



# Generic Frameworks for Nonparametric and Parametric Model Trees

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# Overview

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- Conditional inference trees
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- Summary

# Motivation: Trees

Breiman (2001, *Statistical Science*) distinguishes two cultures of statistical modeling.

- **Data models:** Stochastic models, typically parametric.
- **Algorithmic models:** Flexible models, data-generating process unknown.

**Example:** Recursive partitioning models dependent variable  $Y$  by “learning” a partition w.r.t explanatory variables  $Z_1, \dots, Z_l$ .

## Key features:

- Predictive power in nonlinear regression relationships.
- Interpretability (enhanced by visualization), i.e., no “black box” methods.

# Motivation: Leaves

**Typically:** Simple models for univariate  $Y$ , e.g., mean or proportion.

**Examples:** CART and C4.5 in statistical and machine learning, respectively.

**Problems:** For classical tree algorithms.

- No concept of “significance”, possibly biased variable selection.
- No complex (parametric) models in leaves.
- Many different tree algorithms for different types of data.

**Solutions:** Flexible generic frameworks based on statistical inference.

- Nonparametric: Employ only empirical distribution for inference.
- Parametric: Synthesis of parametric data models and algorithmic tree models.

# Motivation: Branches

**Base algorithm:** Growth of branches from the roots to the leaves of the tree typically follows a simple *recursive partitioning* algorithm.

- 1 Fit a (possibly very simple) model for the response  $Y$ .
- 2 Assess association of  $Y$  and each  $Z_j$ .
- 3 Split sample along the  $Z_{j^*}$  with strongest association: Choose breakpoint with highest improvement of the model fit.
- 4 Repeat steps 1–3 recursively in the subsamples until some stopping criterion is met.

**Generally:** Tree algorithms differ w.r.t. choice of model (1), association measure (2), split strategy (3) and stopping criterion or “pruning” strategy (4).

# Conditional inference trees

**Idea:** Fully nonparametric approach using a modern framework unifying classical nonparametric tests.

## Algorithm

- 1 Model: Nonparametric, empirical distribution of  $Y$ .
- 2 Association measure: Permutation test (i.e., conditional inference) for independence of  $Y$  and each  $Z_j$ .
- 3 Split strategy: Maximize two-sample contrast of  $Y$  along  $Z_j^*$ .
- 4 Stopping criterion: Significance of test in step 2.

**Note:** Both model and tests condition on the observed data.

# Conditional inference trees

**Model:** Predictions can be computed from any quantity of the empirical distribution of  $Y$  in the relevant node, e.g., the mean/median/etc. for numeric  $Y$ , proportion of “successes” for binary  $Y$ , Kaplan-Meier survivor function for censored  $Y$ , etc.

**Association measure:** Independence tests derived from general correlation of  $Y$  and  $Z_j$ .

$$t_j = \text{vec} \left( \sum_{i=1}^n h(Y_i) \cdot g(Z_{j,i}) \right),$$

with  $p$ -dimensional transformation  $g(\cdot)$  and  $q$ -dimensional influence function  $h(\cdot)$ .

# Conditional inference trees

**Test statistics:** Scalar standardized statistic based on conditional expectation  $\mu_j$  and covariance matrix  $\Sigma_j$  (given the data).

$$s_{\max}(t, \mu, \Sigma) = \max_k \left| \frac{(t - \mu)_k}{\sqrt{\Sigma_{k,k}}} \right|,$$
$$s_{\text{quad}}(t, \mu, \Sigma) = (t - \mu)\Sigma^+(t - \mu).$$

Under independence, all permutations of  $Y$  yield the conditional distribution of  $t_j$ . Taking expectations w.r.t. this yields:

$$\mu_j = E(t_j) = \text{vec} \left( \left( \sum_{i=1}^n g(Z_{j,i}) \right) E(h)^\top \right),$$
$$E(h) = n^{-1} \sum_i h(Y_i),$$



