Randomised Decision Forests for Regression

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References:

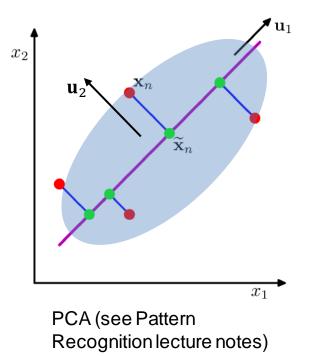
A. Criminisi et al., Decision forests, an unified framework, Foundations and Trends in Computer Graphics and Vision, 7:2-3. Breiman L, Random Forests. Machine Learning, 45 (1), pp 5-32, 2001.

Regression forests

- We discuss the use of random decision forests for the probabilistic estimation of continuous variables.
- Regression forests are used for the nonlinear regression of dependent variables given independent input.
- Both input and output may be multi-dimensional.
- The output can be a point estimate or a full probability density function.
- Regression forests are less popular than their classification counterpart.
- The main difference is that the output label to be associated with an input data is continuous.
- Therefore, the training labels are continuous.
- Consequently the objective function has to be adapted appropriately.
- Regression forests share many of the advantages of classification forests such as <u>efficiency</u> and <u>flexibility</u>.

Nonlinear regression

- Given a set of noisy input data and associated continuous measurements, least squares techniques (closely related to principal component analysis) can be used to fit a linear regressor which minimizes some error computed over all training points.
- Under this model, given a new test input the corresponding output can be efficiently estimated.
- The limitation of this model is in its linear nature, when we know that most natural phenomena have nonlinear behaviour.
- Another well known issue with linear regression techniques is their sensitivity to input noise.
- In geometric computer vision, a popular technique for achieving robust regression via randomization is RANSAC. select few rehable in-layer does points for modelling.



split in recursions

Notations

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Input data vector: \mathbf{v} = (x_1, \dots, x_d) \in \mathbb{R}^d
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Output/label: $\mathbf{y} \in \mathcal{Y} \subset \mathbb{R}^n$

Subset of training points reaching node j: S_j

Subset of points going to the left child node: $\mathcal{S}_{i}^{\mathrm{L}}$

Subset of points going to the right child node: $\mathcal{S}_j^{\mathtt{R}}$, s.t. $\mathcal{S}_j = \mathcal{S}_j^{\mathtt{L}} \cup \mathcal{S}_j^{\mathtt{R}}$

Node test parameters: $\theta \in \mathcal{T}$

Node weak learner: $h(\mathbf{v}, \boldsymbol{\theta}_j) \in \{\text{true}, \text{false}\}\$

Node objective function: $I = I(S_j, \theta)$

Stopping criteria: e.g. max tree depth = D

Leaf predictor model: $p(y|\mathbf{v})$

Forest size i.e. number of trees: T

Ensemble model: $p(y|\mathbf{v}) = \frac{1}{T} \sum_{t}^{T} p_t(y|\mathbf{v})$

Decision forest model for regression

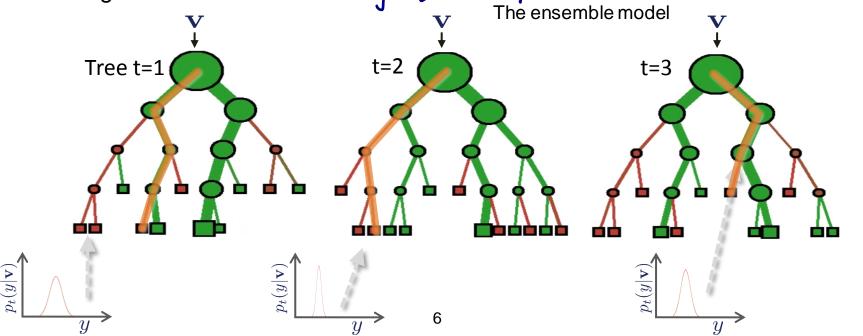
- The regression task: Given a labelled training set learn a general mapping which associates previously unseen independent test data with their correct continuous prediction.
- Like classification the regression task is inductive, with the main difference being the continuous nature of the output.
- In general, a training point is denoted as a labelled pair (v,y). A previously unseen test input (unavailable during training) is shown as a light gray circle on the x axis.

