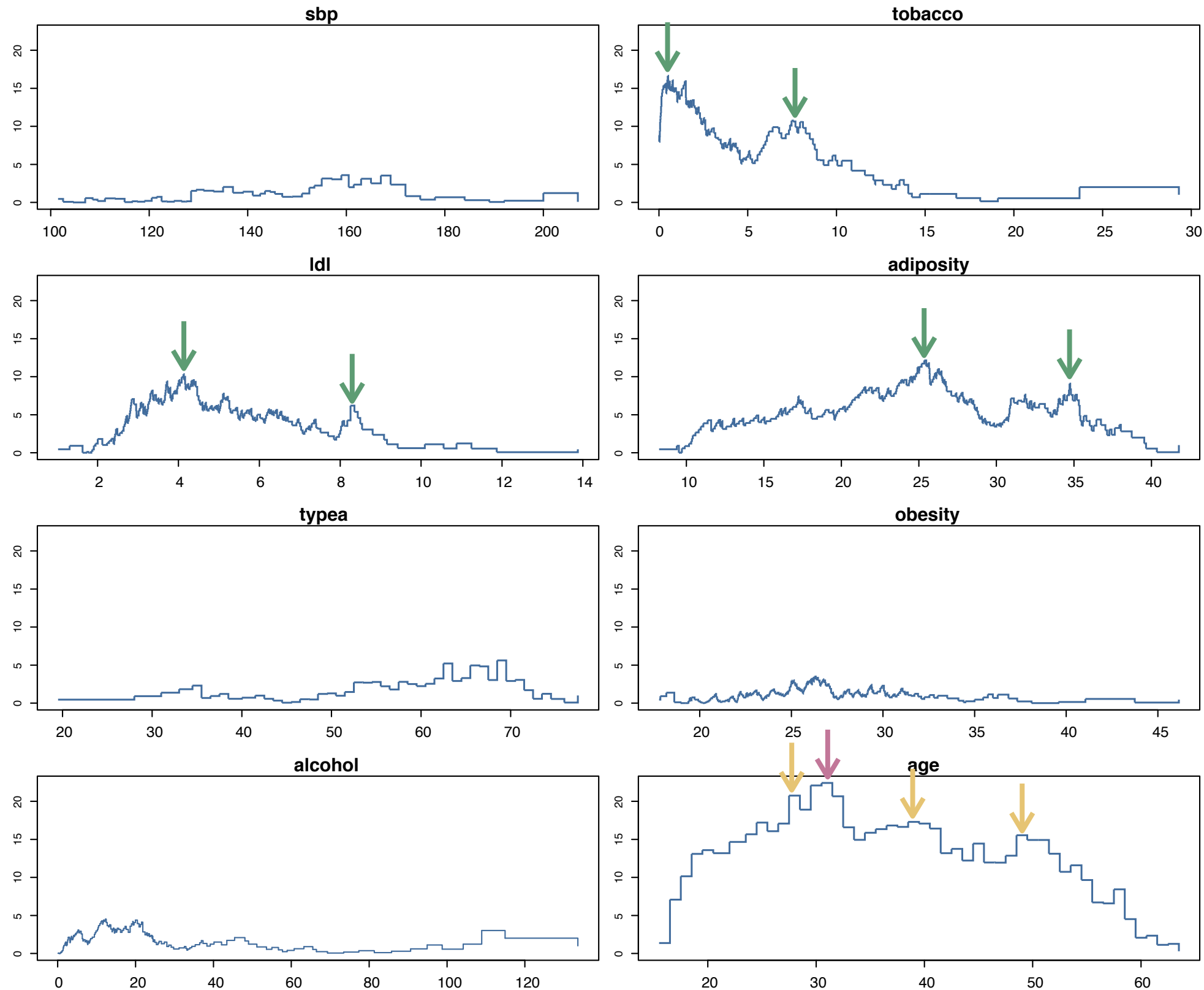
- Since the greedy CART algorithm not necessarily finds the “optimal” tree, try second best splits.
- Use these forests for bagging
- Expect better results for both single trees and aggregations

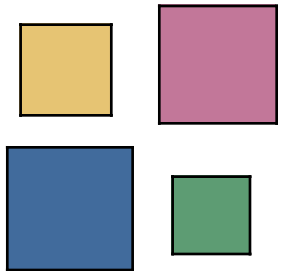
Problems

- How to find “good” candidates for second best splits?
- Number of inner nodes grows exponentially with the number of levels in the tree
⇒ so does the number of alternative trees



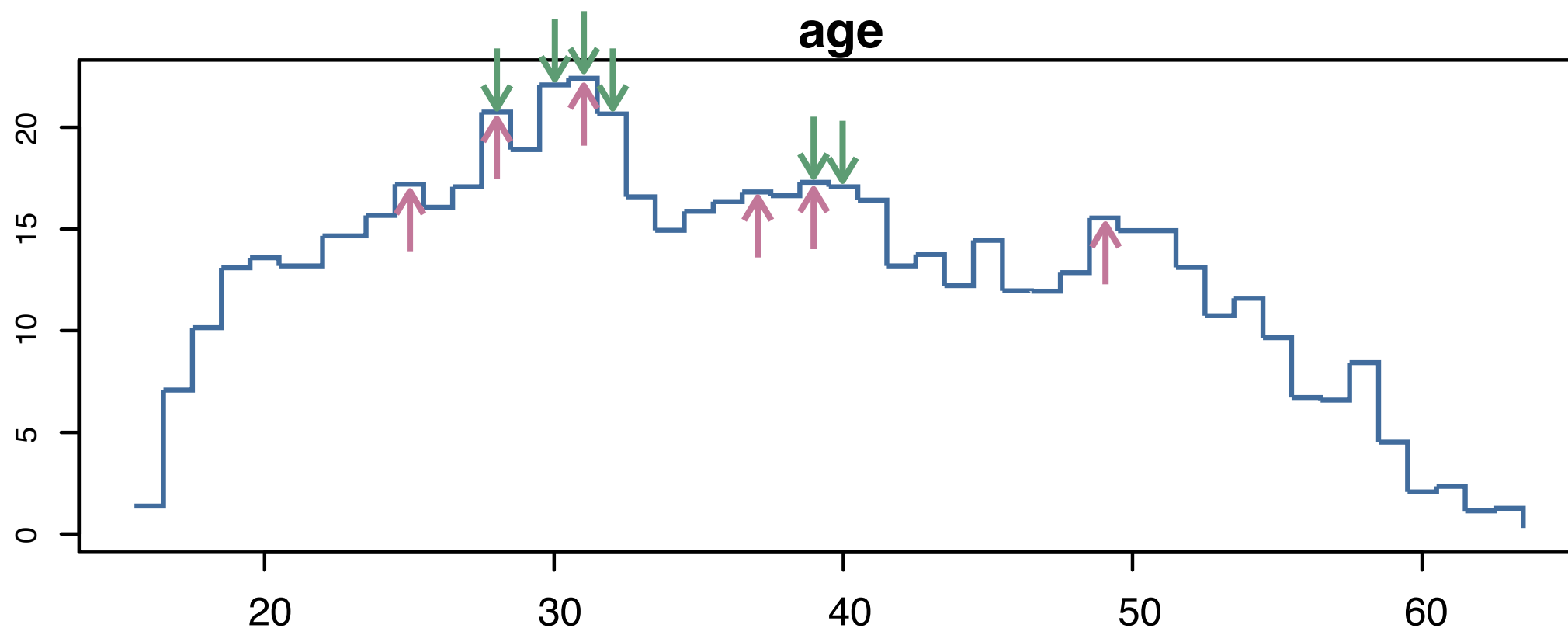
Second Best Splits: South African Heart Data

