

Ludwig-Maximilians-Universität München Institut für Informatik Lehr- und Forschungseinheit für Datenbanksysteme



Knowledge Discovery in Databases SS 2016

Chapter 7: Numerical Prediction

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Knowledge Discovery in Databases I: Numerical Prediction





1) Introduction

- Numerical Prediction problem, linear and nonlinear regression, evaluation measures
- 2) Piecewise Linear Numerical Prediction Models
 - Regression Trees, axis parallel splits, oblique splits
 - Hinging Hyperplane Models
- 3) Bias-Variance Problem
 - Regularization , Ensemble methods

Numerical Prediction



predicted

value

Wind speed

query

Delay of flight

- Related problem to classification: numerical prediction
 - Determine the numerical value of an object
 - Method: e.g., regression analysis
 - Example: prediction of flight delays



- Classification refers to predict categorical class label
- Numerical prediction models continuous-valued functions
- Numerical prediction is *similar* to classification
 - First, construct a model
 - Second, use model to predict unknown value
 - Major method for numerical prediction is regression
 - Linear and multiple regression
 - Non-linear regression







- Housing values in suburbs of Boston
- Inputs
 - number of rooms
 - Median value of houses in the neighborhood
 - Weighted distance to five Boston employment centers
 - Nitric oxides concentration
 - Crime rate per capita

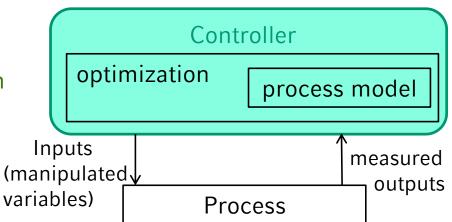


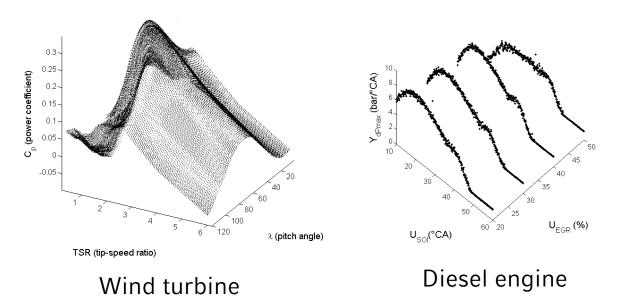
• Goal: compute a model of the housing values, which can be used to predict the price for a house in that area





- Control engineering:
 - Control the inputs of a system in order to lead the outputs to a given reference value
 - Required: a model of the process



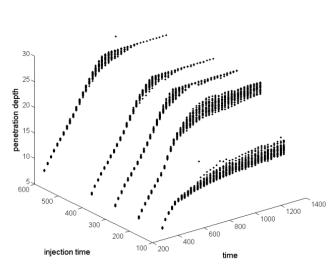


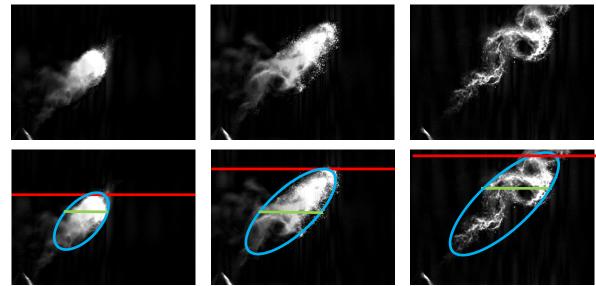




- Fuel injection process:
 - database of spray images
 - Inputs: settings in the pressure chamber
 - Outputs: spray features, e.g., penetration depth, spray width, spray area

compute a model which predicts the spray features, for input settings which have not been measured



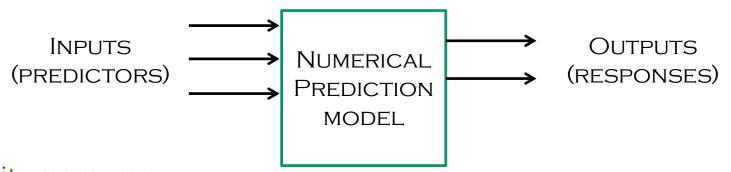




Numerical Prediction



- Given: a set of observations
- Compute: a generalized model of the data which enables the prediction of the output as a continuous value



- Quality measures:
 - Accuracy of the model
 - Compactness of the model
 - Interpretability of the model
 - Runtime efficiency (training, prediction)