## Conditional inference trees

**Similarly:**  $pq \times pq$  conditional covariance matrix  $\Sigma_j$  computed from the permutation distribution under independence:

$$\begin{aligned}\Sigma_j = \text{Var}(t_j) &= \frac{n}{n-1} \text{Var}(h) \otimes \left( \sum_i g(Z_{j,i}) \otimes g(Z_{j,i})^\top \right) - \\ &\quad \frac{1}{n-1} \text{Var}(h) \otimes \left( \sum_i g(Z_{j,i}) \right) \otimes \left( \sum_i g(Z_{j,i}) \right)^\top, \\ \text{Var}(h) &= n^{-1} \sum_i (h(Y_i) - E(h)) (h(Y_i) - E(h))^\top,\end{aligned}$$

where  $\otimes$  denotes the Kronecker product.

# Conditional inference trees

**Significance:** Various approaches can be used to assess the significance of the test statistic  $s(t_j, \mu_j, \Sigma_j)$ :

- Exact: Direct computation of the statistic for all permutations. Typically burdensome.
- Approximate: Compute statistics for a sufficiently large number of permutations, drawn using Monte Carlo methods.
- Asymptotic: Compute the conditional asymptotic distribution of  $s$  based on the asymptotic conditional distribution of  $t_j$ .  
 $t_j \sim \mathcal{N}(\mu_j, \Sigma_j)$ .

# Conditional inference trees

**Choice of transformations:** Based on scale of  $Y$  and  $Z_j$  and type of dependence.

- Categorical: Indicator functions for all  $C$  categories  
 $h(y) = (I_1(y), \dots, I_C(y))^T$ .
- Numeric:
  - Location:  $h(y) = y$  or  $h(y) = \text{rank}(y)$ .
  - Scatter:  $h(y) = (y - \bar{y})^2$  or  $h(y) = (\text{rank}(y) - (n + 1)/2)^2$ .
  - Threshold:  $h(y) = I(y > \zeta)$ .
- Survival: Log rank scores

**Special cases:** Choice of  $h(\cdot)$  and analogously  $g(\cdot)$  yields many classical tests as special cases. Wilcoxon-Mann-Whitney, Spearman, Pearson's  $\chi^2$ , Cochran-Armitage, log rank, Kruskal-Wallis, and many more.

# Conditional inference trees

**Split strategy:** Maximize two-sample contrast of  $Y$  along  $Z_j$ .

- Employ threshold transformation  $g(Z_j) = I(Z_j > \zeta)$  for all possible thresholds  $\zeta$ .
- Choose split  $\zeta^*$  that maximizes the associated test statistic.

**Stopping criterion:** Non-significance of Bonferroni-adjusted  $p$  values from permutation tests.

# Application: Pima Indians diabetes

**Task:** Classification of diabetes in Pima Indian women.

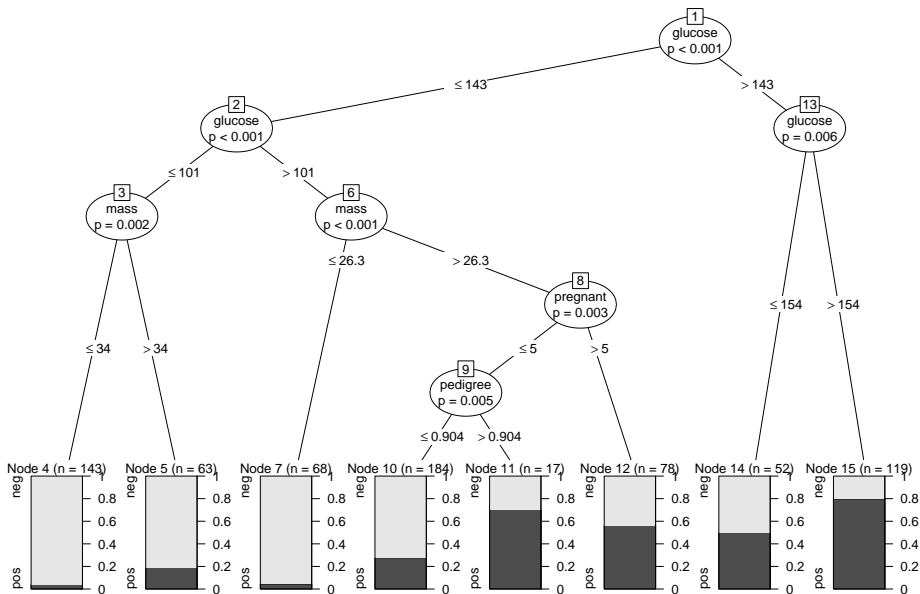
**Source:** Asuncion & Newman (2007), UCI Repository of Machine Learning Databases.

