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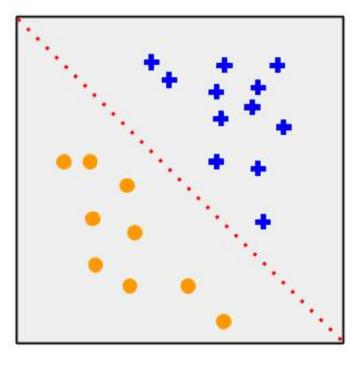
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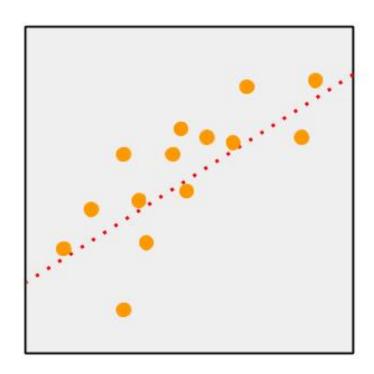


I. Regression vs. Classification

[&]quot;Regression" is the task of predicting a continuous quantity.



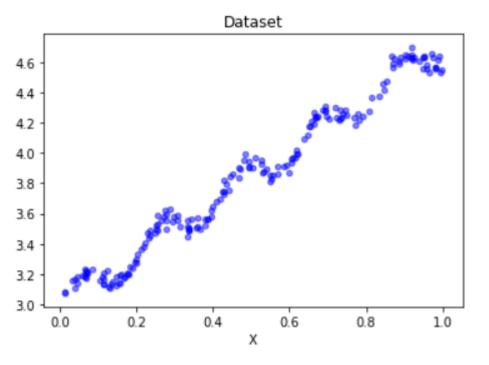




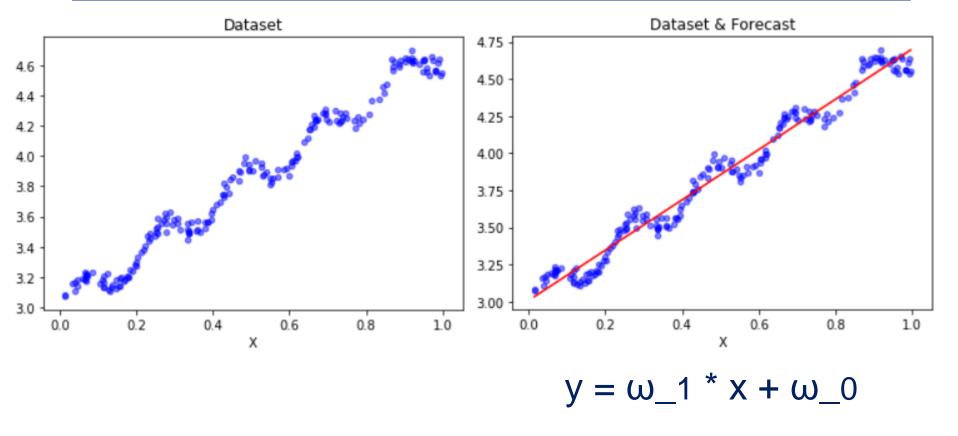
Regression



[&]quot;Classification" is the task of predicting a discrete class label.



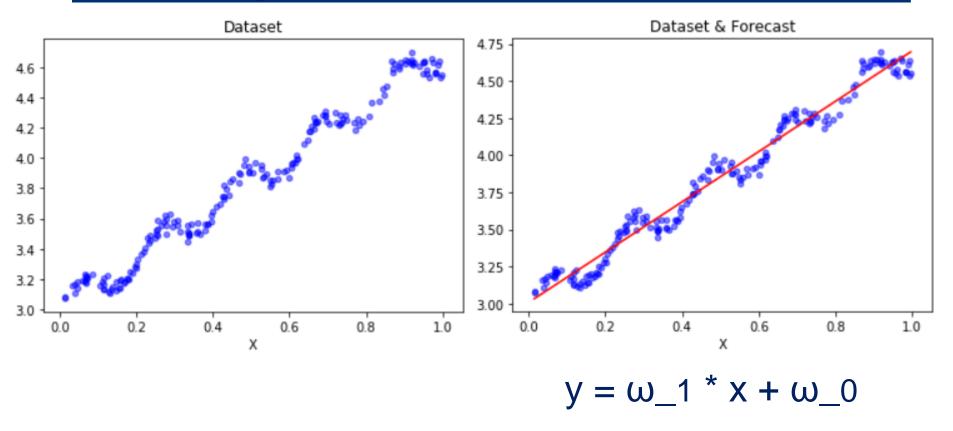




 $\omega_{1}: 1.7$

 $\omega_{-}0:3.0$





how to find the best weight?

$$\omega_{-}1:1.7$$
 $\omega_{-}0:3.0$



How to find the best **weight**:

find the *least squared error*

$$\sum_{i=1}^{m} (y_i - x_i^T \omega)^2$$

The expression in *matrix* is

$$(Y - X\omega)^T (Y - X\omega)$$

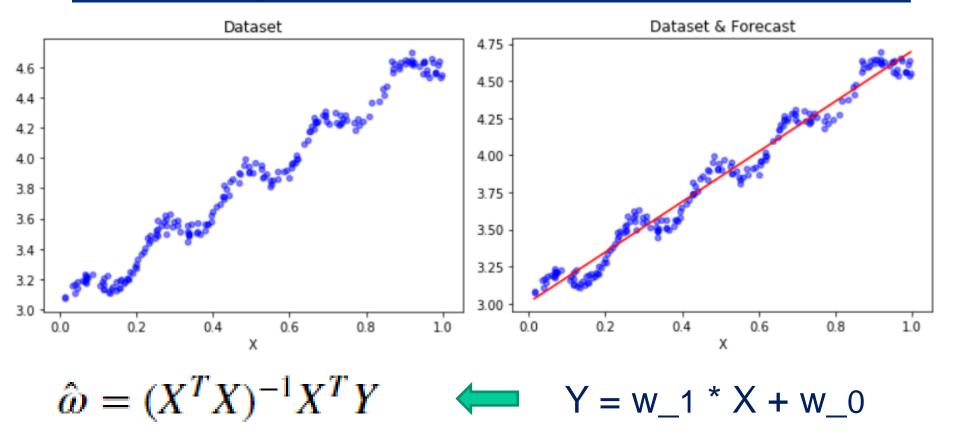
After the <u>derivation</u> with respect to ω , we get