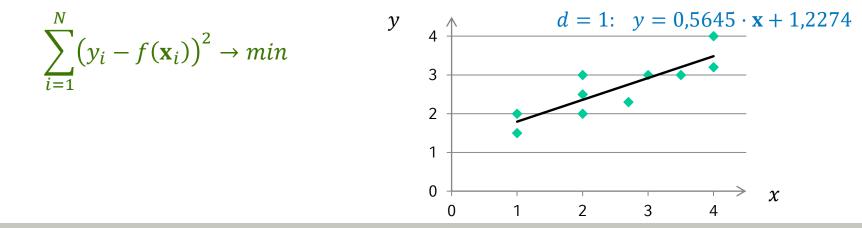
Linear Regression



- Given a set of N observations with inputs of the form x = [x₁, ..., x_d] and outputs y ∈ ℝ
- Approach: minimize the **Sum of Squared Errors (SSE)**
- Numerical Prediction: describe the outputs y as a linear equation of the inputs $\begin{bmatrix} \beta_0 \end{bmatrix}$

$$\hat{y} = f(\mathbf{x}) = \beta_0 + \beta_1 \cdot x_1 + \dots + \beta_d \cdot x_d = \begin{bmatrix} 1 \ x_1 \ \dots \ x_d \end{bmatrix} \cdot \begin{vmatrix} \beta_1 \\ \vdots \end{vmatrix} = \begin{bmatrix} 1 \ x_1 \ \dots \ x_d \end{bmatrix} \cdot \boldsymbol{\beta}$$

• Train the parameters $\boldsymbol{\beta} = [\beta_0 \ \beta_1 \ \dots \beta_d]$:



 $\left[\beta_{d}\right]$



Linear Regression



• Matrix notation: let $X \in \mathbb{R}^{N \times (d+1)}$ be the matrix containing the inputs, $Y \in \mathbb{R}^N$ the outputs, and β the resulting coefficients:

$$\mathbf{X} = \begin{bmatrix} 1 & x_{11} & \dots & x_{1d} \\ \vdots & \ddots & \vdots & \vdots \\ 1 & x_{N1} & \dots & x_{Nd} \end{bmatrix}, \qquad \mathbf{Y} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_N \end{bmatrix} \quad \Rightarrow \quad \boldsymbol{\beta} = \begin{bmatrix} \boldsymbol{\beta}_0 \\ \boldsymbol{\beta}_1 \\ \vdots \\ \boldsymbol{\beta}_d \end{bmatrix}$$

• Goal: find the coefficients β , which minimize the SSE:

$$\min_{\beta} g(\beta) = \min_{\beta} \|X\beta - Y\|_2^2 = \min_{\beta} (X\beta - Y)^T (X\beta - Y)$$

$$= \min_{\beta} (\beta^T X^T X \beta - 2Y^T X \beta + Y^T Y)$$



Linear Regression



- Set the first derivative of $g(\beta) = \beta^T X^T X \beta 2Y^T X \beta + Y^T Y$ to zero: $X^T X \beta = X^T Y$
- If $X^T X$ is non-singular then:

$$\beta = (X^T X)^{-1} \cdot X^T Y$$

• For d = 1, the regression coefficients β_0 and β_1 can be computed as:



Polynomial Regression



[0]

• Second order polynomial for d = 1:

$$\hat{y} = \beta_0 + \beta_1 \cdot x_1 + \beta_2 \cdot x_1^2 = x_d = \begin{bmatrix} 1 & x_1 & x_1^2 \end{bmatrix} \cdot \begin{bmatrix} \beta_0 \\ \beta_1 \\ \beta_2 \end{bmatrix}$$

with
$$X = \begin{bmatrix} 1 & x_{11} & x_{11}^2 \\ \vdots & \vdots & \vdots \\ 1 & x_{N1} & x_{N1}^2 \end{bmatrix}$$
 and $\beta = (X^T \cdot X)^{-1} \cdot X^T \cdot Y$

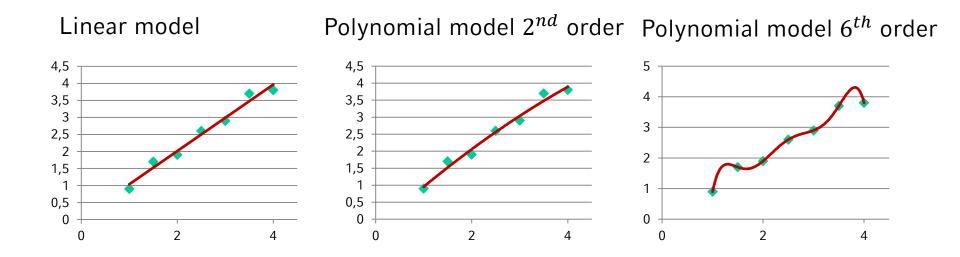
• Second order polynomial for d = 2:

$$\hat{y} = \beta_0 + \beta_1 \cdot x_1 + \beta_2 \cdot x_2 + \beta_3 \cdot x_1^2 + \beta_4 \cdot x_2^2 + \beta_5 \cdot x_1 \cdot x_2$$





- The number of coefficients increases exponentially with k and d
- Model building strategies: forward selection, backward elimination
- The order of the polynomial should be as low as possible, high order polynomials tend to overfit the data





Nonlinear Regression



- Different nonlinear functions can be approximated
- Transform the data to a linear domain

 $\hat{y} = \boldsymbol{\alpha} \cdot e^{\boldsymbol{\gamma} x} \Rightarrow \ln(\hat{y}) = \ln(\boldsymbol{\alpha}) + \boldsymbol{\gamma} x$