 Legression: Supervised (parring (given training sets)



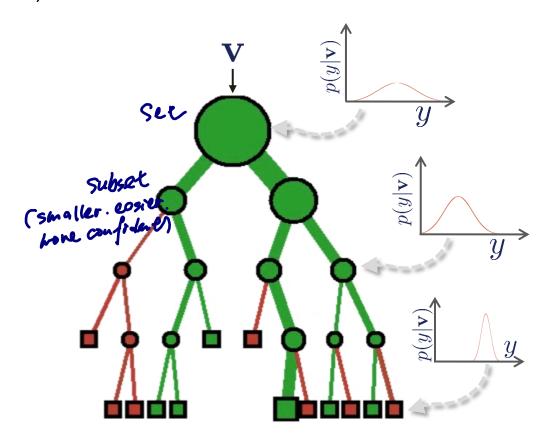
Decision forest model for regression

- Formally, given a multi-variate input v we wish to associate a continuous multi-variate label $\mathbf{y} \in \mathcal{Y} \subset \mathbb{R}^n$
- More generally, we wish to estimate the probability density function p(y|v).
- As usual the input is represented as a multi-dimensional feature response vector $\mathbf{v}=(x_1,\cdots,x_d)\in\mathbb{R}^d$
- A regression forest is a collection of randomly trained regression trees.
- Like in classification it can be shown that a forest generalizes better than a single over-trained tree. Single tree: overfitting



Decision forest model for regression

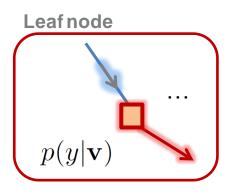
 A regression tree splits a complex nonlinear regression problem into a set of smaller problems which can be more easily handled by simpler models (e.g., linear ones).





The prediction model

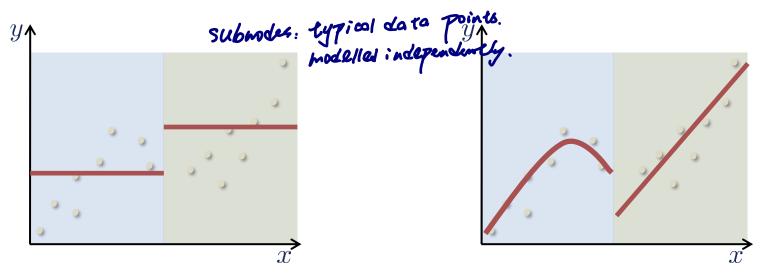
- The first job of a decision tree is to decide which branch to direct the incoming data to.
- When the data reaches a terminal node then that leaf needs to make a prediction.
- The actual form of the prediction depends on the prediction model.
- In classification we have used the pre-stored empirical class posterior as model.
- In regression forests we have a few alternatives.



The prediction model

- For instance we could use a polynomial function of a subspace of the input
 v.
- In the low dimensional example a generic polynomial model is given below.
- This simple model captures both the linear and constant models.

Examples of leaf (predictor) models



Predictor model: constant

$$y = const$$

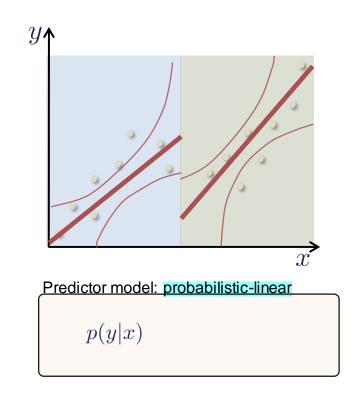
Predictor model: polynomial (/ thear/modises)

$$y=\sum_{i=0}^n w_i x^i$$
 (note: linear for n=1, constant for n=0)



The prediction model

- In this we are interested in output confidence as well as its actual value.
- Thus for prediction we can use a probability density function over the continuous variable y.
- So, given the t-th tree in a forest and an input point v, the associated leaf output takes the form p_t(y/v).
- In the low-dimensional example, we assume an underlying linear model and each leaf yields the conditional p(y|x).



The ensemble model

Like in classification, the forest output is the average of all tree outputs

$$p(\mathbf{y}|\mathbf{v}) = \frac{1}{T} \sum_{\mathbf{var}}^{T} p_{\mathbf{v}}(\mathbf{y}|\mathbf{v}) \quad \text{input} \quad$$

- Randomness model:
 - As in classification here we use a randomized node optimization model.
 - Therefore, the amount of randomness is controlled during training by the parameter $\rho = |\mathcal{T}_i|$.
 - The random subsets of split parameters T_j can be generated on the fly when training the j-th node.