$$X^{T}(Y-X\omega)$$

Setting this to zero and solve for ω to get the following equation

$$\hat{\omega} = (X^T X)^{-1} X^T Y$$





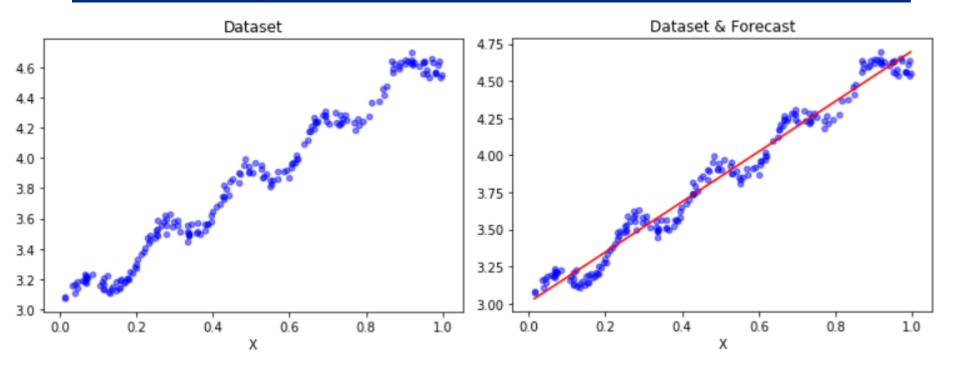
weights: [[3.00774324]

[1.69532264]]

 $w_0: 3.0$

w_1 : 1.7



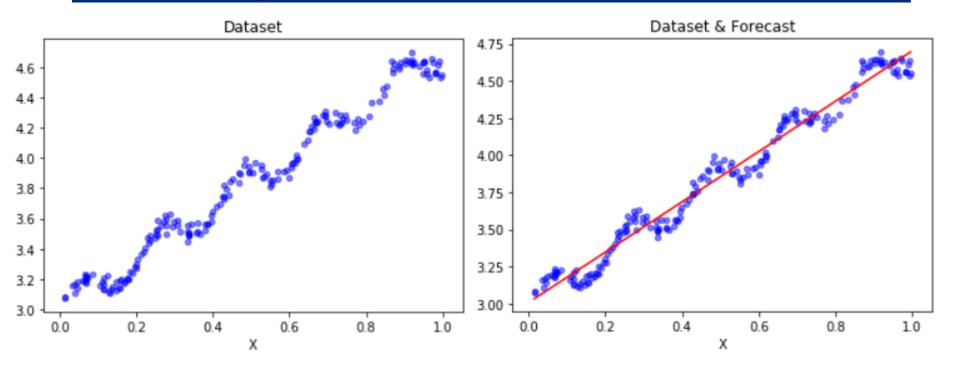


But how far do they differ from each other?

numpy.corrcoef(pred, actual): array([[1. , 0.98647356],

[0.98647356, 1.]])

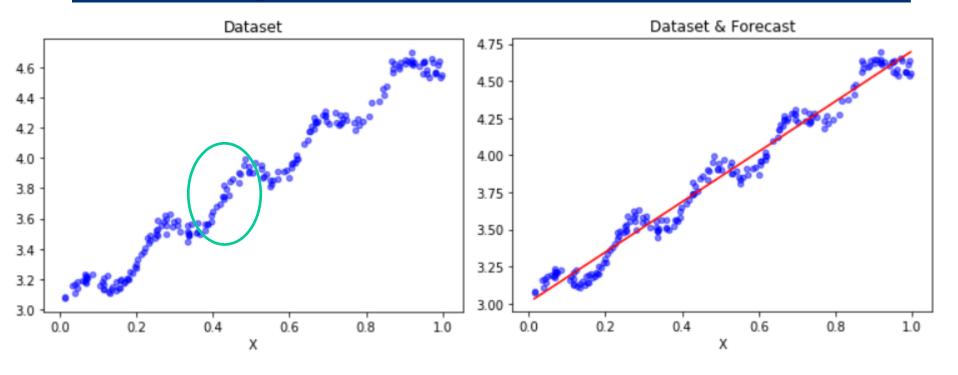




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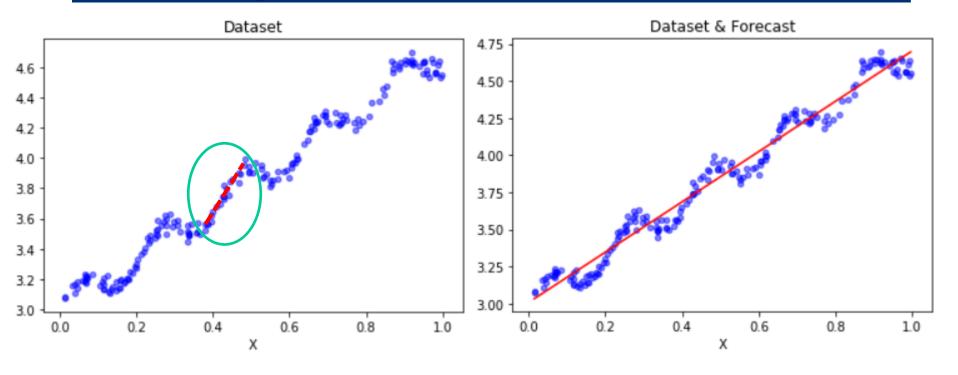




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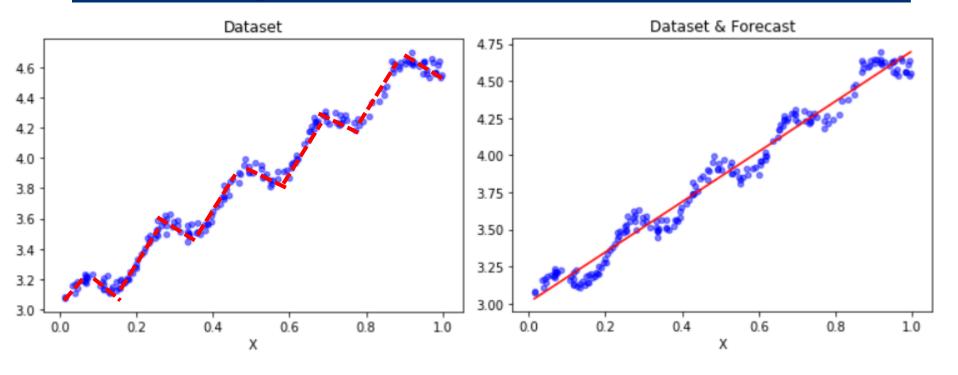