 $\Rightarrow \hat{y}' = \beta_0 + \beta x$

(for $\hat{y}' = \ln(\hat{y})$, $\beta_0 = \ln(\alpha)$, and $\beta_1 = \gamma$)

- The parameters β_0 and β are estimated with LS
- The parameters α and γ are obtained, describing an exponential curve which passes through the original observations
- Problem: LS determines normally distributed errors in the transformed space ⇒ skewed error distribution in the original space



Nonlinear Regression



- Different nonlinear functions can be approximated
- Outputs are estimated by a function with nonlinear parameters, e.g., exponential, trigonometric
- Example type of function:

$$\hat{y} = \beta_0 + \beta_1 e^{\beta_2 x} + \sin(\beta_3 x)$$

- Approach: the type of nonlinear function is chosen and the corresponding parameters are computed
- No closed form solution exists ⇒ numerical approximations:
 - Gauss Newton, Gradient descent, Levenberg-Marquardt





- Problems:
 - Linear regression most of the real world data has a nonlinear behavior
 - Polynomial regression limited, cannot describe arbitrary nonlinear behavior
 - General nonlinear regression the type of nonlinear function must be specified in advance



Piecewise Linear Regression



• Piecewise linear functions:

$$f(\mathbf{x}) = \begin{cases} \beta_{00} + \beta_{01} \cdot x_1 + \dots + \beta_{0d} \cdot x_d, & \mathbf{x} \in \mathcal{D}_1 \\ \vdots \\ \beta k_0 + \beta_{k1} \cdot x_1 + \dots + \beta_{kd} \cdot x_d, & \mathbf{x} \in \mathcal{D}_k \end{cases}$$

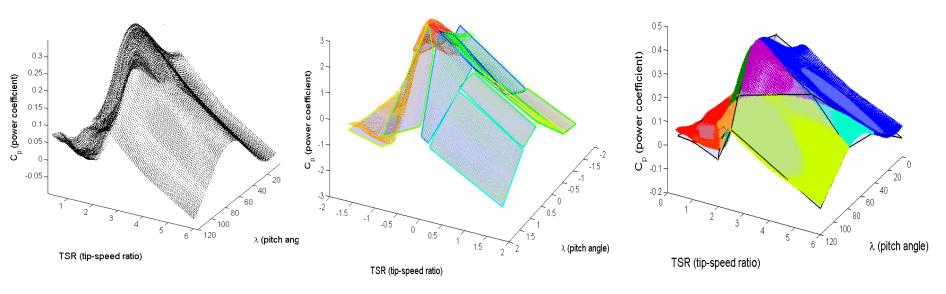
- Simple approach
- Able to describe arbitrary functions
- The **accuracy** is increasing with an increasing number of partitions/linear models
- The **compactness & interpretability** is increasing with a decreasing number of partitions/ linear models
- Challenge: find an appropriate partitioning in the input space (number and shapes)

 \wp_1, \dots, \wp_k are partitions in the input space





1. Introduction of different learning techniques for piecewise linear models



Training set

Piecewise linear model with regression trees Continuous piecewise linear model with HHmodels

2. Discussion of the bias-variance problem, regression and *ensemble techniques*





1) Introduction

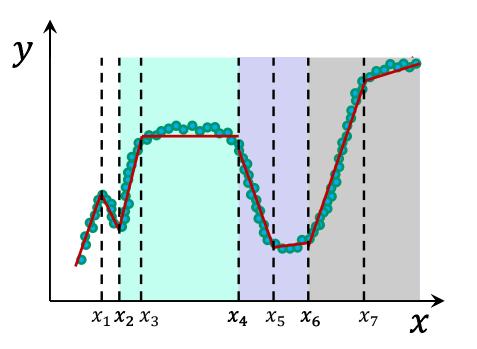
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Regression Trees



- Greedy divide and conquer: recursive partitioning of the input space
- Example with input *x* and output *y* :