The training objective function

- Forest training happens by optimizing an energy over a training set S_0 of data and associated continuous labels.
- Training a split node j happens by optimizing the parameters of its weak learner:

$$oldsymbol{ heta}_j^* = rg \max_{oldsymbol{ heta}_j \in \mathcal{T}_j} I_j$$

- Now, the main difference between classification and regression forest is in the form of the objective function I.
- We employ a continuous formulation of information gain.

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London Information gain on continuous distribution

- Entropy and information gain can also be defined for continuous-valued labels and distributions.
- The definition of the information gain remains the same but this time, instead of using the Shannon entropy, the differential entropy is used

$$H(S) = -\int_{y \in \mathcal{Y}} p(y) \log(p(y)) dy$$

Here *y* is a continuous label and *p* is the probability density function estimated from the training points in the set *S*.

- From a practical point of view, in the discrete case, the distribution p(c) was defined as the empirical distribution (i.e. class histogram) computed from the training set.
- Similarly in the continuous distribution p(y) can be defined either using parametric distributions or non-parametric methods.

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London Information gain on continuous distribution

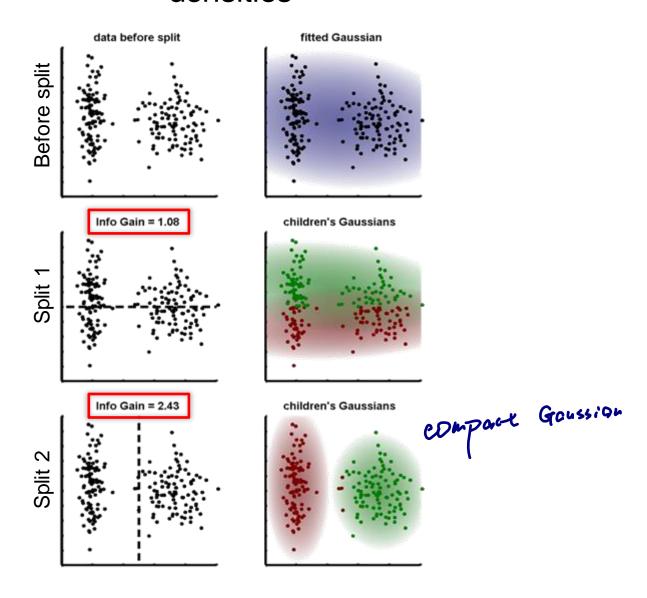
- One of the most popular choice in various applications is to use Gaussianbased models to approximate the density p(y) due to their simplicity.

where
$$\Lambda(S)$$
 is the data covariance matrix in differentiation: avoid high complexity in differentiations

- A toy example in the next slide illustrates the role of the continuous information gain in training.
- /->Information gain for discrete categorical distribution is for classification, continuous distribution of y for regression, continuous distribution of x for clustering.
- This time we wish to cluster similar points according to their features (again, depicted as the coordinates of a 2D space).
- Given an arbitrary input data point we wish the tree to predict its associated cluster.

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London Atoy example, Information gain on continuous, parametric densities



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LorAlton example, Information gain on continuous, parametric densities

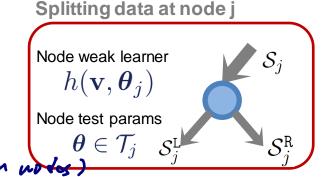
- Fitting a Gaussian to the entire initial set S produces the density shown in blue, which has a high differential entropy.
- Splitting the data horizontally produces two largely overlapping and slightly smaller Gaussians (in red and green).
- The large overlap indicates a suboptimal separation and is associated with a relatively low information gain (I = 1.08).
- Splitting the data points vertically yields better separation, with peakier Gaussians and a correspondingly higher value of information gain (I = 2.43).

The training objective function

The following definition of information gain is used:

$$I_j = \sum_{\mathbf{v} \in \mathcal{S}_j} \log(|\Lambda_{\mathbf{y}}(\mathbf{v})|) - \sum_{i \in \{\mathtt{L},\mathtt{R}\}} \left(\sum_{\mathbf{v} \in \mathcal{S}_j^i} \log(|\Lambda_{\mathbf{y}}(\mathbf{v})|) \right) \\ \text{Node test params} \\ \boldsymbol{\theta} \in \mathcal{T}_j \quad \mathcal{S}_j^\mathtt{L}$$

Where Λ_y the covariance matrix computed from probabilistic fitting.