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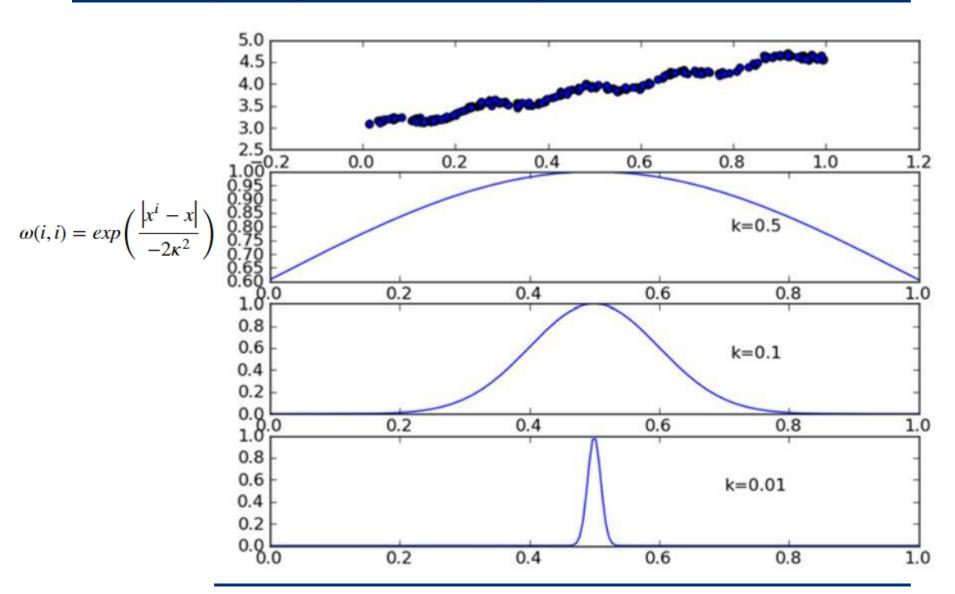
More weights are given to those data points which are close to the data point of interest, then the least-squares regression similar to the linear regression will be carried out.

$$\hat{\omega} = (X^T X)^{-1} X^T Y \longrightarrow \hat{\omega} = (X^T W X)^{-1} X^T W Y$$

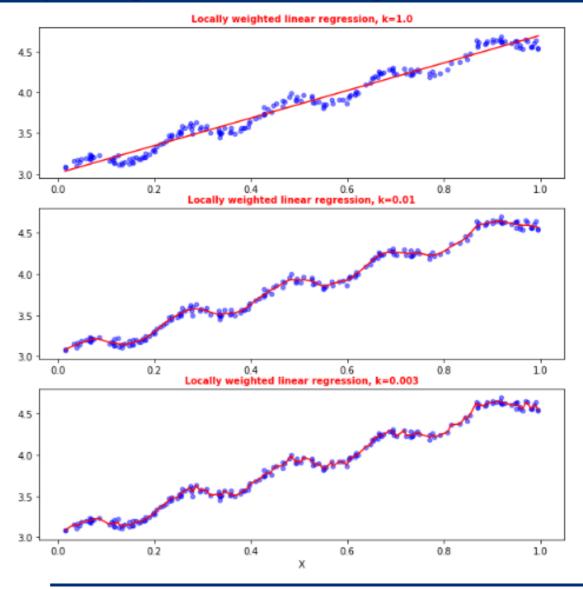
W is a matrix and will be generated by a <u>kernel</u> function, which shall give nearby points more weights than other points. The mostly used kernel is <u>Gaussian</u> and assigns the weights by

$$\omega(i,i) = exp\left(\frac{\left|x^{i} - x\right|}{-2\kappa^{2}}\right)$$

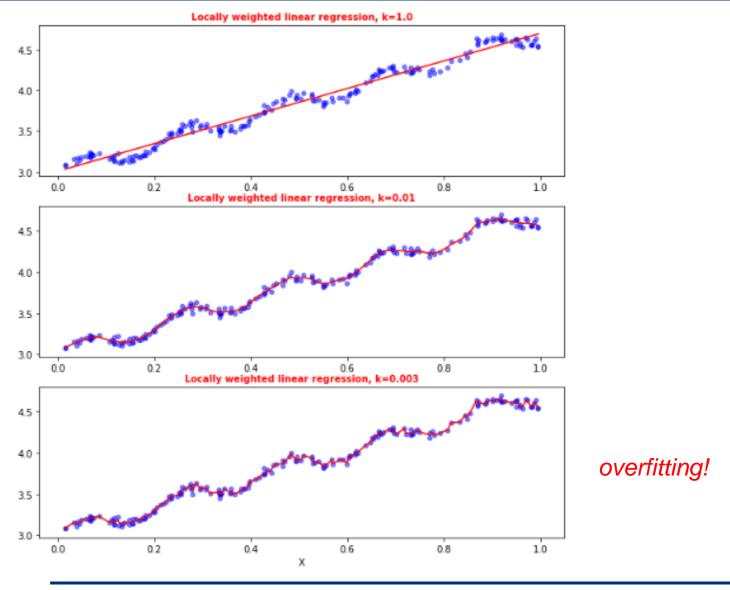














Example: predicting the age of an abalone

Testdata: abalone.txt

Feature_X:								Age_Y:
[[1.	0.455	0.365	0.095	0.514	0.2245	0.101	0.15]	[[15.]
[1.	0.35	0.265	0.09	0.225	0.0995	0.048	0.07]	[7.]
[-1.	0.53	0.42	0.135	0.677	0.2565	0.141	0.21]	[9.]
[]	[10.]
[0.	0.33	0.255	0.08	0.205	0.0895	0.039	0.05]]	[7.]]