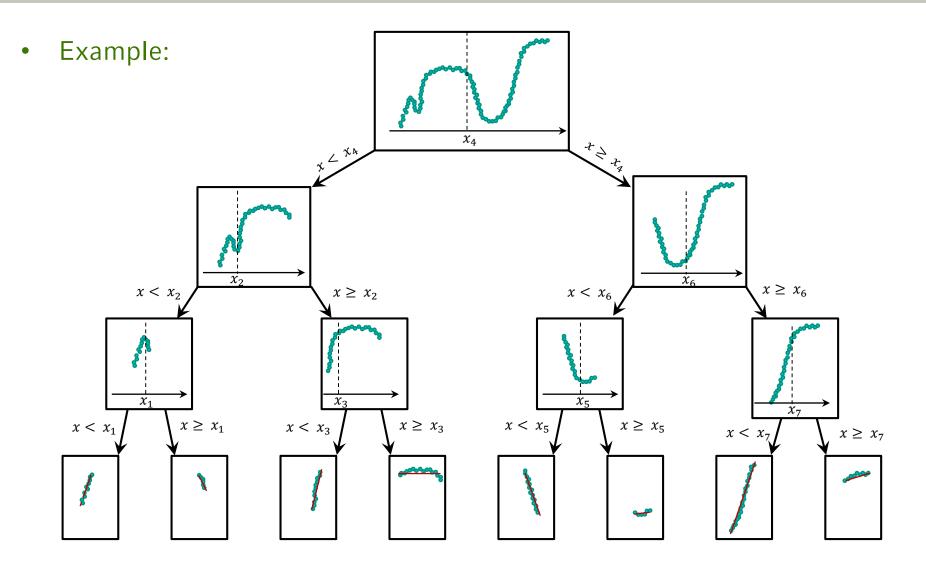


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Regression Trees







Regression Trees

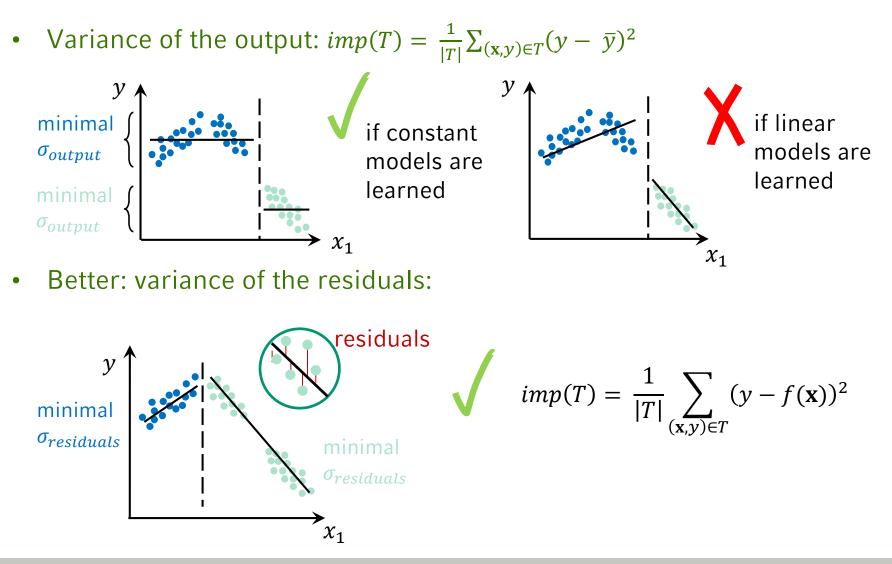


- General approach of learning a regression tree:
 - Given: set of observations T
 - Find a split of T in T_1 and T_2 with minimal summed impurity $imp(T_1) + imp(T_2)$
 - If the stopping criterion is not reached: repeat for T_1 and T_2
 - If the stopping criterion is reached: undo the split
- Internal node denotes a test in the input space
- Branch represents an outcome of the test
- Leaf nodes contain a linear function, used to predict the output



Impurity Measure









- The recursive splitting is stopped if:
 - a) The sample size of a node is below a specified threshold
 - b) The split is not significant:
 - If the relative impurity ratio induced by a split is higher than a given threshold, then the split is not significant $imp(T_1) + imp(T_2)$.

$$\tau = \frac{imp(T_1) + imp(T_2)}{imp(T)} > \tau_0$$

- As the tree grows the resulting piecewise linear model gets more accurate. τ increases, becoming higher than τ_0
- Choosing the parameter $\tau_0 \Leftrightarrow$ trading accuracy with overfitting
- stopping too soon \Rightarrow model is not accurate enough
- stopping too late \Rightarrow model overfits the observations

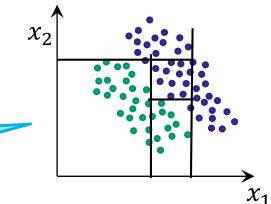


Split Strategy



- The split strategy determines how the training samples are partitioned, whether the split is actually performed is decided by the stopping criterion
- The most common splits are axis parallel:
 - Split = a value in one input dimension
 - Compute the impurity of all possible splits in all input dimensions and choose at the end the split with the lowest impurity
 - For each possible split compute the two corresponding models and their impurity ⇒ expensive to compute

4 axis parallel splits in the 2D input space, in order to separate the red from the blue samples





Strategy for Oblique Splits



- More intuitive to use oblique splits
- An oblique split is a linear separator in the input space instead of a split value in an input dimension
- The optimal split (with minimal impurity measure) cannot be efficiently computed