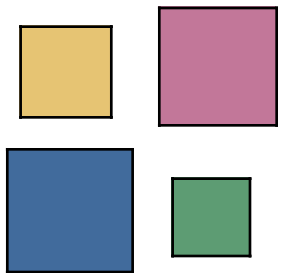




Second Best Splits: Global vs. Local

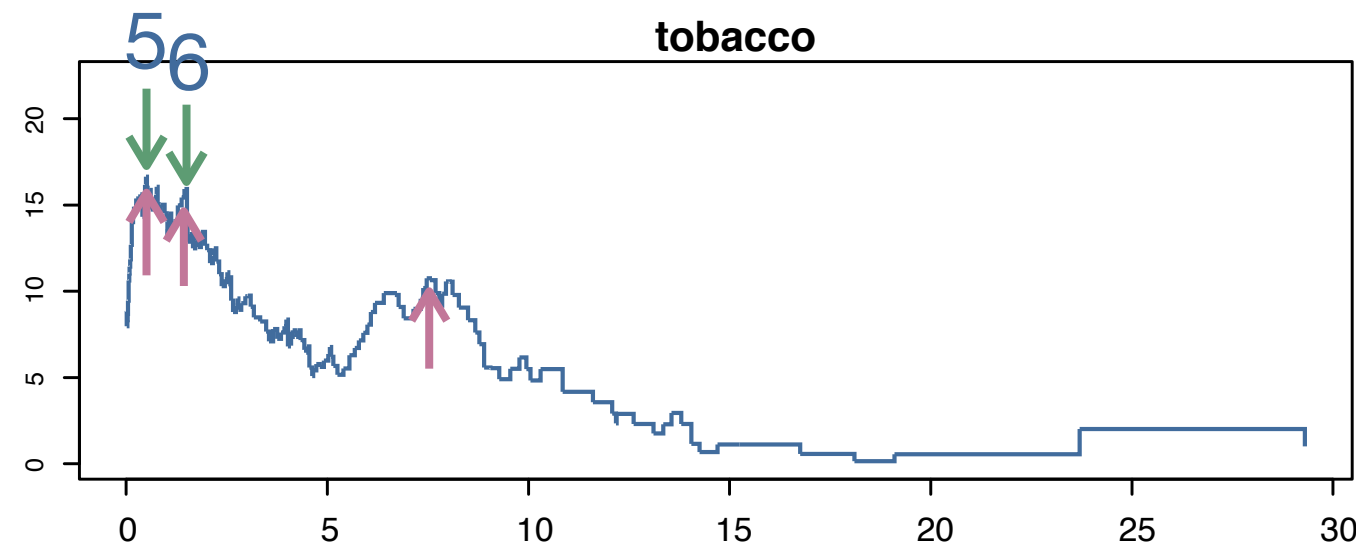
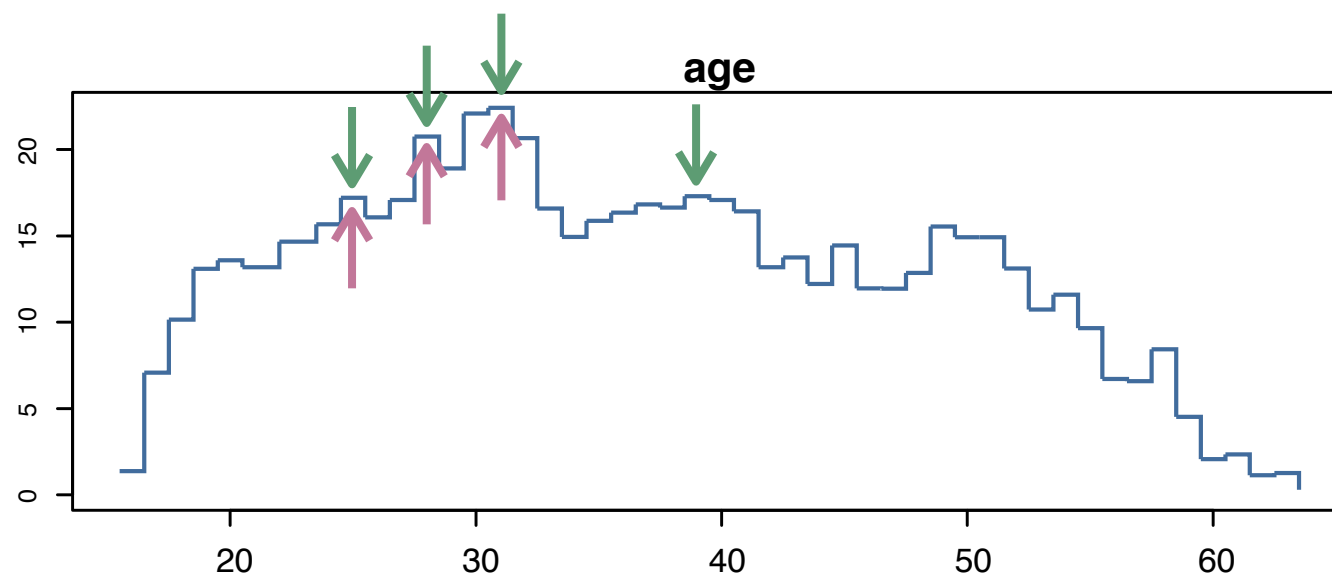
- When searching for a “best” split point, we can either look for
 - all top n greatest deviance gains, or
 - only look for local maxima
- Example
Top 6 splits

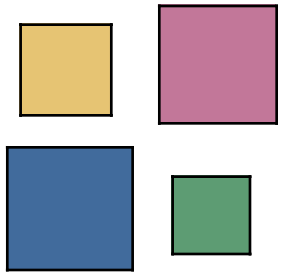




Second Best Splits: Forcing Variables

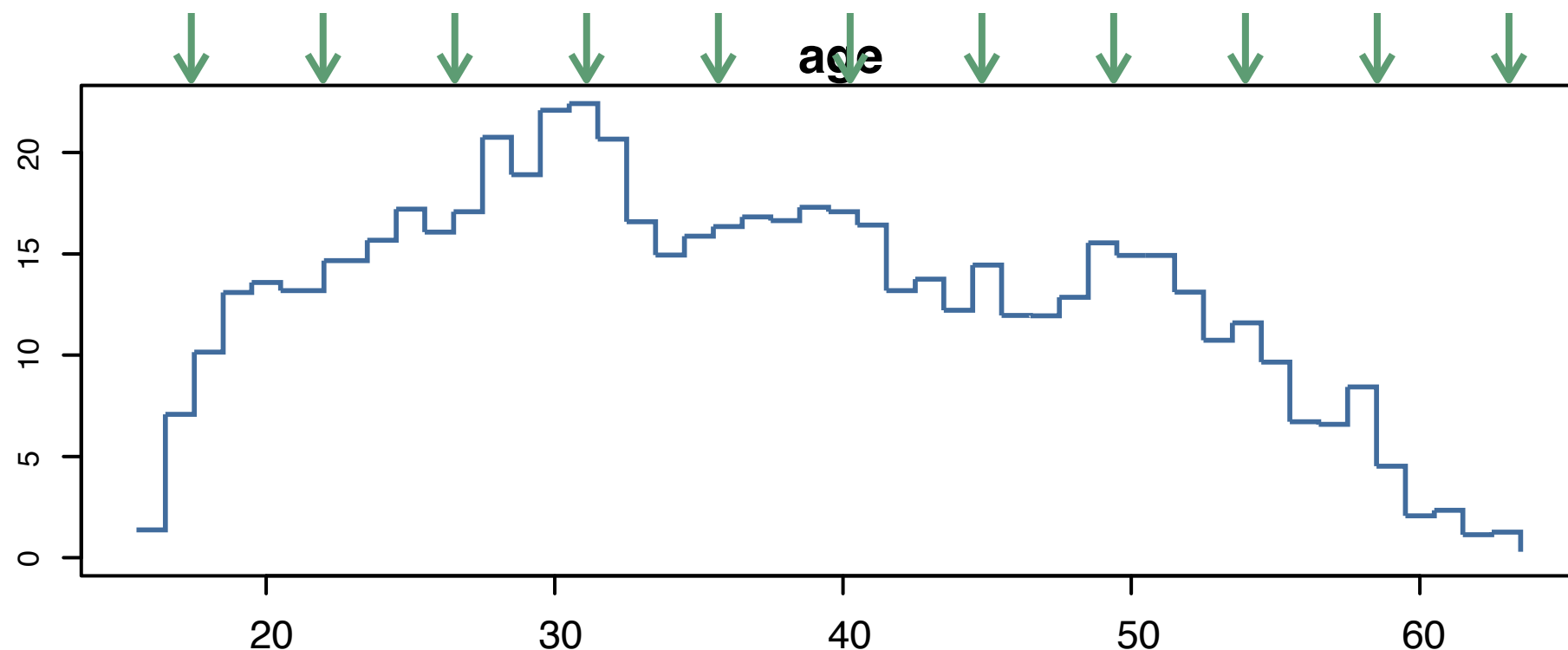
- Often a single variable dominates the potential deviance gain, and shadows all other variables
⇒ Many probably good split points are lost.
- Solution:
Force a minimum number of split points for **each** variable.
- Example: top 6 vs. top 3

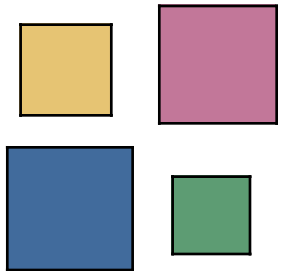




Second Best Splits: Grid Search

- In some situations good split points might not even be associated with some (local) maximum in deviance gain.
(Remember the XOR Example)
- Grid searches are most exhaustive, but also most expensive.