<u>Training</u> set and <u>test</u> set are *identical*:

k=0.1, squared error: 56.82523568972884 k=1.0, squared error: 429.8905618700651 k=10, squared error: 549.1181708826451



Example: predicting the age of an abalone

<u>Training</u> set and <u>test</u> set are *identical*:

k=0.1, squared error: 56.82523568972884 k=1.0, squared error: 429.8905618700651 k=10, squared error: 549.1181708826451

<u>Training</u> set and <u>test</u> set are *different*.

k=0.1, squared error: 41317.161723642595

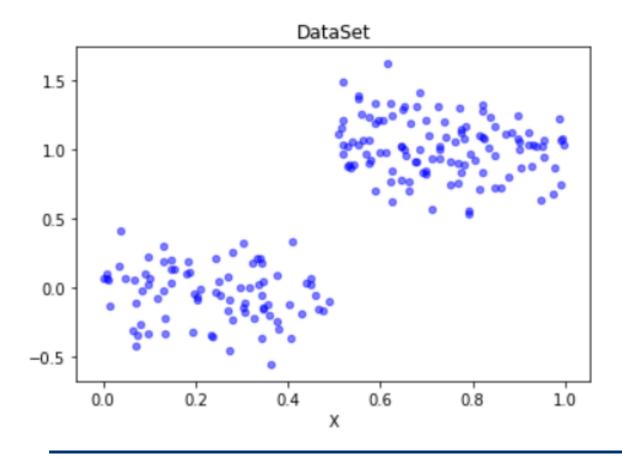
k=1.0, squared error: 573.526144189767

k=10, squared error: 517.5711905387598

Linear regression: squared error: 518.6363153249638



How to deal with the *nonlinearities* in real life?



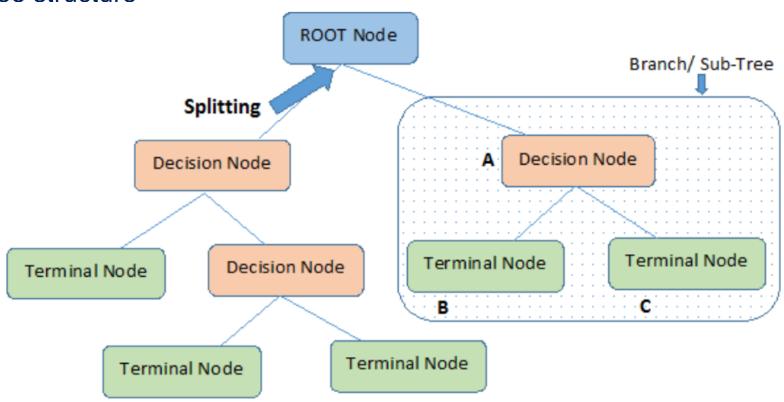


How to deal with the *nonlinearities* in real life?

Tree Regression

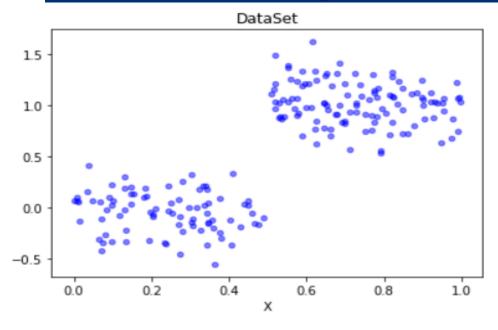


Tree structure



Note:- A is parent node of B and C.





Feature Index:

Threshold value:

Leaf node value:

Number of Samples:

0

0.5

0.0 or 1.0

100

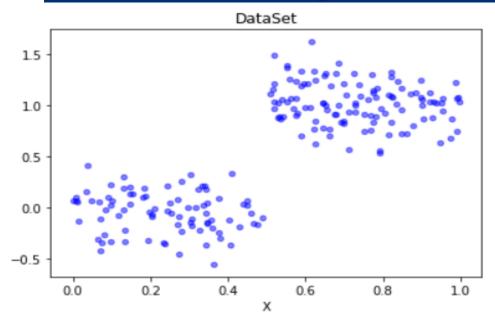
Y = **0.0**100

 $X_0 <= 0.5$?

Y = **1.0**

100





How to find the splitting feature and splitting point?

1

 $X_0 <= 0.5$?

Feature Index:

Threshold value:

Leaf node value:

Number of Samples:

0

0.5

0.0 or 1.0

100



Y = **1.0**

100



How to deal with the *nonlinearities* in real life?

Tree regression:

It implements a new algorithm called <u>CART</u> (Classification And Regression Trees). It is well-known and well-documented tree-building algorithm that makes <u>binary splits</u> to handle continuous variables.

By doing this we choose a **feature** and make **values** <u>greater</u> than the desired go on the **right** side of the tree and all the other values go on the **left** side.



How to make a binary split?

We need to select: splitting <u>feature 'xj'</u> and a splitting <u>point 's'</u> so that we can divide data into two regions R1 and R2:

$$R_1(j,s) = \{x | x^{(j)} \le s\}, \quad R_2(j,s) = \{x | x^{(j)} > s\}$$

Then we calculate the <u>average</u> value for each generated region by

$$\hat{c}_1 = ave(y_i | x_i \in R_1(j, s)), \quad \hat{c}_2 = ave(y_i | x_i \in R_2(j, s))$$

Goal: find such (ĉ 1, ĉ 2) which gives the **minimum** of total squared error as follows

$$\min_{j,s} \left[\min_{\hat{c}_1} \sum_{x_i \in R_1(j,s)} (y_i - \hat{c}_1)^2 + \min_{\hat{c}_2} \sum_{x_i \in R_2(j,s)} (y_i - \hat{c}_2)^2 \right]$$



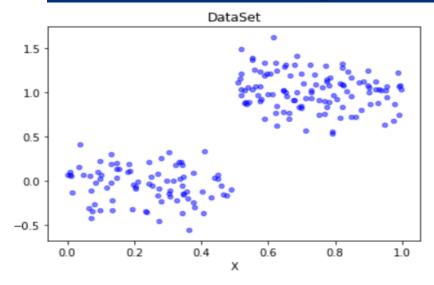
Example: binary split in the tree regression

Let's split it by the value of the second feature, threshold value: 0.5

```
matrix([[1., 0., 0., 0.], [0., 1., 0.], [0., 0., 0.], [0., 0., 1., 0.], [0., 0., 0., 1.]])
```

Left: [[1. 0. 0. 0.] [0. 0. 1. 0.] [0. 0. 0. 1.]] Right: [[0. 1. 0. 0.]]