<http://www.ics.uci.edu/~mllearn/MLRepository.html>.

**Response:** Test result for *diabetes* (positive/negative).

**Explanatory variables:** Plasma *glucose* concentration, number of times *pregnant*, diastolic blood *pressure* (mm Hg), body *mass* index, diabetes *pedigree* function, *age* (in years).

# Application: Pima Indians diabetes



## Application: Pima Indians diabetes

**Inference:** In each node, an asymptotic permutation test for independence of *diabetes* ( $Y$ ) and each of the variables *glucose*,  $\dots$ , *age* ( $Z_1, \dots, Z_6$ ) is carried out.

**Transformations:** Indicator function for categorical variables (response) and identity for numeric variables (all regressors).

$$\begin{aligned}h(Y_i) &= (I_{\text{pos}}(Y_i), I_{\text{neg}}(Y_i))^{\top}, \\g(Z_{j,i}) &= Z_{j,i}.\end{aligned}$$

**Interpretation:** Corresponds to two-sample  $t$  test with pooled one-sample standard deviation.

# Model-based recursive partitioning

**Idea:** More complex models for multivariate  $Y$ , e.g., multivariate normal model, regression models, etc.

## Goal:

- Synthesis of parametric data models and algorithmic tree models.
- Fitting local models by partitioning of the sample space.

## Algorithm

- ① Model: Parametric model for  $Y$  with additive objective function.
- ② Association measure: Parameter instability tests.
- ③ Split strategy: Model segmentation.
- ④ Stopping criterion: Significance of test in step 2.



# Model-based recursive partitioning: Estimation

**Models:**  $\mathcal{M}(Y, \theta)$  with (potentially) multivariate observations  $Y \in \mathcal{Y}$  and  $k$ -dimensional parameter vector  $\theta \in \Theta$ .

**Parameter estimation:**  $\hat{\theta}$  by optimization of objective function  $\Psi(Y, \theta)$  for  $n$  observations  $Y_i$  ( $i = 1, \dots, n$ ):

$$\hat{\theta} = \operatorname{argmin}_{\theta \in \Theta} \sum_{i=1}^n \Psi(Y_i, \theta).$$

**Special cases:** Maximum likelihood (ML), weighted and ordinary least squares (OLS and WLS), quasi-ML, and other M-estimators.

**Central limit theorem:** If there is a true parameter  $\theta_0$  and given certain weak regularity conditions,  $\hat{\theta}$  is asymptotically normal with mean  $\theta_0$  and sandwich-type covariance.

# Model-based recursive partitioning: Estimation

**Estimating function:**  $\hat{\theta}$  can also be defined in terms of

$$\sum_{i=1}^n \psi(Y_i, \hat{\theta}) = 0,$$

where  $\psi(Y, \theta) = \partial\Psi(Y, \theta)/\partial\theta$ .

**Idea:** In many situations, a single global model  $\mathcal{M}(Y, \theta)$  that fits **all**  $n$  observations cannot be found. But it might be possible to find a partition w.r.t. the variables  $Z = (Z_1, \dots, Z_l)$  so that a well-fitting model can be found locally in each cell of the partition.