Implementation: The Grid

- If we allow s_j splits per node on level j of the tree, we get a maximum of

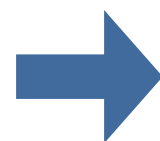
$$S = \prod_{i=1}^k s_i^{2^{i-1}}$$

trees for a tree with no more than k levels. Example:

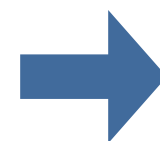
$$s = (7, 4, 2) \Rightarrow S = 7^{2^0} \cdot 4^{2^1} \cdot 2^{2^2} = 7 \cdot 16 \cdot 16 = 1792$$

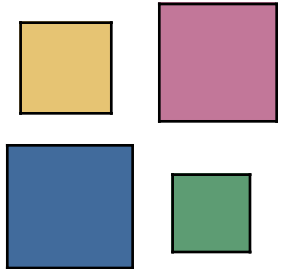
$\Rightarrow$ Work on a grid of computers

Controller



PVM





Implementation: The R-Package

TWIX {TWIX}

R Documentation

Top Classification Trees

Description

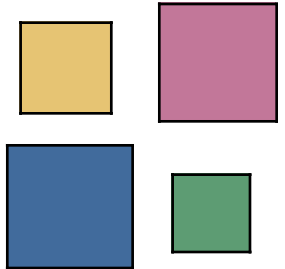
Top Classification Trees

Usage

```
TWIX(formula, data = NULL, test.data = NULL, subset = NULL,  
      method = "deviance", topn.method = "complete", cluster = NULL,  
      minsplit = 20, minbucket = round(minsplit/3), Devmin = 0.01,  
      topN = 1, level = 6, st = 1, cl.level = 2, tol = 0.01, ...)
```

Arguments

<code>formula</code>	formula of the form $y \sim x_1 + x_2 + \dots$, where y must be a factor and $x_1, x_2, \dots$ are numeric.
<code>data</code>	an optional data frame containing the variables in the model(training data).
<code>test.data</code>	a data frame containing new data.
<code>subset</code>	an optional vector specifying a subset of observations to be used.
<code>method</code>	Which split points will be used? This can be "deviance" (default), "grid" or "local". If the method is set to: <i>local</i> the program uses the local maxima of the split function(entropy), <i>deviance</i> all values of the entropy, <i>grid</i> grid points.
<code>topn.method</code>	one of "complete"(default) or "single". A specification of the consideration of the split points. If set to "complete" it uses split points from all variables, else it uses split points per variable.
<code>cluster</code>	name of the cluster, if parallel computing will be used.
<code>minsplit</code>	the minimum number of observations that must exist in a node.
<code>minbucket</code>	the minimum number of observations in any terminal <leaf> node.
<code>Devmin</code>	the minimum improvement on entropy by splitting.
<code>topN</code>	integer vector. How many splits will be selected and at which level? If length 1, the same size of splits will be selected at each level. If length > 1, for example <code>topN=c(3,2)</code> , 3 splits will be chosen at first level, 2 splits at second level and for all next levels 1 split.
<code>level</code>	maximum depth of the trees. If <code>level</code> set to 1, trees consist of root node.
<code>st</code>	step parameter for method "grid".
<code>cl.level</code>	parameter for parallel computing.
<code>tol</code>	parameter, which will be used, if <code>topn.method</code> is set to "single".
<code>...</code>	further arguments to be passed to or from methods.



“Driving the Beast”

- The most important tuning parameters are

- `method`

Which split points will be used? This can be "deviance" (default), "grid" or "local". If the `method` is set to: *local* the program uses the local maxima of the split function(entropy), *deviance* all values of the entropy, *grid* grid points.

- `topn.method`

one of "complete"(default) or "single". A specification of the consideration of the split points. If set to "complete" it uses split points from all variables, else it uses split points per variable.

- `topN`

integer vector. How many splits will be selected and at which level? If length 1, the same size of splits will be selected at each level. If length > 1, for example `topN=c(3,2)`, 3 splits will be chosen at first level, 2 splits at second level and for all next levels 1 split.

- `level`

maximum depth of the trees. If `level` set to 1, trees consist of root node.

- Stopping Rules:

- `minsplit`

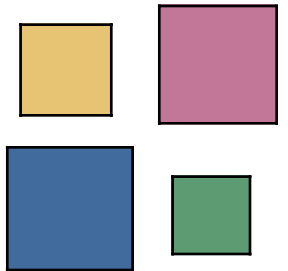
the minimum number of observations that must exist in a node.

- `minbucket`

the minimum number of observations in any terminal <leaf> node.

- `Devmin`

the minimum improvement on entropy by splitting.



South African Heart Desease Data cont.

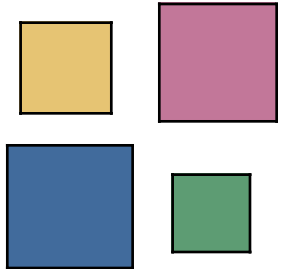
- To get a “fair”, i.e. generalizable and not too overfitted classifier, we usually split the data into 3 chunks:



- **Training**
All models are trained using the training data
- **Validation**
The “best” model is selected using the validation data
(The chosen model is then estimated with training+validation)
- **Test**
The performance is then assessed with the test data

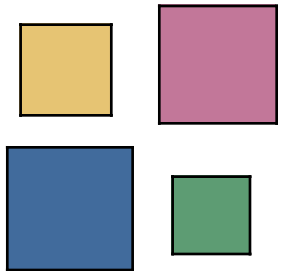
What about trees?

Model Structure = Model parameters

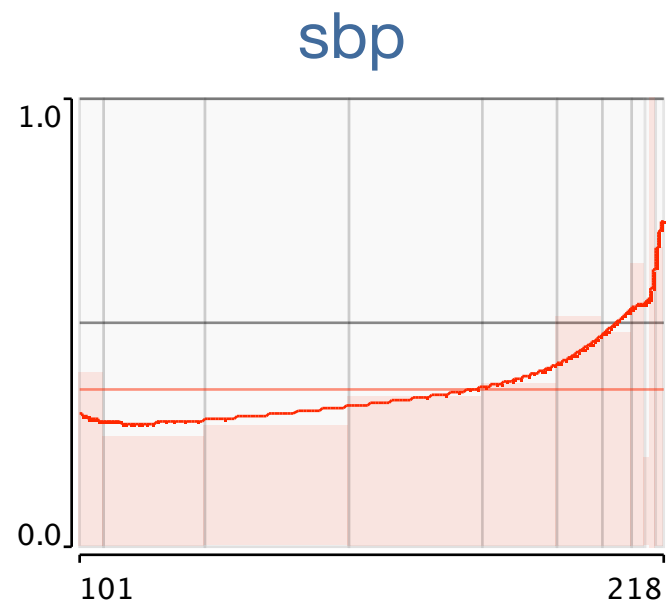


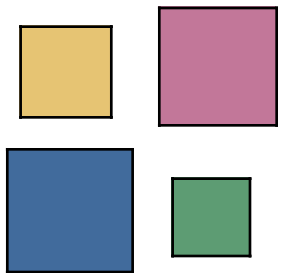
The Dataset

- 10 Variables, 462 Observations
- Target:
Coronary Heart Disease (chd), 34,63% = 160 cases
- Inputs:
 - continuous
 - sbp systolic blood pressure
 - tobacco cumulative tobacco (kg)
 - ldl low density lipoprotein cholesterol
 - adiposity
 - typea type-A behavior
 - obesity
 - alcohol current alcohol consumption
 - age age at onset
 - discrete
 - *omitted* famhist family history of heart disease (Present, Absent)

