Dataset: Wisconsin Breast Cancer (Original)

Applying numerous machine learning algorythms

Structure of the presentation

- ▶ What I have done?
 - ► Logistic Regression
 - ► Neural Networks
 - ► Error, bias vs variance
- ▶ What does the future brings?
 - **SVM**
 - Diagnostic
 - Prognostic

First steps

- Number of instances: 699
- Number of attributes: 10 + class atribute
 - Sample code number
 - > 9 attributes from 1 to 10 values.
 - Class attribute: 2 for benign, 4 for malignant
- Missing attributes: 16 instances
- Class distribution:
 - Benign: 458 (65.5%)
 - Malignant: 241 (34.5%)

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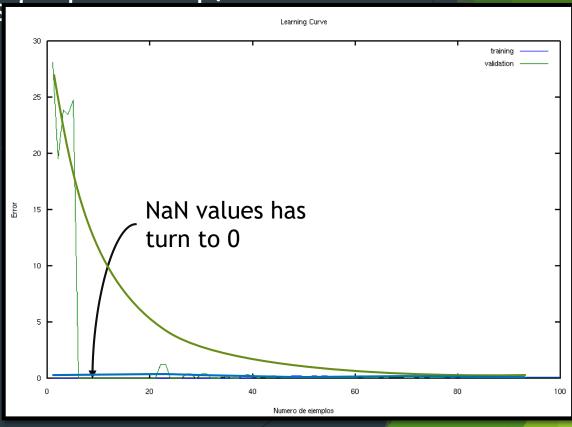
How do we work with this dataset:

- Reading the dataset file
 .data
- 2. Removing the "Sample code number attribute"
- 3. Transforming the output from 2 and 4 to 0 and 1 Y = (Y == 4);

- With the data recently read, let's try how good is our hypothesis.
- Without regularization.
- Splitting the dataset in two: 70% is for training data, 30% is for cross-validation data.

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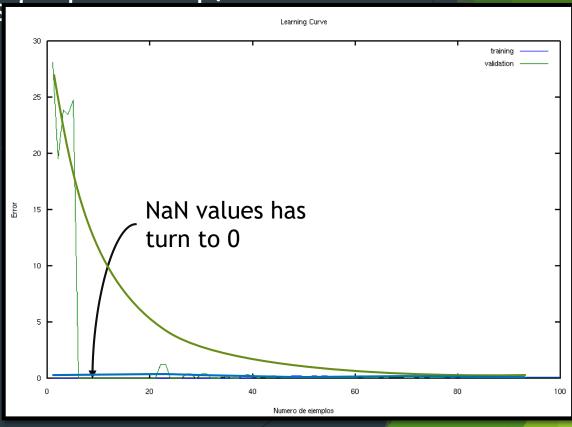
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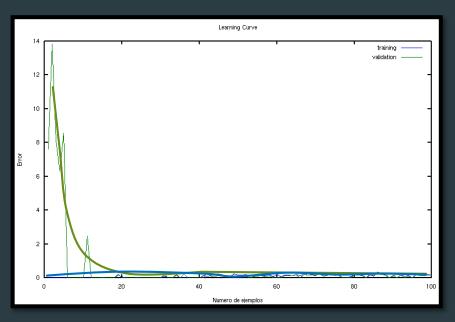
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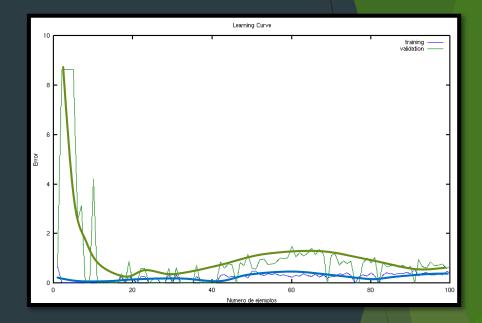
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We can do it better...



- Increasing the degree of our hypothesis (by increasing the number of features)
- Using a function that combine the features geometrically
 - ► X1,X2,X1*X2,X1^2,X2^2...