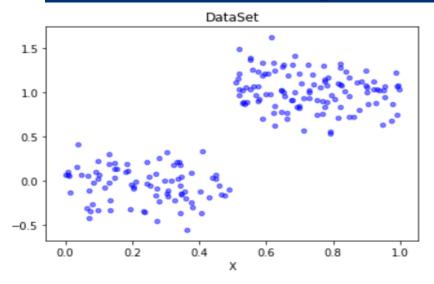




Stop Conditions:

Mind. error reduction: 1





print(createTree(data_mat))

{'spInd': 0,

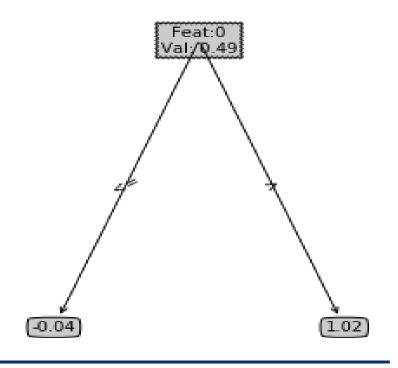
'spVal': 0.48813,

'left': -0.04465028571428572,

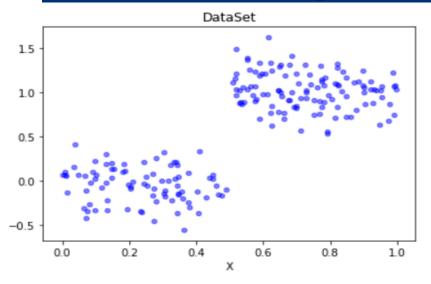
'right': 1.0180967672413792 }

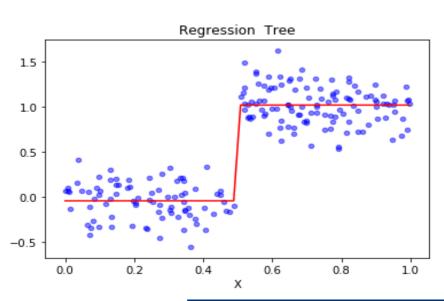
Stop Conditions:

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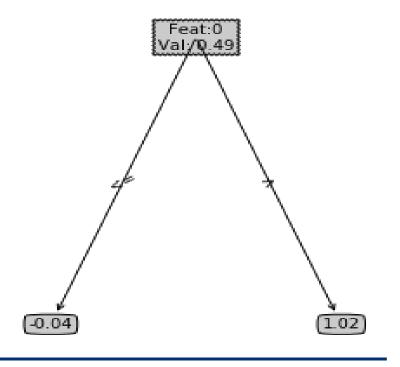




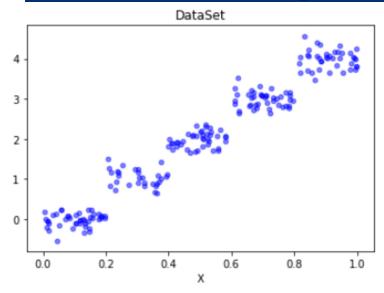


Stop Conditions:

Mind. error reduction: 1



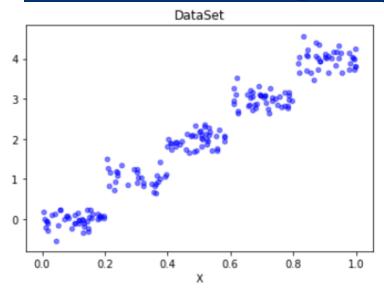




Stop Conditions:

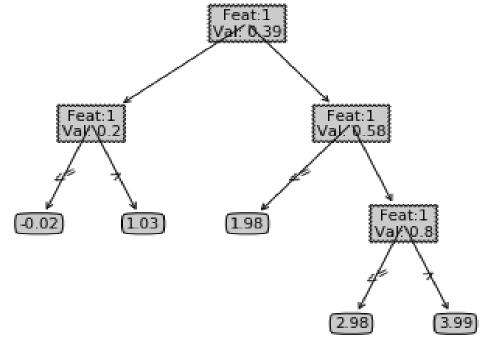
Mind. error reduction: 1



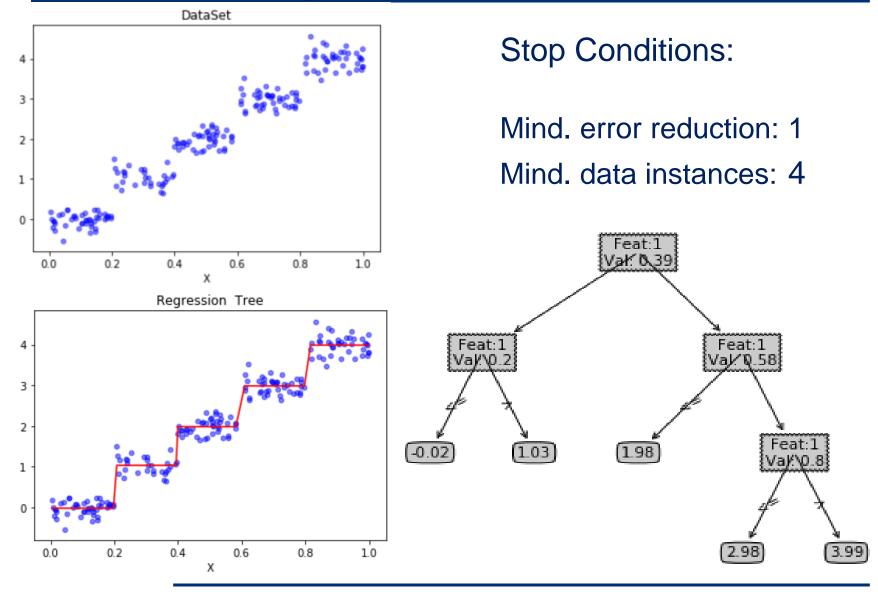


Stop Conditions:

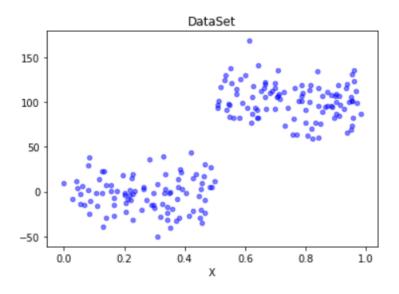
Mind. error reduction: 1







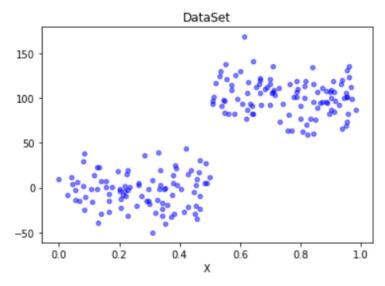




Stop Conditions:

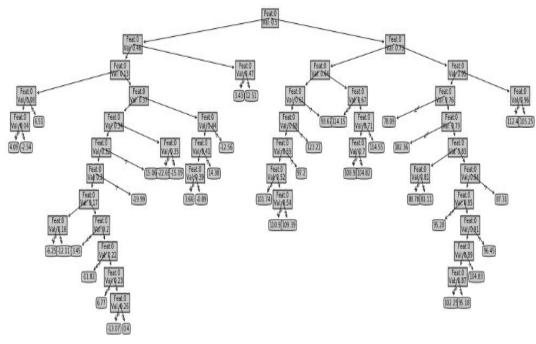
Mind. error reduction: 1



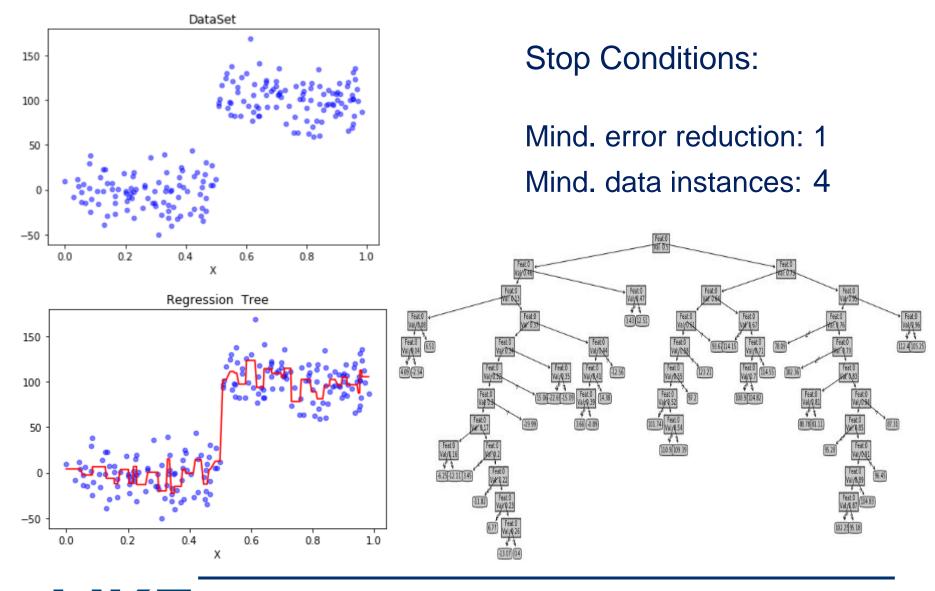


Stop Conditions:

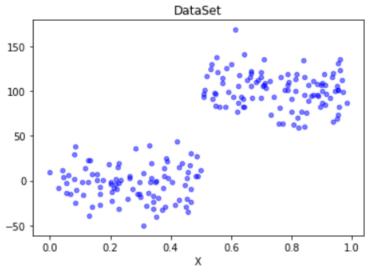
Mind. error reduction: 1

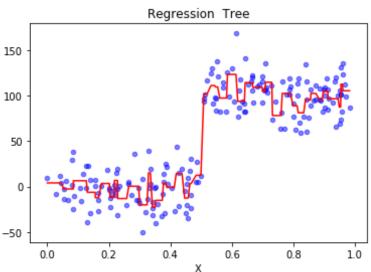








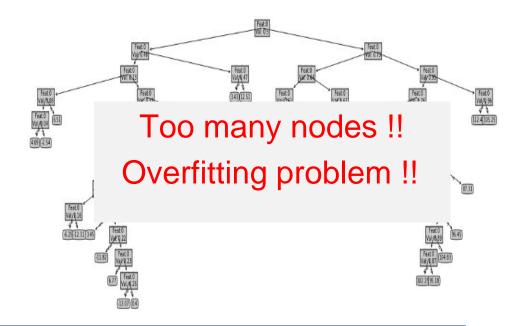




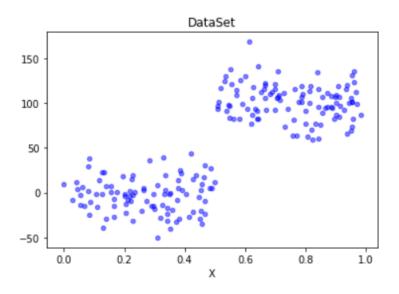
Stop Conditions:

Mind. error reduction: 1

Mind. data instances: 4





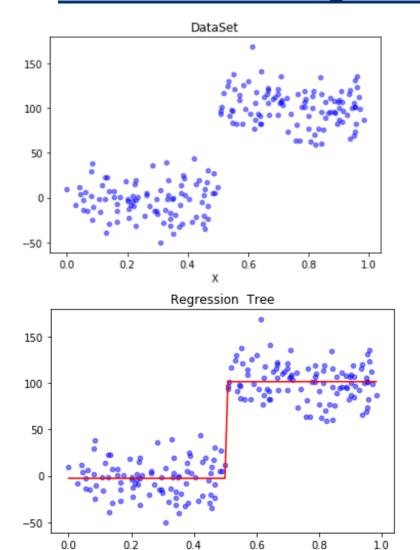


Stop Conditions:

Mind. error reduction: 4 10000

Mind. data instances: 4

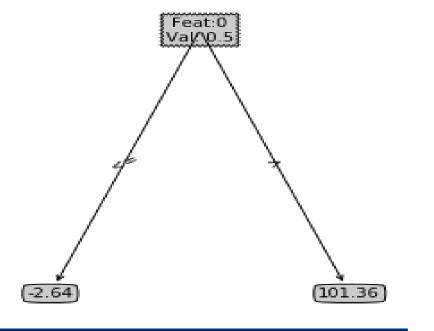




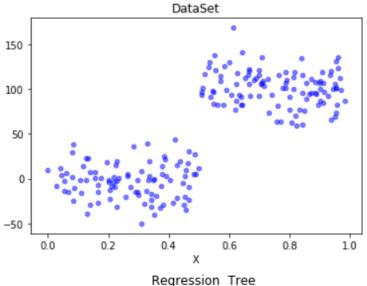
Stop Conditions:

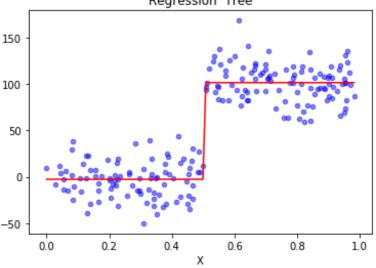
Mind. error reduction: 4 10000

Mind. data instances: 4









Stop Conditions:

Mind. error reduction: 4 10000

Mind. data instances: 4

Is there any solution without *user intervention*?