**Tool:** Assess parameter instability w.r.t to partitioning variables  $Z_j \in \mathcal{Z}_j$  ( $j = 1, \dots, l$ ).

# Model-based recursive partitioning: Tests

Generalized M-fluctuation tests capture instabilities in  $\hat{\theta}$  for an ordering w.r.t  $Z_j$ .

**Basis:** Empirical fluctuation process of cumulative deviations w.r.t. to an ordering  $\sigma(Z_{ij})$ .

$$W_j(t, \hat{\theta}) = \hat{V}^{-1/2} n^{-1/2} \sum_{i=1}^{\lfloor nt \rfloor} \psi(Y_{\sigma(Z_{ij})}, \hat{\theta}) \quad (0 \leq t \leq 1)$$

**Functional central limit theorem:** Under parameter stability  $W_j(\cdot, \hat{\theta}) \xrightarrow{d} W^0(\cdot)$ , where  $W^0$  is a  $k$ -dimensional Brownian bridge.

# Model-based recursive partitioning: Tests

**Test statistics:** Scalar functional  $\lambda(W_j)$  that captures deviations from zero.

**Null distribution:** Asymptotic distribution of  $\lambda(W^0)$ .

**Special cases:** Class of test encompasses many well-known tests for different classes of models. Certain functionals  $\lambda$  are particularly intuitive for numeric and categorical  $Z_j$ , respectively.

**Advantage:** Model  $\mathcal{M}(Y, \hat{\theta})$  just has to be estimated once. Empirical estimating functions  $\psi(Y_i, \hat{\theta})$  just have to be re-ordered and aggregated for each  $Z_j$ .

# Model-based recursive partitioning: Tests

**Splitting numeric variables:** Assess instability using sup $LM$  statistics.

$$\lambda_{\text{sup}LM}(W_j) = \max_{i=\underline{i}, \dots, \bar{i}} \left( \frac{i}{n} \cdot \frac{n-i}{n} \right)^{-1} \left\| W_j \left( \frac{i}{n} \right) \right\|_2^2.$$

**Interpretation:** Maximization of single shift  $LM$  statistics for all conceivable breakpoints in  $[\underline{i}, \bar{i}]$ .

**Limiting distribution:** Supremum of a squared,  $k$ -dimensional tied-down Bessel process.

# Model-based recursive partitioning: Tests

**Splitting categorical variables:** Assess instability using  $\chi^2$  statistics.

$$\lambda_{\chi^2}(W_j) = \sum_{c=1}^C \frac{n}{|I_c|} \left\| \Delta_{I_c} W_j \left( \frac{i}{n} \right) \right\|_2^2$$

**Feature:** Invariant for re-ordering of the  $C$  categories and the observations within each category.

**Interpretation:** Captures instability for split-up into  $C$  categories.

**Limiting distribution:**  $\chi^2$  with  $k \cdot (C - 1)$  degrees of freedom.

# Model-based recursive partitioning: Segmentation

**Goal:** Split model into  $b = 1, \dots, B$  segments along the partitioning variable  $Z_j$  associated with the highest parameter instability. Local optimization of

$$\sum_b \sum_{i \in I_b} \Psi(Y_i, \theta_b).$$

$B = 2$ : Exhaustive search of order  $O(n)$ .

$B > 2$ : Exhaustive search is of order  $O(n^{B-1})$ , but can be replaced by dynamic programming of order  $O(n^2)$ . Different methods (e.g., information criteria) can choose  $B$  adaptively.

**Here:** Binary partitioning.

# Model-based recursive partitioning: Pruning

**Pruning:** Avoid overfitting.

**Pre-pruning:** Internal stopping criterion. Stop splitting when there is no significant parameter instability.

**Post-pruning:** Grow large tree and prune splits that do not improve the model fit (e.g., via cross-validation or information criteria).