 χ_1

• Heuristic approach required

 χ_2

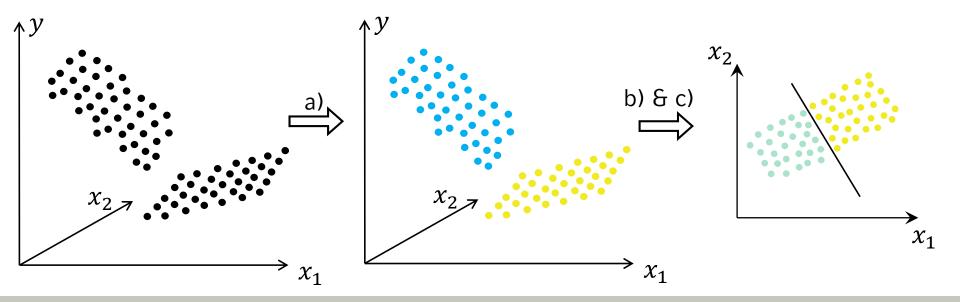
1 oblique split in the 2D input space, in order to separate the red from the blue samples



Strategy for Oblique Splits



- Heuristic approach:
 - a) Compute a clustering in the full (input + output) space, such that the samples are as well as possible described by linear equations
 - b) Project the clusters onto the input space
 - c) Use the clusters to train a linear classifier in the input space. Split = separating hyperplane in input space

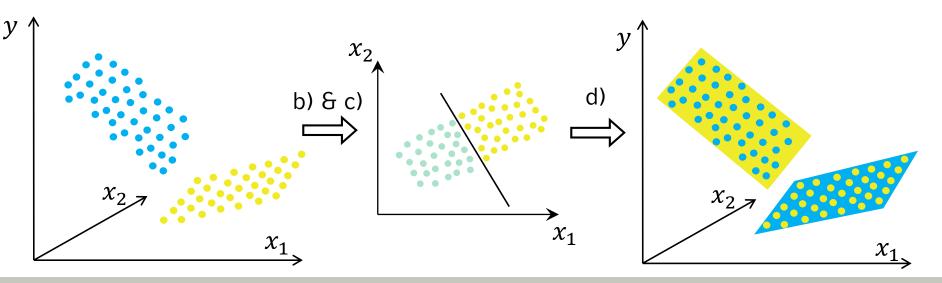




Strategy for Oblique Splits



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 - b) Project the clusters onto the input space
 - c) Use the clusters to train a linear classifier in the input space. Split = separating hyperplane in input space
 - d) Compute linear models for the two linearly separated clusters

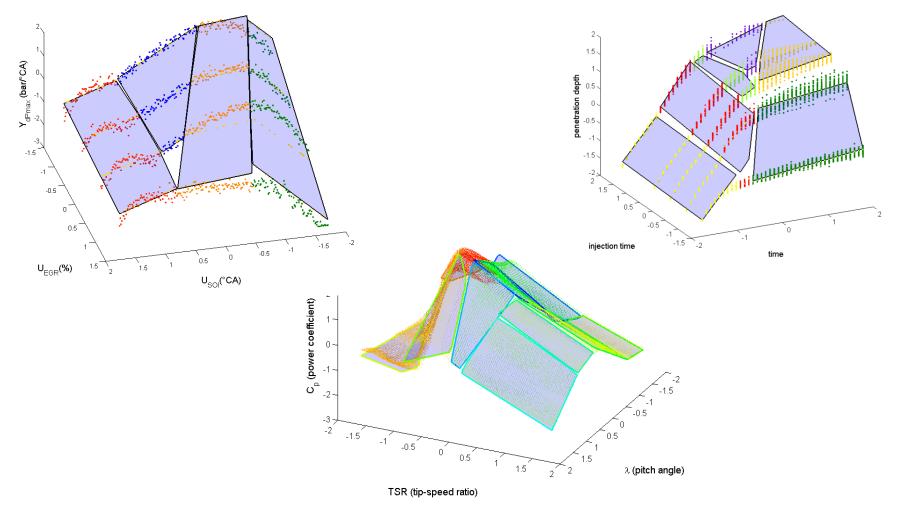




Example Models



• Example piecewise linear models (with oblique splits in the input space):







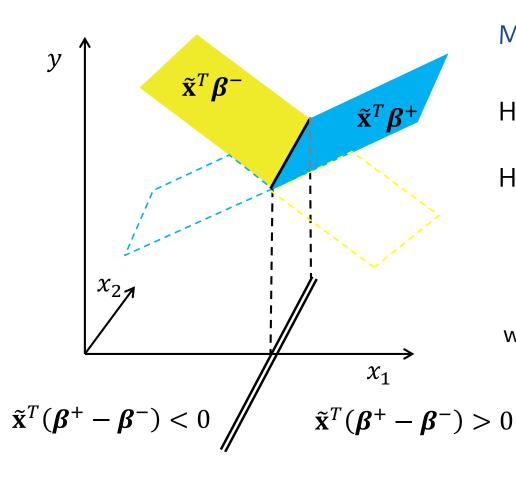
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• Hinging Hyperplane Model (HH-model) for continuous models



Model:
$$f(\mathbf{x}) = \sum_{i=1}^{K} h_i(\mathbf{x})$$

Hinge: $\Delta = \beta^+ - \beta^-$
Hinge function:
 $(\mathbf{\tilde{x}}^T \beta^+ - \mathbf{\tilde{x}}^T \cdot \mathbf{A})$