→ Postpruning

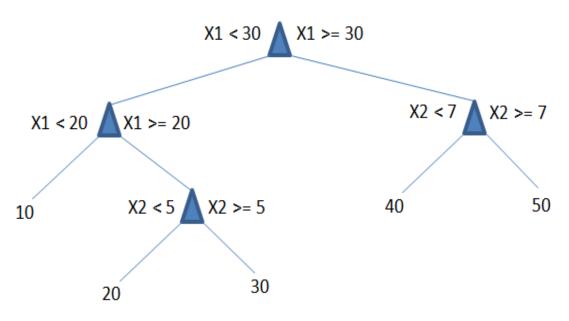




- Make the regression tree to a large depth
- Start at the bottom and combine every two leaf nodes(left and right) into a new terminal node, if such error after merge is smaller than the original one

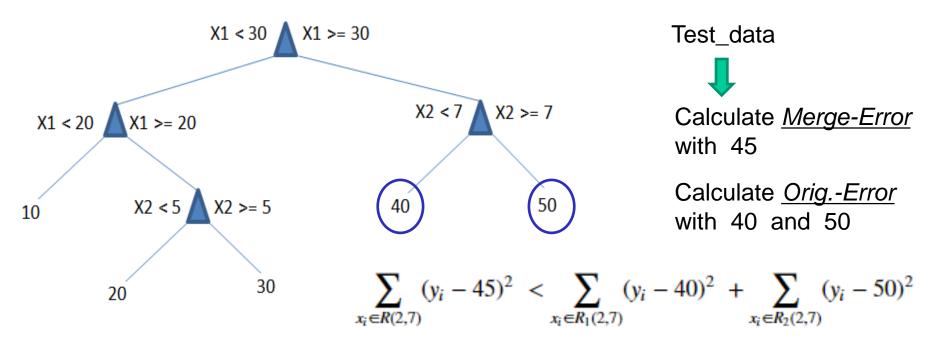


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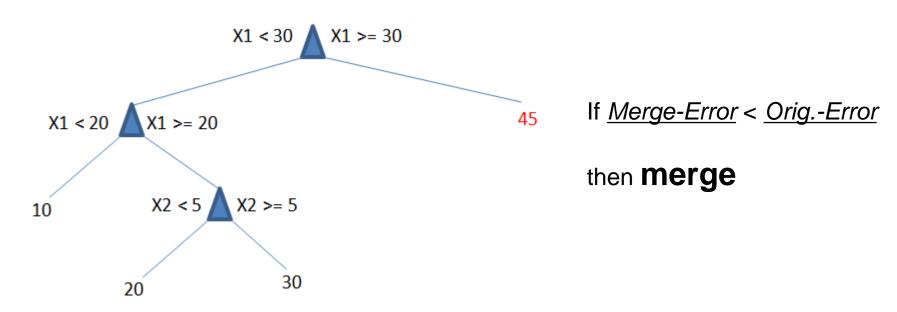


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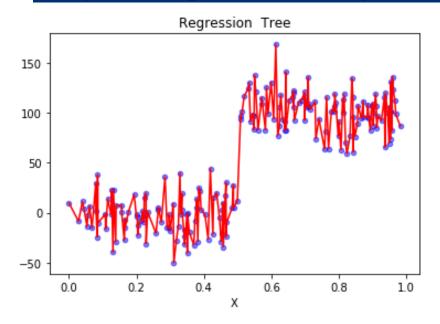


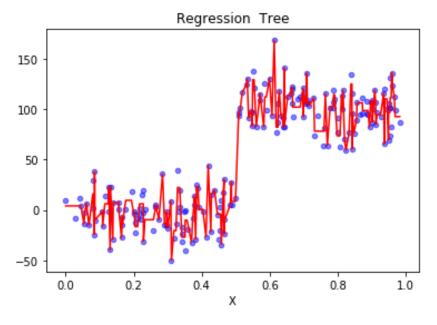
- Make the regression tree to a large depth
- Start at the bottom and combine every two leaf nodes(left and right) into a new terminal node, if such error after merge is smaller than the original one





Prepruning vs. Postpruning





Prepruning

Sum of leaf nodes: 200

tree depth: 25

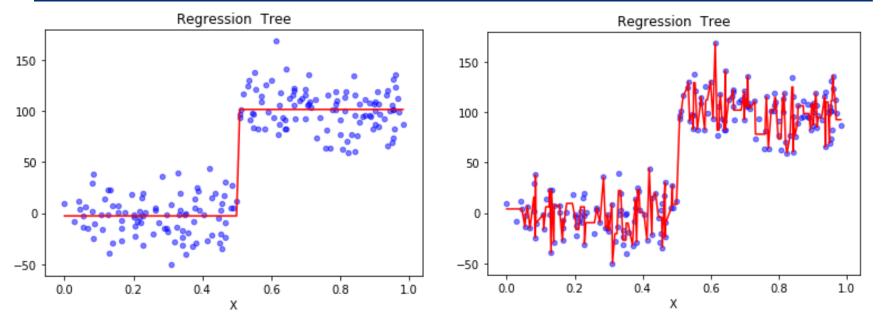
Postpruning

Sum of leaf nodes: 141

tree depth: 23



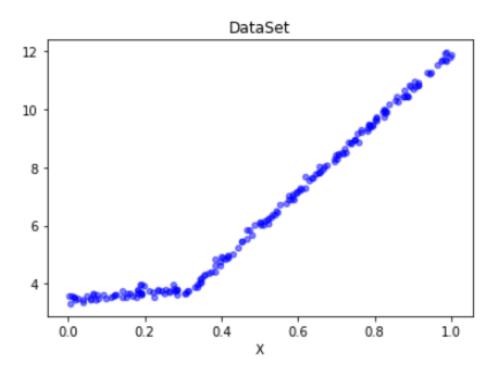
Prepruning vs. Postpruning



Some nodes were pruned off the tree, but it wasn't reduced to two nodes as we had hoped. It turns out that **postpruning** isn't as effective as **prepruning**.

We can employ **both** to give the best possible model.