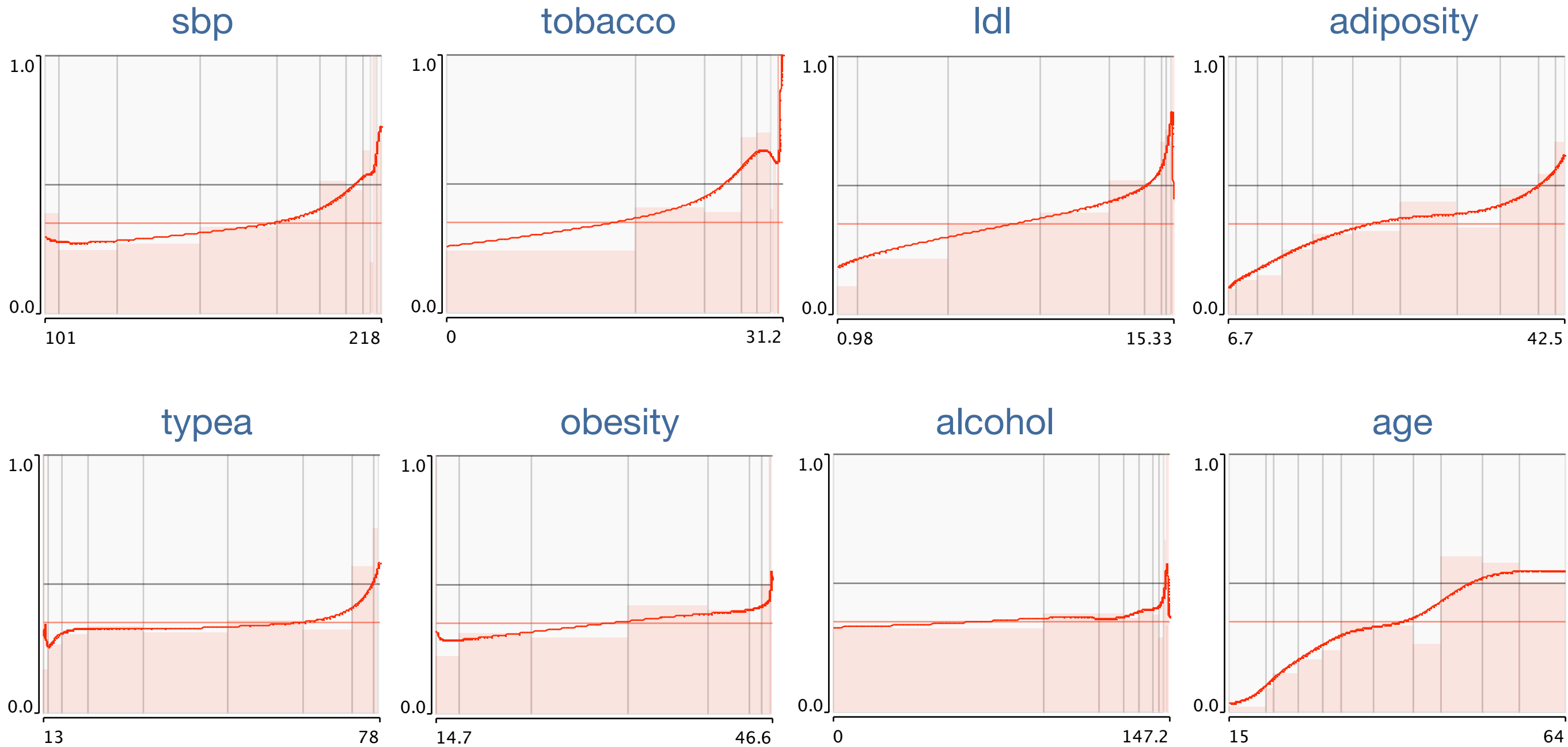


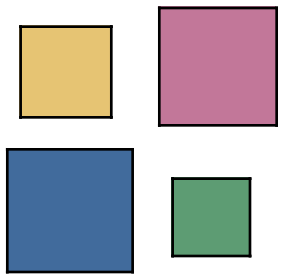
The Dataset: Univariate



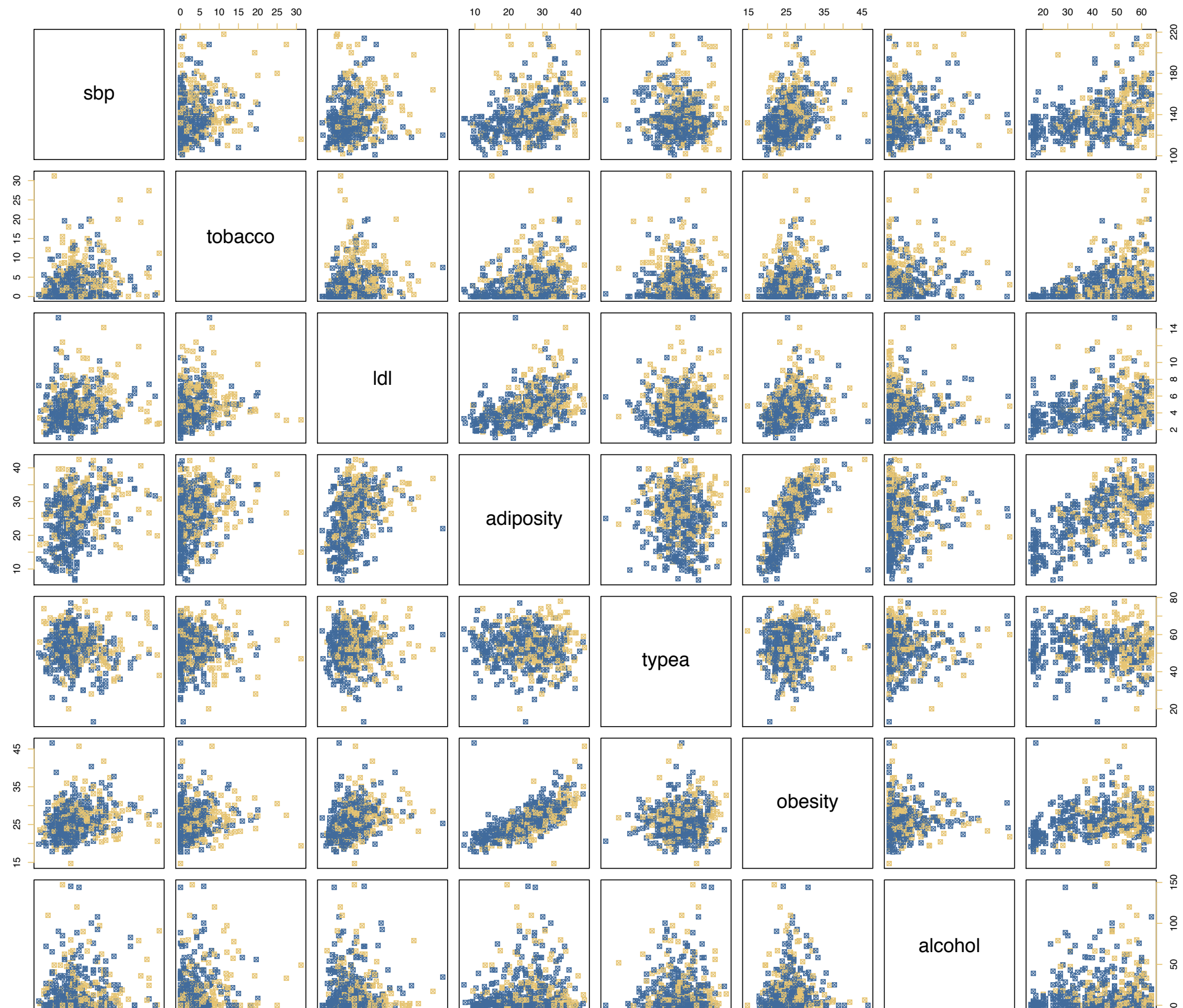


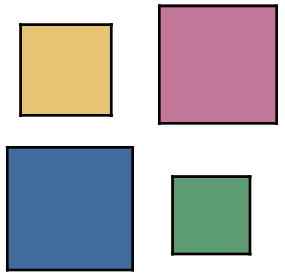
The Dataset: Univariate



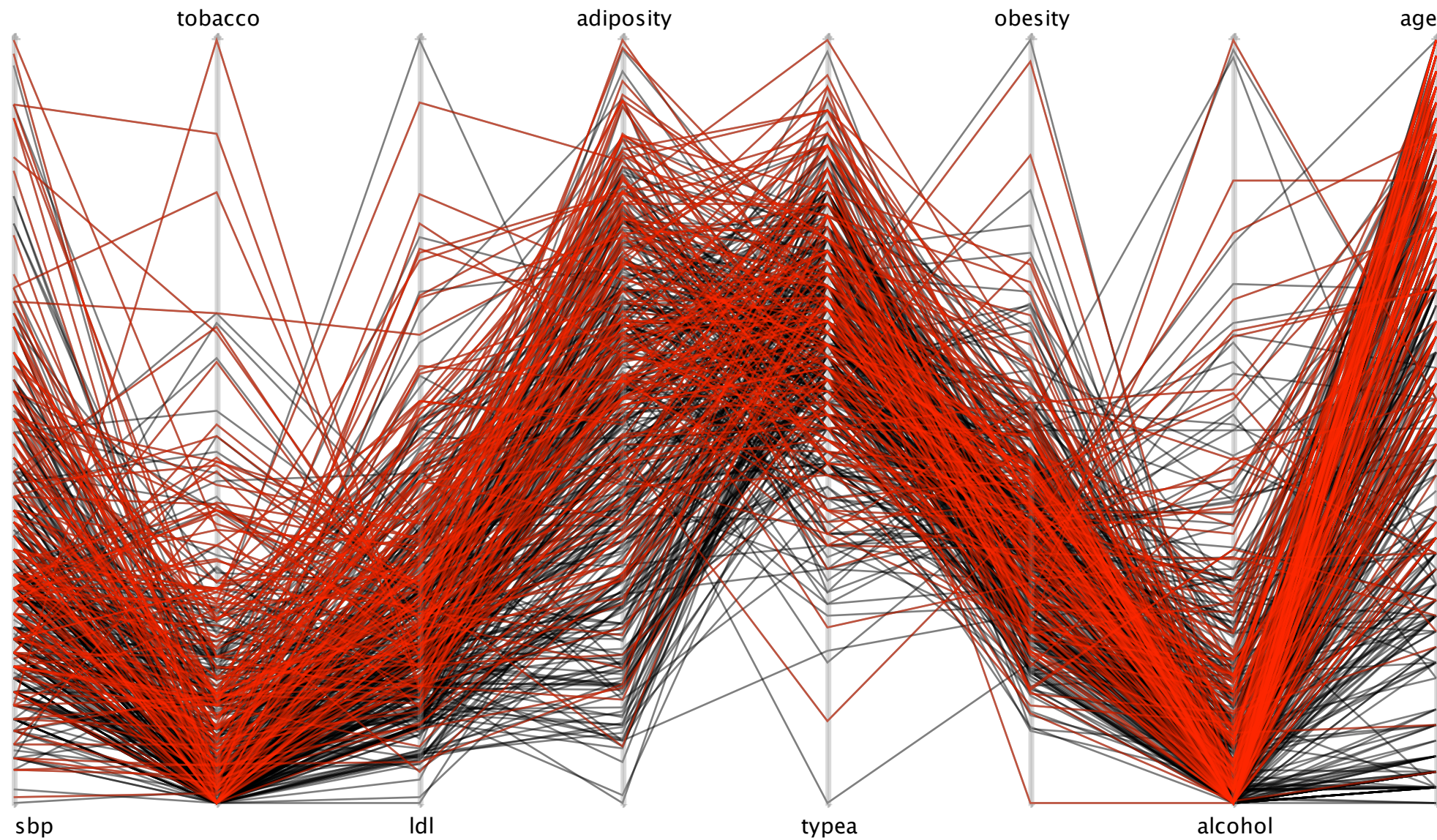


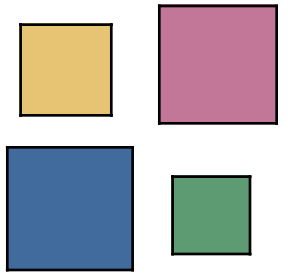
The Dataset: Bivariate





The Dataset: Multivariate





The Competitors: On 100 random samples

- Logistic Regression

```
glm(response~., data=dataTrain, family="binomial")
```