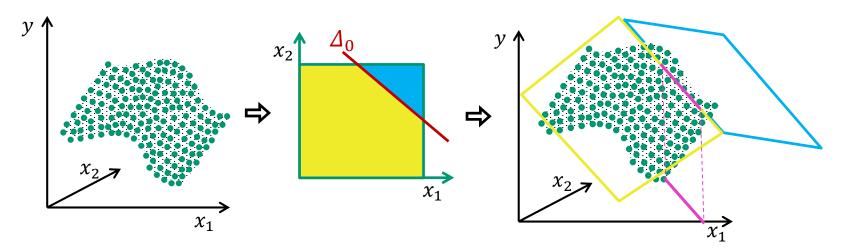
 $h(\mathbf{x}) = \begin{cases} \tilde{\mathbf{x}}^T \boldsymbol{\beta}^+, & \tilde{\mathbf{x}}^T \cdot \boldsymbol{\Delta} > 0\\ \\ \tilde{\mathbf{x}}^T \boldsymbol{\beta}^-, & \tilde{\mathbf{x}}^T \cdot \boldsymbol{\Delta} \le 0 \end{cases}$ with $\tilde{\mathbf{x}}^T = (1, \mathbf{x}_1, \dots, \mathbf{x}_n).$

[L. Breiman (1993)]





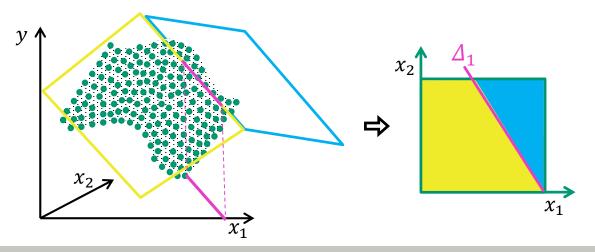
- Hinge Finding Algorithm (HFA)
 - 1) Start with a random partitioning of the input space: Δ_j (j = 0)
 - 2) Determine the two corresponding partitions:
 - $S_j^- = \{ \mathbf{x} \mid \tilde{\mathbf{x}}^T \boldsymbol{\Delta}_j \leq 0 \}$ and $S_j^+ = \{ \mathbf{x} \mid \tilde{\mathbf{x}}^T \boldsymbol{\Delta}_j > 0 \}$
 - 3) Compute the regression hyperplanes for S_j^+ and S_j^-







- Hinge Finding Algorithm (HFA)
 - 3) Compute the regression hyperplanes for S_j^+ and S_j^-
 - 4) Compute the hinge Δ_{j+1} from the regression coefficients β_j^- and β_j^+
 - 5) Determine the new partitions S_{j+1}^+ and S_{j+1}^- determined by Δ_{j+1}
 - 6) If $S_{j+1}^+ = S_j^+$ or $S_{j+1}^+ = S_j^-$, then stop, else return to step 3).





Line Search for the Hinge Finding Algorithm



 x_1

• The Hinge Finding Algorithm (HFA) might not converge – a hinge might induce a partitioning outside the defined input space x_2

- Line search: binary search to guarantee convergence (to a local minimum)
- Instead of updating the hinge directly $\Delta_j \rightarrow \Delta_{j+1}$, first check the accuracy improvement brought by Δ_{j+1}
- If Δ_{j+1} does not improve the model impurity, then perform a binary search after the linear combination of Δ_j and Δ_{j+1} yielding Δ_j the lowest impurity

$$\boldsymbol{\Delta}'_{j+1} = \boldsymbol{\Delta}_i + \lambda \left(\boldsymbol{\Delta}_{j+1} - \boldsymbol{\Delta}_j \right), \qquad \lambda \in \left\{ \frac{1}{2}, \frac{1}{4}, \frac{1}{8}, \frac{1}{16} \right\}$$



Fit Multiple Hinges



• Goal: Describe the target function *y* as a sum of *N* hinge functions

$$y = \sum_{i=1}^{N} h_i(x)$$

• Each hinge function h_i can be seen as fitted to the residual function

$$y_{[i]} = y - \sum_{j \neq i} h_j(x),$$

since then $h_i(x) = y - \sum_{j \neq i} h_j(x)$.

- Fit multiple hinges iteratively in *N* steps:
 - Start with $h_1(x) = \cdots = h_N(x) = 0$
 - Step *n*: Fit the *n*-th hinge h_n to $y_{[n]} = y (h_1(x) + \dots + h_{n-1}(x))$. Then repeatedly refit h_1 to $y_{[1]}$, ..., and h_{n-1} to $y_{[n-1]}$ until the hinges do not change anymore.

Fit Multiple Hinges



- Example:
 - Step 1: Fit the first hinge function h_1 to $y_{[1]} = y 0$.
 - Step 2: Compute the residual outputs

$$y_{[2]} = y - h_1(x).$$

Fit the second hinge function h_2 to $y_{[2]}$. Then refit the first hinge to $y_{[1]} = y - h_2(x)$.