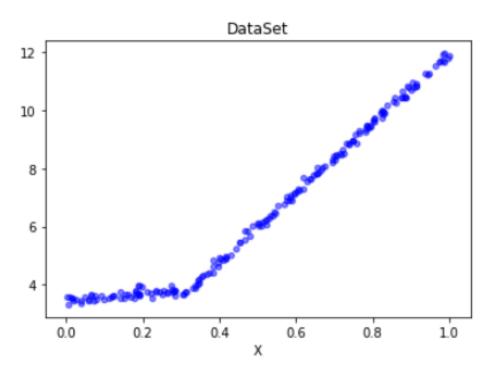




Would it be better to model this dataset as

- a bunch of constant values (many leaf nodes) or
- two straight lines (1: from <u>0.0</u> to <u>0.3</u>; 2: from <u>0.3</u> to <u>1.0</u>)

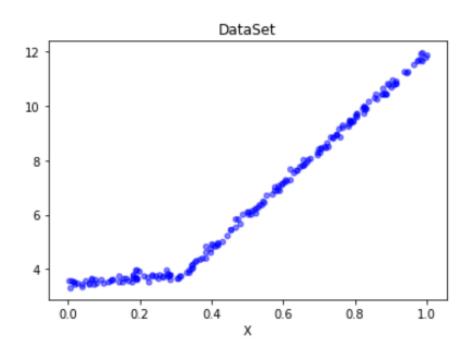




Model Tree = Tree + Linear regression

A way to model dataset as a <u>piecewise</u> linear model at each leaf node. Piecewise linear means that the model consists of multiple linear segments





Result:

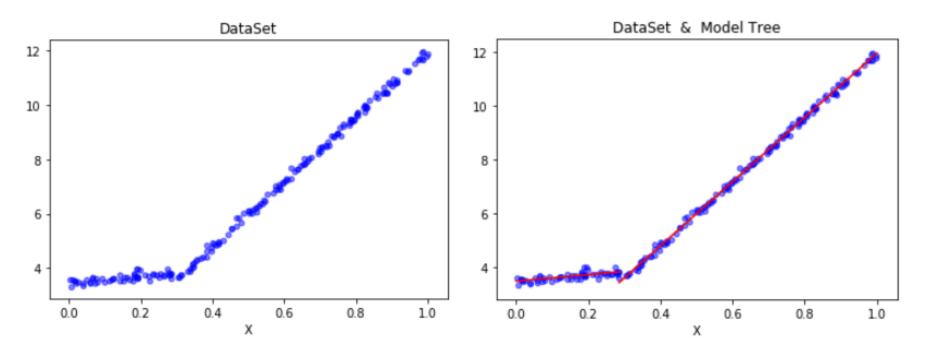
{'spInd': 0, 'spVal': 0.285477,

'left': matrix([[3.46877936], [1.18521743]]),

'right': matrix([[1.69855694e-03], [1.19647739e+01]])}

$$y = 0 + 11.96 * x$$
 and $y = 3.47 + 1.20 * x$





Result:

{'spInd': 0, 'spVal': 0.285477,

'left': matrix([[3.46877936], [1.18521743]]),

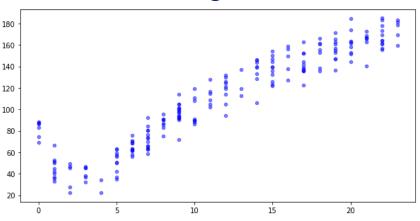
'right': matrix([[1.69855694e-03], [1.19647739e+01]])}

$$y = 0 + 11.96 * x$$
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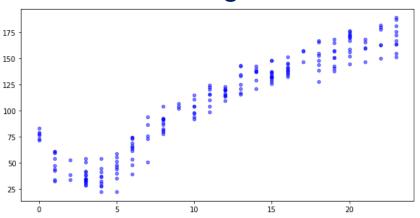


V: Linear Regression vs. Tree Regression

Training-data



Testing-data



Evaluation method:

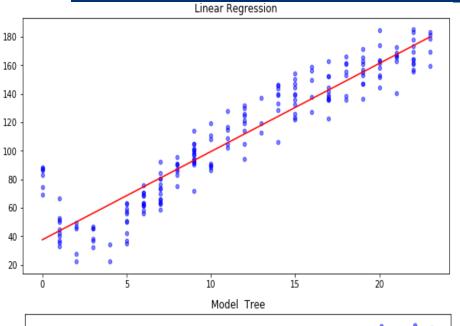
Correlation coefficients

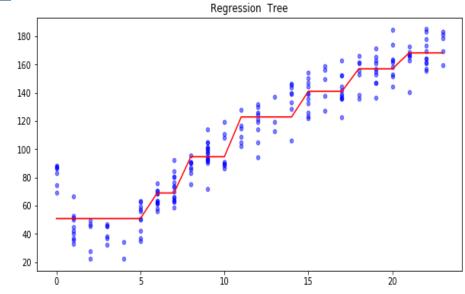
Regression methods:

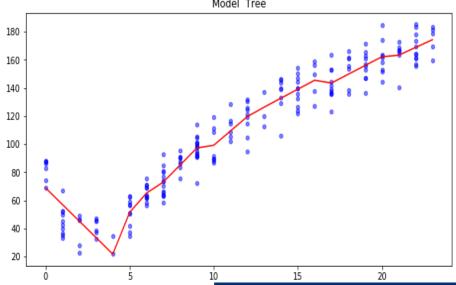
- Linear Regression(LR)
- Regression Tree (RT)
- Model Tree (MT)



V: Linear Regression vs. Tree Regression







Correlation coefficients:

• LR: 0.94346842356

• RT: 0.96408523182

MT: 0.97604121913



VI: Summary

Regression is the process of predicting a target value, which makes it become one of the most useful tools in statistics.

Linear methods:

- Linear regression: does a good job, when the dataset is simple and linear.
- Locally weighted linear regression: the forecast would be more precise.
 Overfitting problem should be avoided.

Non-linear methods:

- Regression tree: breaks up the predicted value into piecewise constant segments.
- Model Tree: implements the linear regression equations at each leaf node.
- Pre- and Postpruning: can effectively reduce the complexity of tree and help avoid the overfitting problem.



Reference

- Harrington, Peter. 2012. Machine Learning in Action. Shelter Island (N.Y.): Manning Publications Co.
- Li Hang. 2012. Statistic Learning Methods. Tsinghua University Publications Co.
- Analytics Vidhya: A Complete Tutorial on Tree Based Modeling from Scratch. APRIL 12, 2016 https://www.analyticsvidhya.com/blog/2016/04/complete-tutorial-tree-based-modeling-scratch-in-python