- Traditional CART

```
rpart(response ~ ., data=dataTrain,  
       parms=list(split='information'))
```

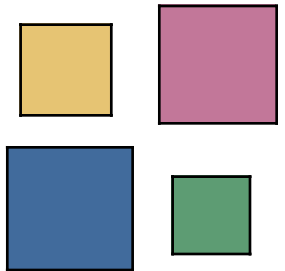
- Bagging

```
bagging(response~., data=dataTrain, nbagg=100)
```

- SVM

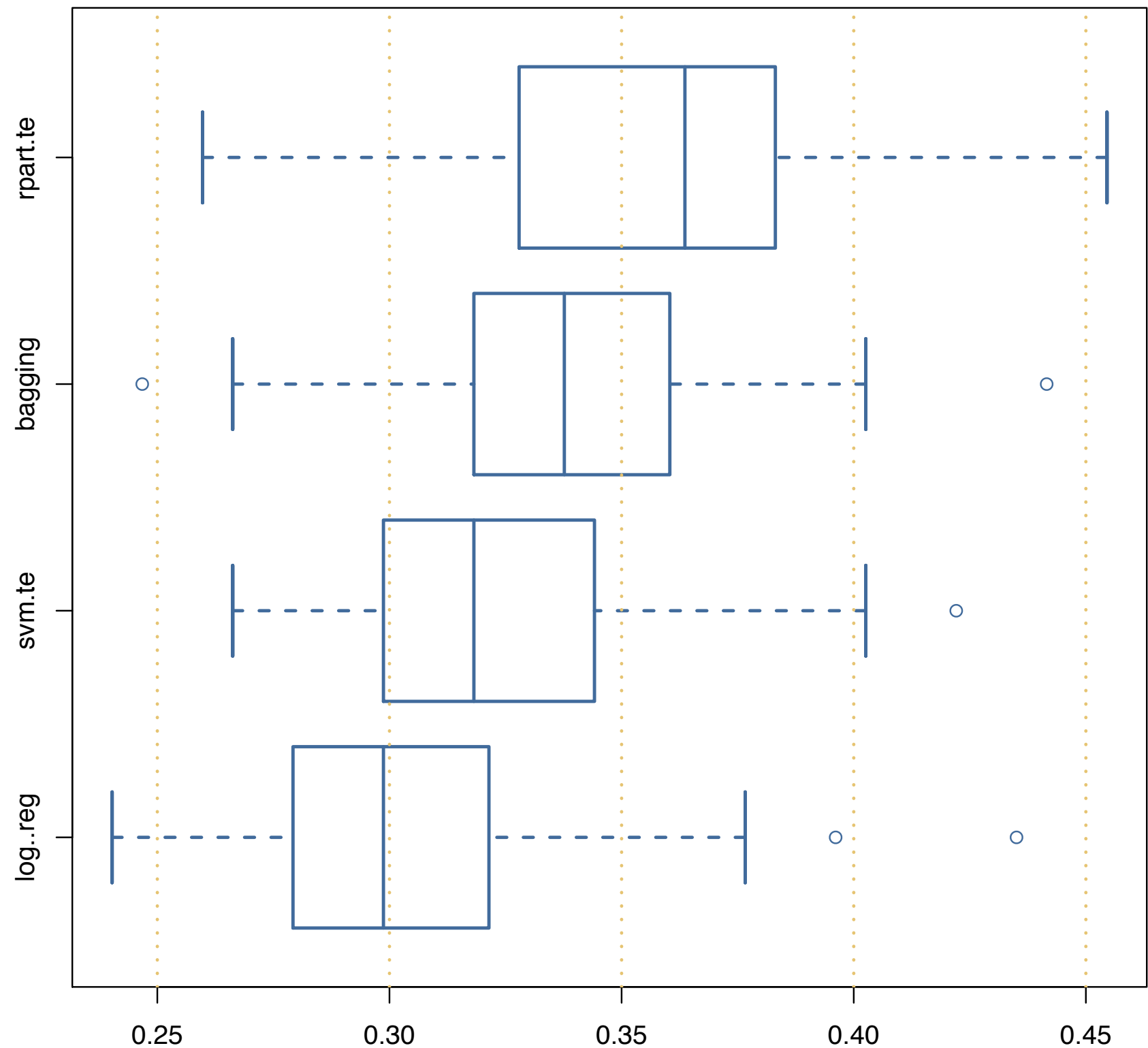
```
svm(response~., data=dataTrain)
```

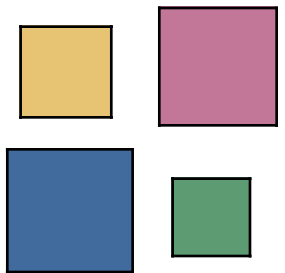
!! None of the methods has been further tuned !!