- Step 3: Compute the residual outputs

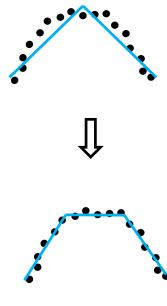
$$y_{[3]} = y - h_1(x) - h_2(x).$$

Fit the third hinge function h_3 to $y_{[3]}$.

Then repeatedly refit

$$h_1$$
 to $y_{[1]} = y - h_2(x) - h_3(x)$ and
 h_2 on $y_{[2]} = y - h_1(x) - h_3(x)$.

until no more changes occur.

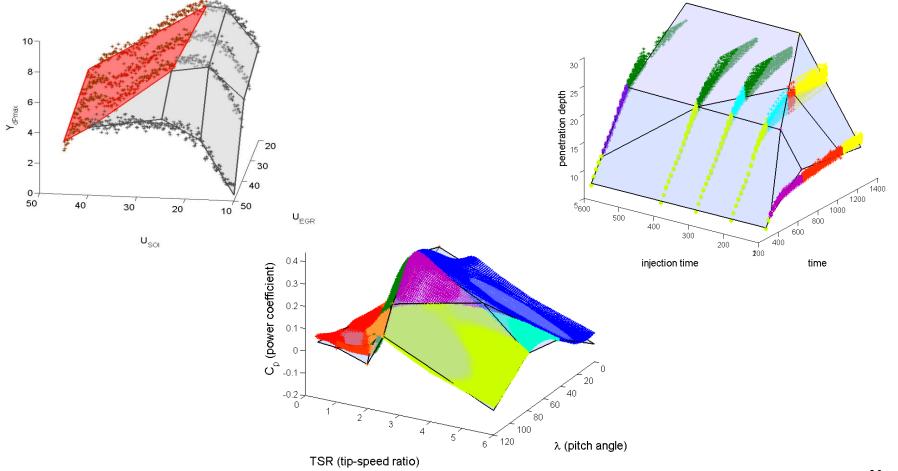




Example Models



• Example piecewise linear models (with oblique splits in the input space):







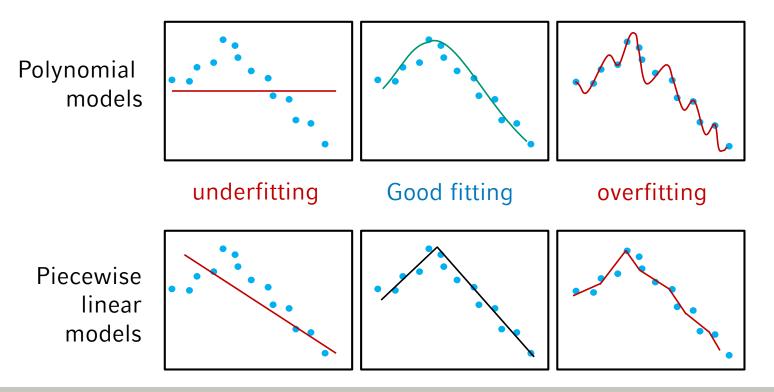
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- Underfitting = a learned model is not flexible enough to capture the underlying trend
- Overfitting = a learned model is too flexible, allowing to capture illusory trends in the data, which appear due to noise





Bias and Variance



- Assuming that the data generation process can be repeated (with a certain amount of randomness) \Rightarrow obtain several datasets D_i & for each D_i a model f_i is learned
- Bias = the difference between the average prediction of these models and the correct value
- Variance = the variability of the predictions of the different models

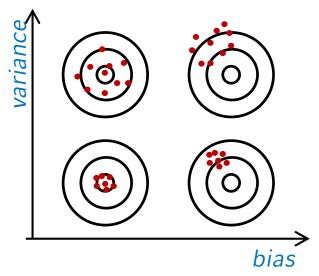


Image after http://scott.fortmann-roe.com/docs/BiasVariance.html



Bias-Variance Tradeoff



- Underfitting = low variance, high bias (e.g. use mean output as estimator)
- *High bias* = a model does not approximate the underlying function well
- Overfitting = high variance, low bias
- When a model is too complex, small changes in the data cause the predicted value to change a lot ⇒ high variance
- Search for the best tradeoff between bias and variance
- Regression: control the bias-variance tradeoff by means of the polynomial order/number of coefficients
- Regression trees: control the bias-variance tradeoff by means of the tree size/number of submodels



Bias and Variance



$$(\star): E[z^2] = E[(z - E[z])^2] + E[z]^2$$

• Consider the expected prediction error of a learned model:

 $Err(\mathbf{x}) = E[(f(\mathbf{x}) - y)^2]$ $Err(\mathbf{x}) = E[f(\mathbf{x})^2 - 2f(\mathbf{x})y + y^2]$ $Err(\mathbf{x}) = E[f(\mathbf{x})^{2}] - 2E[f(\mathbf{x})]E[y] + E[y^{2}]$ (\star) $Err(\mathbf{x}) = E\left[(f(\mathbf{x}) - \overline{f(\mathbf{x})})^2\right] + \overline{f(\mathbf{x})}^2 - 2\overline{f(\mathbf{x})}E[y] + E\left[(y - E[y])^2\right] + E[y]^2$ $Err(\mathbf{x}) = E\left[(f(\mathbf{x}) - \overline{f(\mathbf{x})})^2\right] + \left(\overline{f(\mathbf{x})} - E[y]\right)^2 + E\left[(y - E[y])^2\right]$ error due to the error due to the irreducible variance of *f* bias of *f* error (systematic error)



Bias and Variance



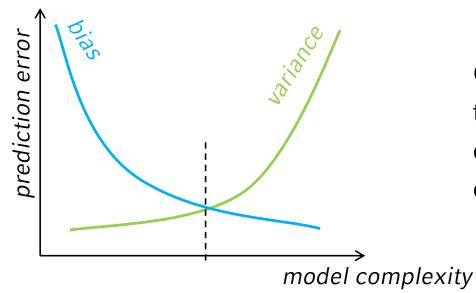
• Consider the expected prediction error of a learned model:

$$Err(\mathbf{x}) = E\left[(f(\mathbf{x}) - \overline{f(\mathbf{x})})^2\right] + \underbrace{\left(\overline{f(\mathbf{x})} - E[y]\right)^2}_{Y} + E\left[(y - E[y])^2\right]$$

error due to the variance of *f*

error due to the bias of *f* (systematic error)

irreducible error



Carefully balance between these two types of error, in order to minimize the total expected prediction error



Regularization



• Minimizing the sum of squared errors

$$\sum_{(x,y)\in T} (f(x) - y)^2 \to \min$$

computes an unbiased linear model with very high variance

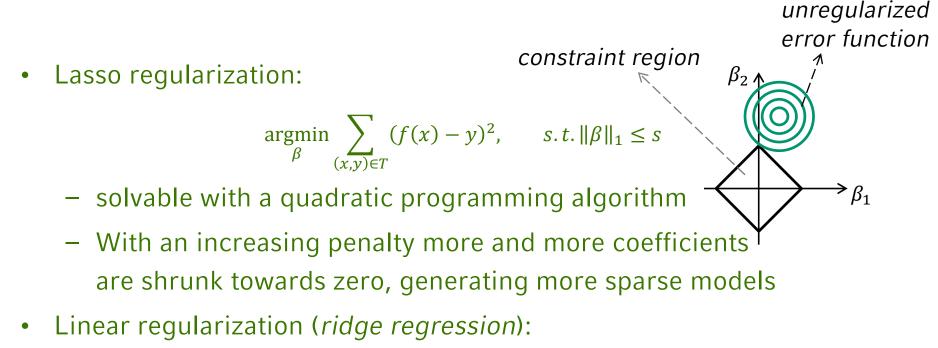
- Idea: give up the unbiasedness and obtain a variance decrease by penalizing the model complexity
- Regularization: simultaneously minimize the sum of squared errors and the norm of the coefficient vector
- Linear regularization (*ridge regression*):

$$\sum_{(x,y)\in T} (f(x) - y)^2 + \lambda \|\beta\|_2^2 \longrightarrow \min$$



Regularization

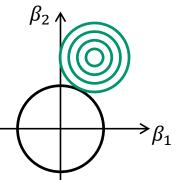




$$\underset{\beta}{\operatorname{argmin}} \sum_{(x,y)\in T} (f(x) - y)^2, \qquad s.t. \|\beta\|_2^2 \le s$$

- solvable similar to SSE: $\beta = (X^T X + \lambda I)^{-1} \cdot X^T Y$

Reduces all coefficients simultaneously







- When discussing the bias-variance tradeoff, we assumed infinitely many replications of our data set, but in practice we have only one training set T
- Simulate multiple training sets $T_1, T_2, ..., T_k$ by constructing bootstrap replicates of the original training set T, by randomly drawing samples from T (with replacement) such that $|T_j| = |T|, j \in \{1, ..., k\}$
- Learn a model f_j for each replicate T_j (use as test set $T_{Sj} = T \setminus T_j$)
- For each input x, we have several predictions y₁,..., y_k ⇒ compute the average prediction

• $f(\mathbf{x}) \approx \overline{f(\mathbf{x})} \Rightarrow (f(\mathbf{x}) - \overline{f(\mathbf{x})})^2 \approx \mathbf{0} \Rightarrow$ the variance is removed/reduced

• Bias: $(\overline{f(\mathbf{x})} - E[y])^2$ is the same as before



Ensemble Methods



- Bagging:
 - use it for models with a low bias
 - If the bias is low, bagging reduces the variance, while bias remains the same
 - use it for complex models, which tend to overfit the training data
 - in practice it might happen that the bagging approach slightly increases the bias
- Boosting:
 - Can be adapted for regression models
 - Reduces the bias in the first iterations
 - Reduces the variance in later iterations