The training objective function

The error or fit objective function for single-variate output y is:

$$I(\mathcal{S}_{j},\theta) = H(\mathcal{S}_{j}) - \sum_{i \in \{L,R\}} \frac{|\mathcal{S}_{j}^{i}|}{|\mathcal{S}_{j}|} H(\mathcal{S}_{j}^{i})$$

$$H(\mathcal{S}) = \frac{1}{|\mathcal{S}|} \sum_{x \in \mathcal{S}} \int_{y} p(y|x) \log p(y|x) \, dy \qquad p(y|x) \sim N\left(y; \overline{y}, \sigma_{y}^{2}(x)\right)$$

$$H(\mathcal{S}) = \frac{1}{|\mathcal{S}|} \sum_{x \in \mathcal{S}} \frac{1}{2} \log\left((2\pi e)^{2} \sigma_{y}^{2}(x)\right)$$

$$I = \sum_{(x,y) \in \mathcal{S}_{j}} \log \left(\sigma_{y}(x)\right) - \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x)\right)\right) + \sum_{i \in \{L,R\}} \left(\sum_{(x,y) \in \mathcal{S}_{j}^{i}} \log \left(\sigma_{y}(x$$



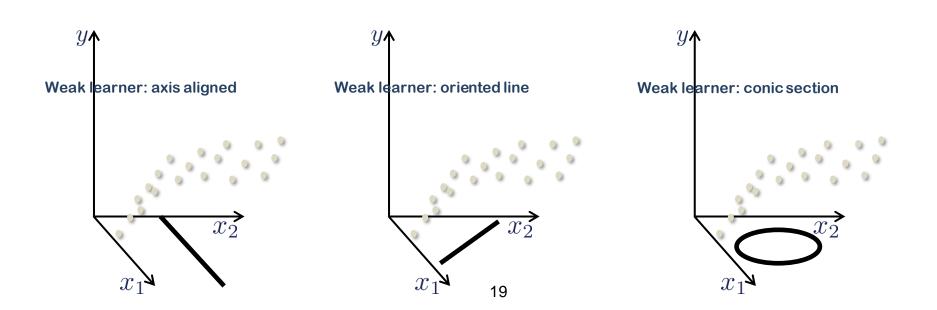
The weak learner model

 As usual, the data arriving at a split node j is separated into its left or right children according to a binary weak learner stored in an internal node, of the following general form:

$$h(\mathbf{v},\theta_i) = \{0 \text{ or } 1\}$$

where 0 and 1 can be interpreted as "false" and "true" respectively.

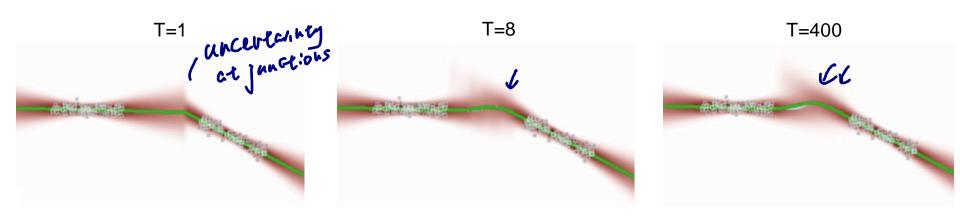
Like in classification here we consider three types of weak learners: (i) axis-aligned, (ii) oriented hyperplane, (iii) quadratic (for an illustration on 2D→1D regression).



Effect of the forest size

- A forest of shallow trees (D = 2) and varying size T is trained.
- We use axis-aligned weak learners, and probabilistic linear predictor models.
- As the number of trees increases both the prediction mean and its uncertainty become smoother.







Effect of the tree depth

- The effect of varying the maximum allowed tree depth D on the same training set is shown.
- A regression forest with D = 1 (top row in figure) corresponds to conventional linear regression (with additional confidence estimation).
- In this case the training data is more complex than a single line and thus such a degenerate forest under-fits.
- In contrast, a forest of depth D = 5 (bottom row in figure) yields over-fitting. This is highlighted in the figure by the high frequency variations in the prediction confidence and the mean of y(x).

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Effect of the tree depth