Competitors Results: Error Rates

- Trad. Trees
- Bagging
- Support Vektor Machines
- Log. Regression

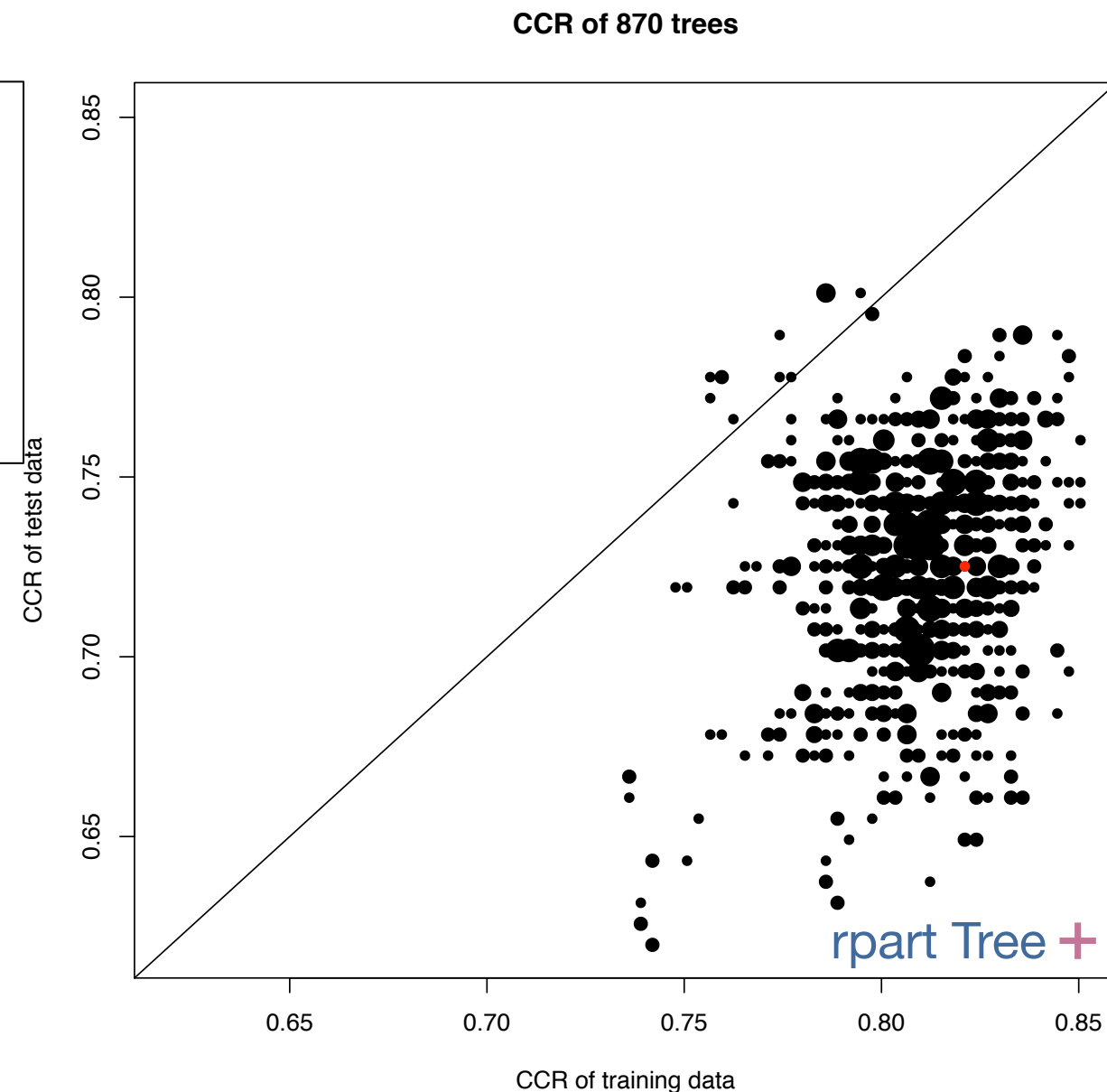
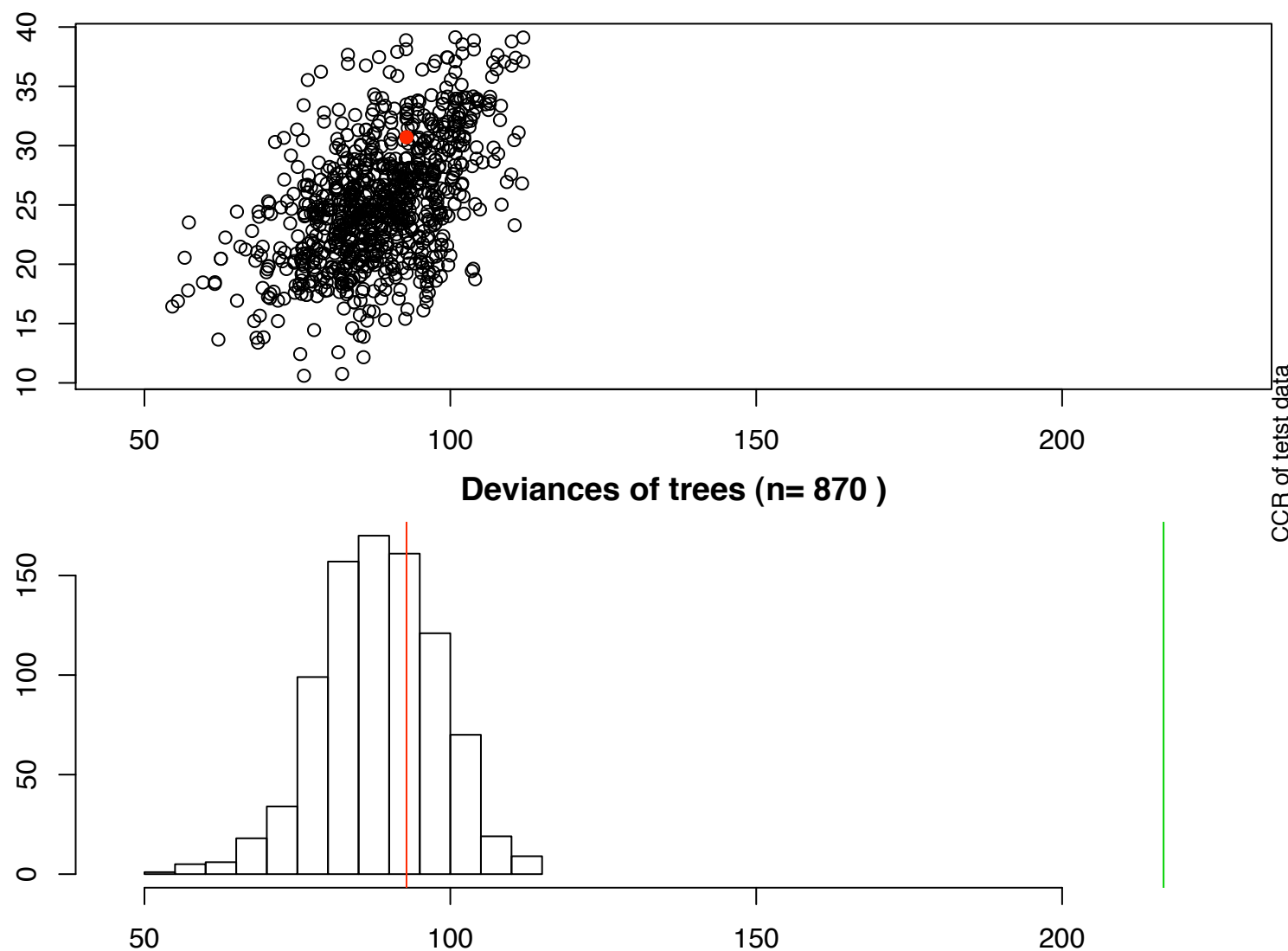


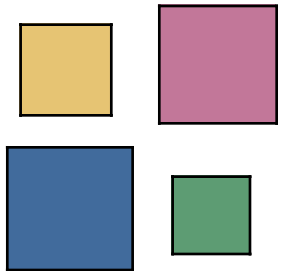


TWIX: Diagnostics

- For a given “Multitree” we can compare deviance and classification rate on training and test/validation data.

Example:





TWIX: Tree Selection

- The CCR (Correct Classification Rate) of the top TWIX trees are far better than those of greedy trees and most other classification methods

Quest:

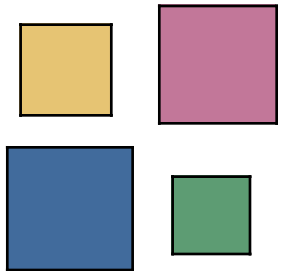
How to find the “best” trees from the validation data?

- Currently:

Sort trees according to

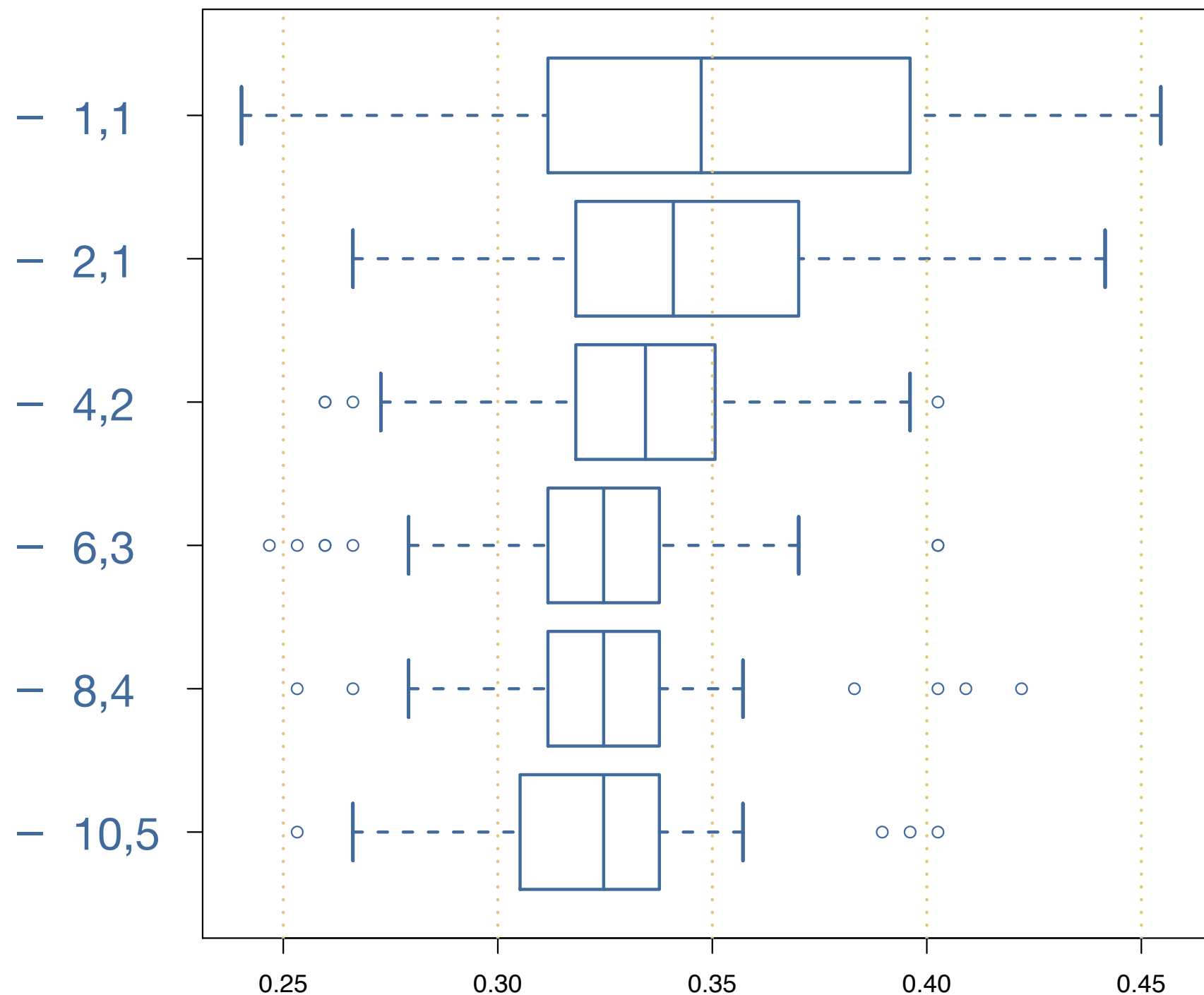
- training deviance
- test deviance
- training CCR
- test CCR

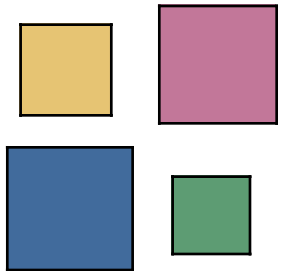
and pick the best!