**Here:** Pre-pruning based on Bonferroni-corrected  $p$  values of the fluctuation tests.



# Application: Pima Indians diabetes

**Task:** Reconsider classification of diabetes in Pima Indian women.

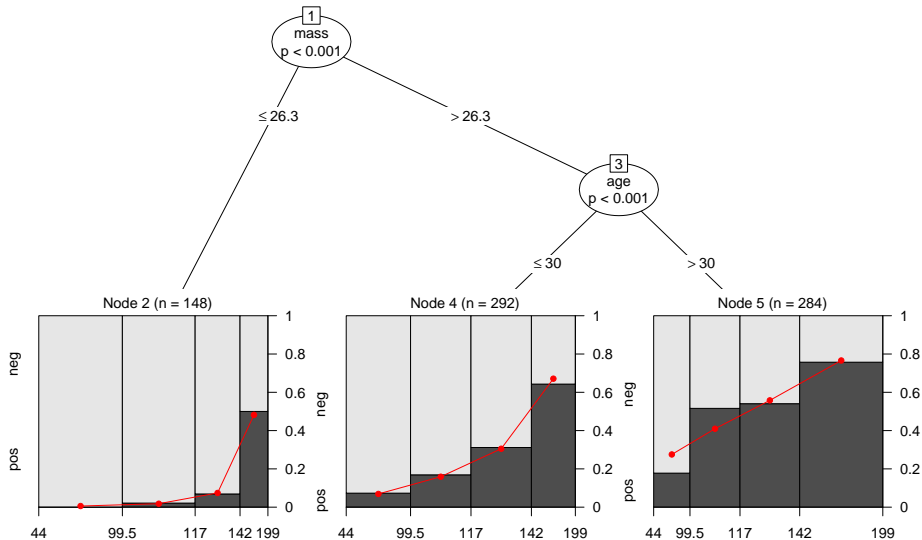
**Idea:**

- Variable *glucose* occurred in many splits in conditional inference tree.
- More parsimonious model may be possible if *glucose* is employed as continuous regressor rather than partitioning variable.

**Model:** Logistic regression of *diabetes* on *glucose*.

**Partitioning variables:** All remaining variables.

# Application: Pima Indians diabetes



# Application: Pima Indians diabetes

## Model-based recursive partitioning:

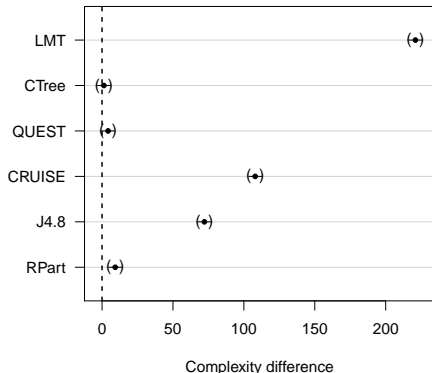
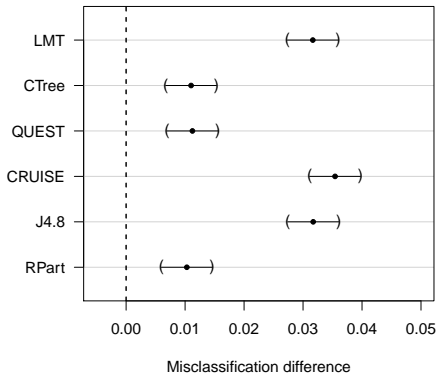
- Coefficient estimates for regressors.
- Parameter instability tests for partitioning variables (bold = significant at adjusted 5% level, underlined = smallest  $p$  value).

	Regressors		Partitioning variables				
	(const.)	glucose	pregnant	pressure	mass	pedigree	age
1	-5.608	0.039	<b>26.49</b>	8.67	<u>43.41</u>	<b>21.04</b>	<b>39.47</b>
2	-10.999	0.065	8.40	4.50	<u>9.31</u>	4.02	4.53
3	-4.958	0.037	<b>24.80</b>	7.63	9.05	<b>19.29</b>	<u>33.71</u>
4	-6.573	0.045	3.46	3.77	5.09	<u>7.20</u>	6.20
5	-3.319	0.027	6.24	1.74	13.34	<u>14.89</u>	10.24

# Application: Pima Indians diabetes

**Benchmark:** Compare predictive performance (misclassification rate) and model complexity (number of parameters/splits) of model-based recursive partitioning with other tree algorithms.