TWIX: Results

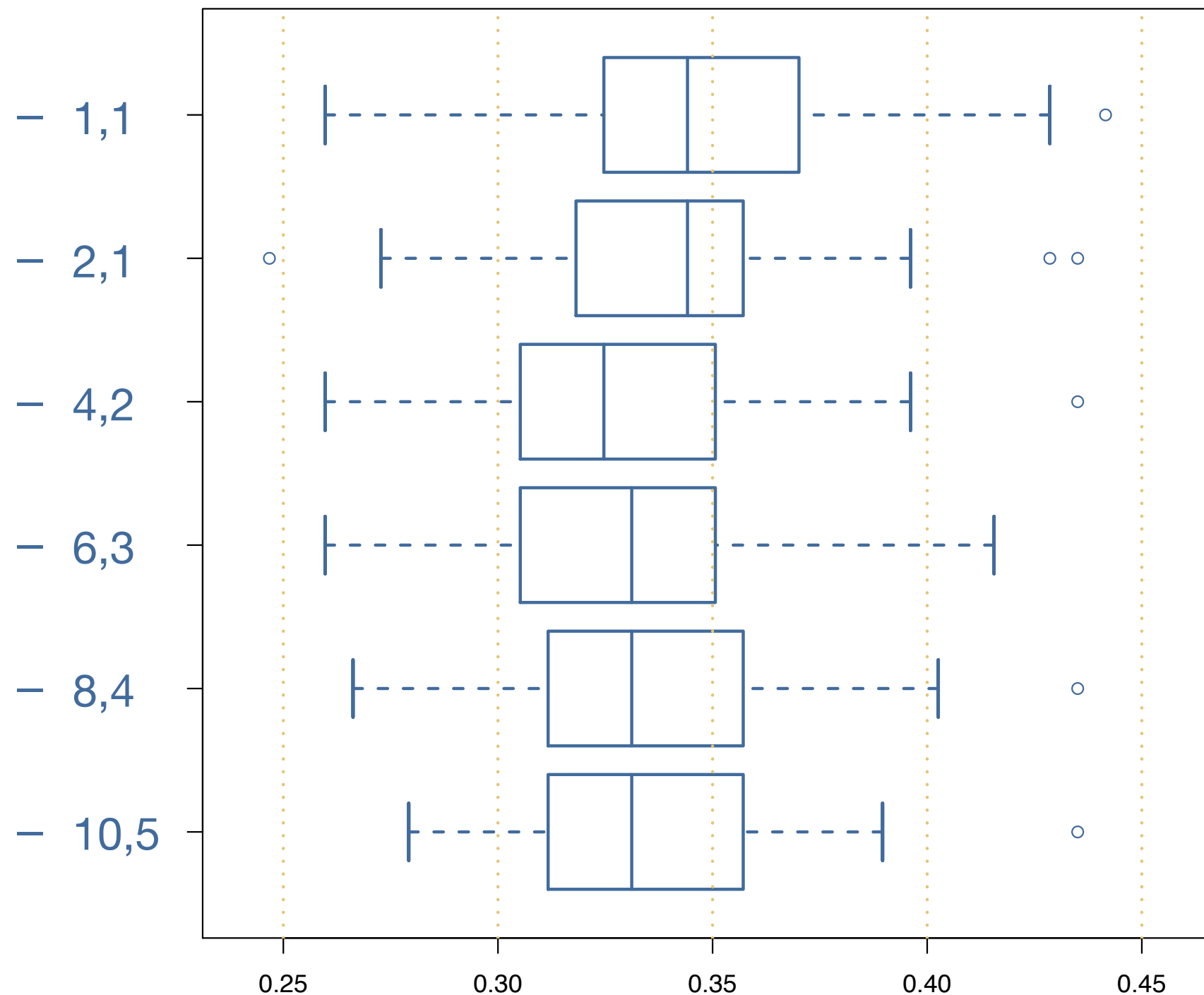
- Local Maxima across all variables (50 samples)

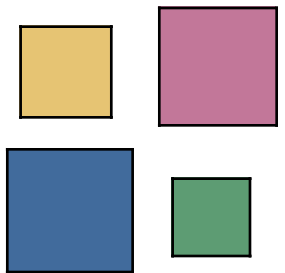




TWIX: Results

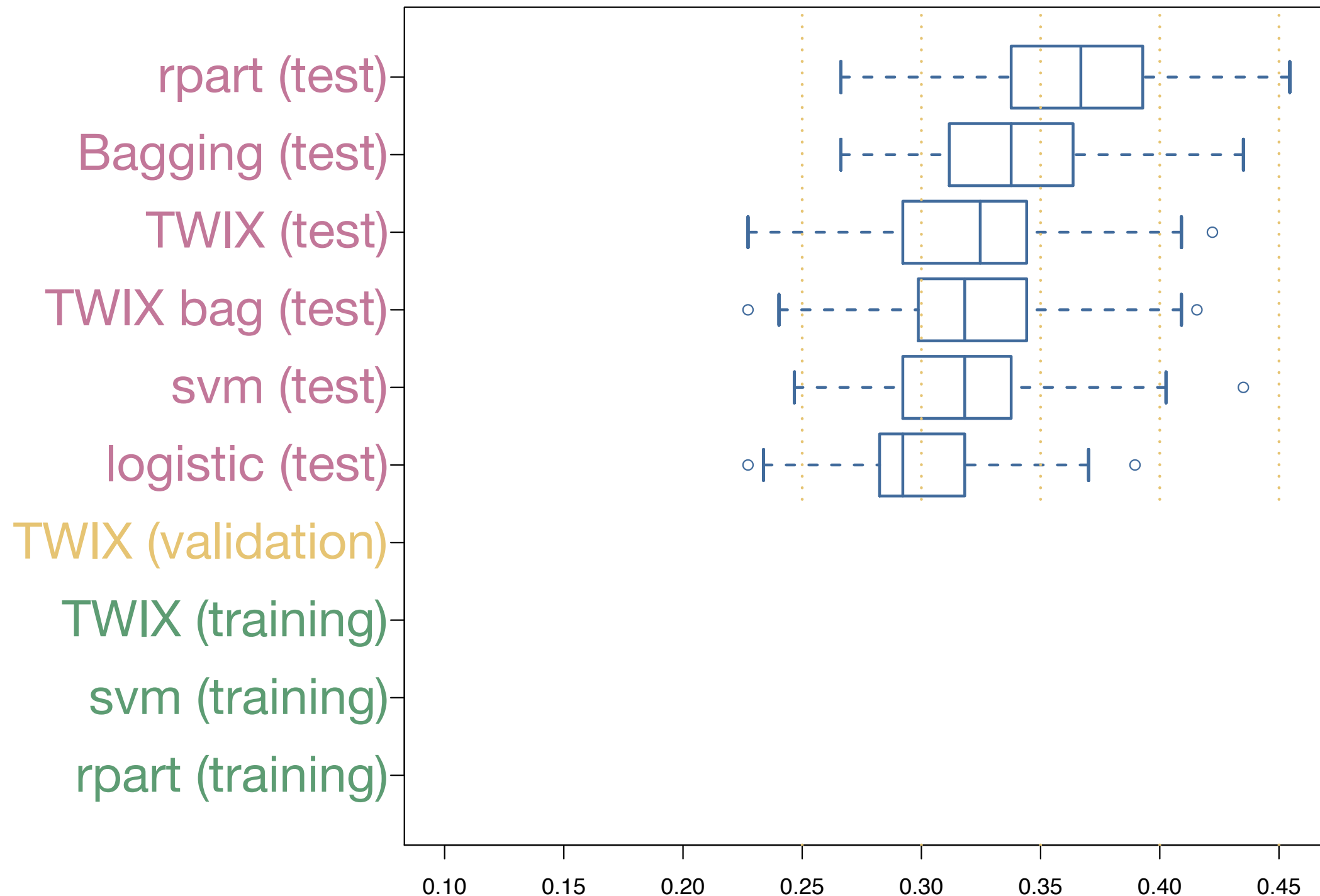
- Local Maxima, within all variables (50 samples)

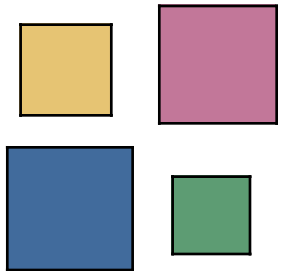




TWIX: Results

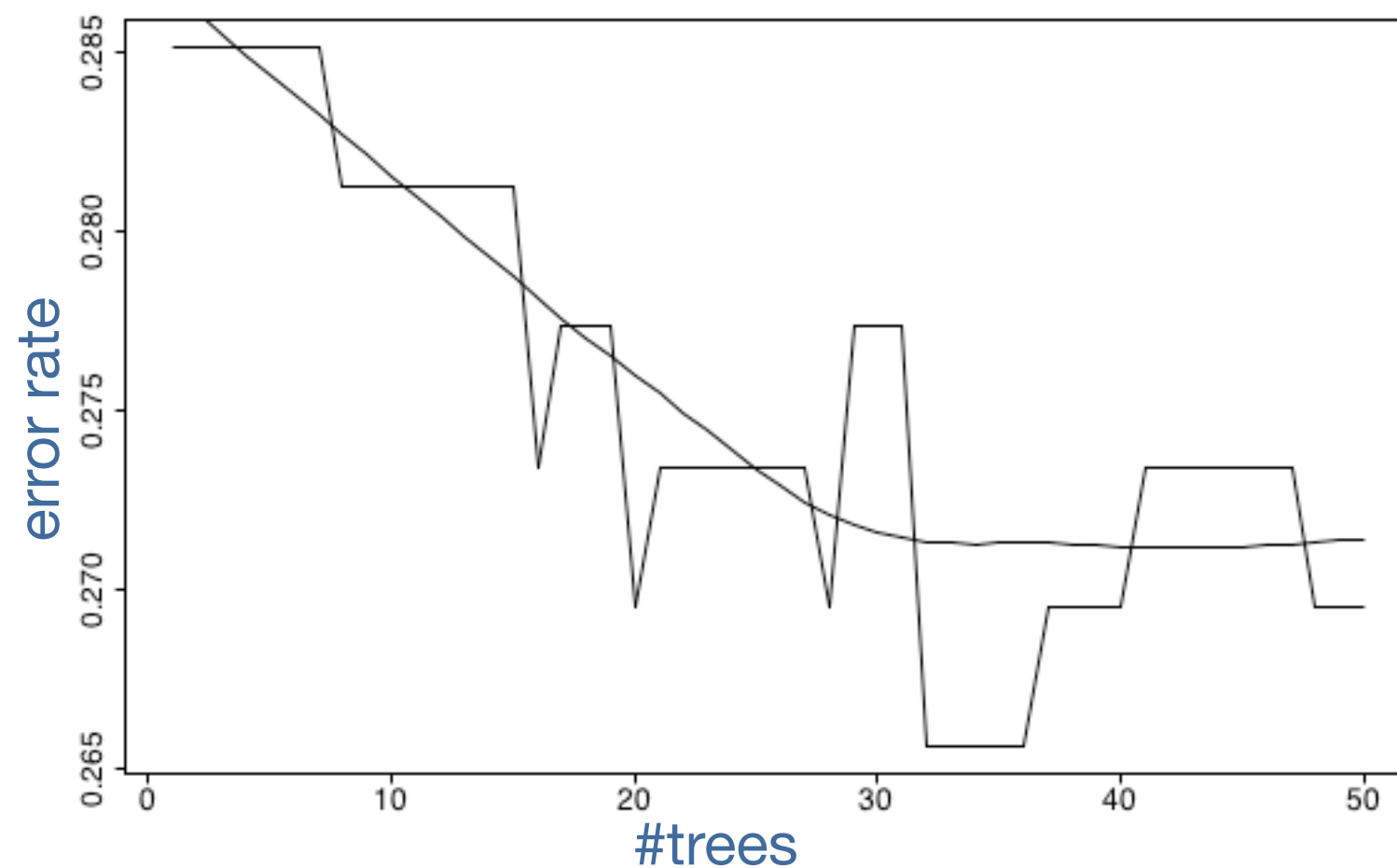
- 100 runs of a (10,5,2) tree

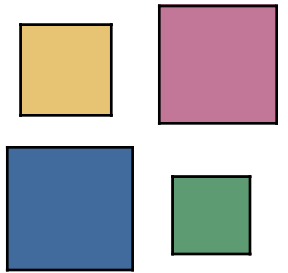




Bagged TWIX

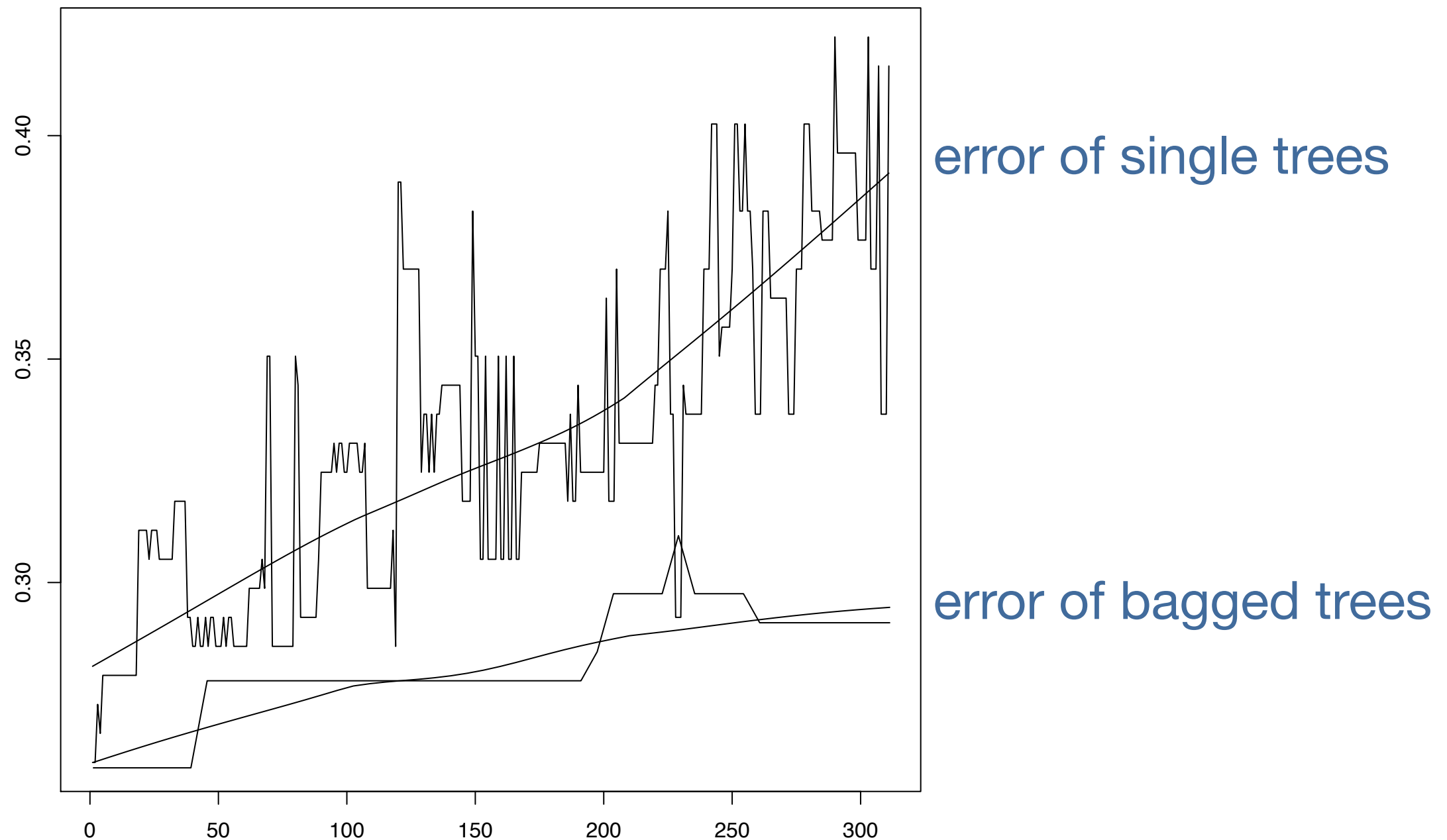
- What bagging should look like:

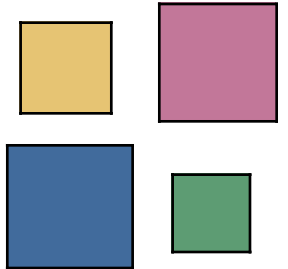




Bagged TWIX: Example

- Bagging TWIX trees does often not improve the CCR.





Conclusion

- For the South-African-Heart Data
 - Single TWIX-trees out-perform traditional tree and usually bagged trees
 - Bagged TWIX beats bagged trees and reaches top performance
 - TWIX gives good **single** alternative tree models
- Still much room for performance improvement
 - Stopping rules and pruning
 - Better tree selection
 - Improved selection of second best splits
 - More tests on more datasets
 - Better understanding of the tree-families
- Computational effort is high,
but parallel computing speeds up dramatically
- Complex methods are hard to implement and hard to test