**Setup:** 250 bootstrap samples and out-of-bag misclassification rate.



# Application: Treatment effect for chronic disease

**Task:** Identify groups of chronic disease patients with different treatment effects.

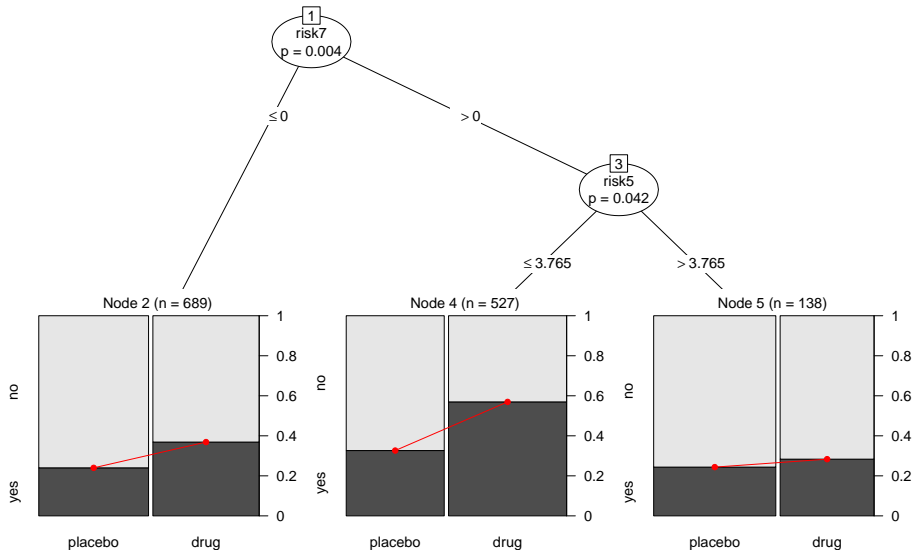
**Source:** Anonymized data from consulting project.

**Model:** Logistic regression.

- Response: Improvement (yes/no) of chronic disease after treatment over several weeks.
- Regressor: Treatment (active drug/placebo).
- Partitioning variables: 11 variables that describe disease status of patients. Lower values indicate more severe forms of the disease.

**Result:** Treatment is most effective for certain intermediate forms of the disease.

# Application: Treatment effect for chronic disease



# Software

All methods are implemented in the R system for statistical computing and graphics. Freely available under the GPL (General Public License) from the Comprehensive R Archive Network:

- Trees/recursive partytioning: `ctree()` in **party** for conditional inference trees, and `mob()` in **party** for model-based recursive partitioning.
- Inference: `independence_test()` in **coin** for permutation tests for independence, and `gefp()` in **strucchange** for structural change tests.

<http://www.R-project.org/>  
<http://CRAN.R-project.org/>

# Summary

## Conditional inference trees:

- Tree models based on nonparametric statistical inference.
- Based on modern class of permutation tests for independence.
- Aims to capture dependence patterns by recursive partitioning.
- Can be adapted to dependent and explanatory variables of arbitrary types, by employing suitable transformations/influence functions.
- Flexible implementation freely available: New transformations/influence functions can be simply plugged in.



# Summary

## Model-based recursive partitioning:

- Synthesis of classical parametric data models and algorithmic tree models.
- Based on modern class of parameter instability tests.
- Aims to minimize clearly defined objective function by greedy forward search.
- Can be applied general class of parametric models.
- Alternative to traditional means of model specification, especially for variables with unknown association.
- Object-oriented implementation freely available: Extension for new models requires some coding but not too extensive if interfaced model is well designed.

# References

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