DEGREE=2

DEGREE=5

➤ You can check that an increasing in the degree of the polynomial implies a low bias between 40 and 80 dataset size, but high bias >80. (Increment of variance)

- ▶ We assume that:
 - $\rightarrow \lambda \uparrow$ fixes high bias Our Problem
 - $\rightarrow \lambda \downarrow$ fixes high variance
- Now splitting in three the dataset:
 - ► 60% Training
 - ▶ 20% Cross-Validation
 - ▶ 20% Testing

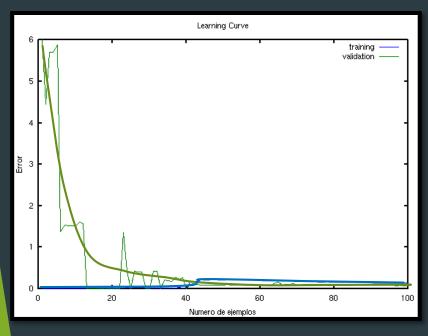
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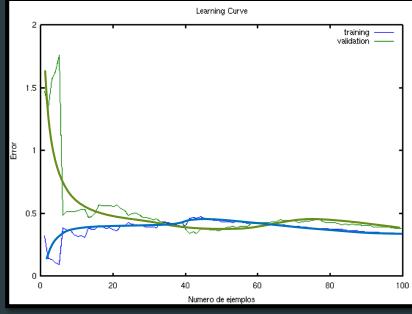
What to do:

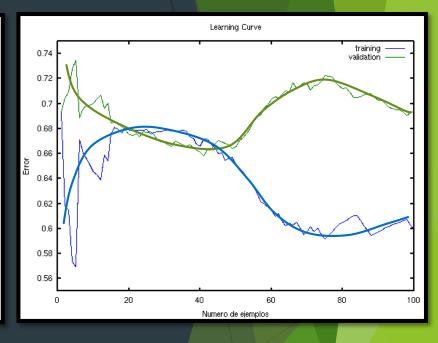
High lambda

Just right one-

Low lambda







- Low lambda (0.001)
- High bias

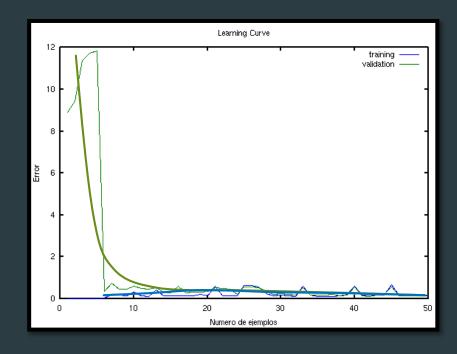
- Just right lambda(3)
- The greatest of the 100% test result

- High lambda (300)
- High variance

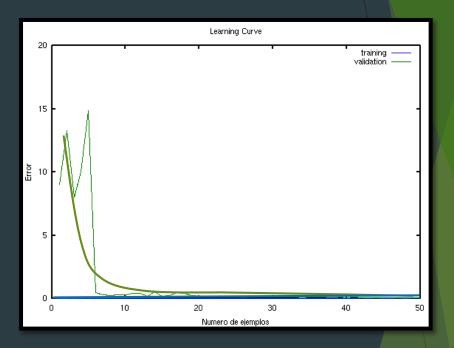
- Just one thing left... Put it all together.
 - ▶ Looking for the best relation bias-variance.
 - ► Testing diferents splittings of the data to know what kind of split its the better.
 - ► Try to diagnose, by giving an hand-made example, if is beginning or malignant.

- Looking for the right structure of our network.
 - ▶ Best number of hidden units
 - ▶ Best number of hidden layers

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Lowest hidden units



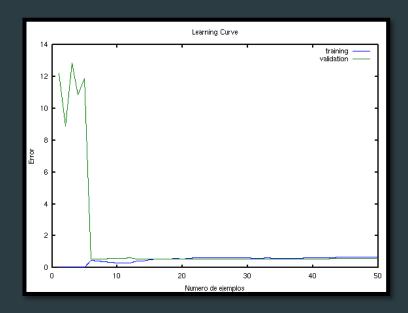
Highest hidden units

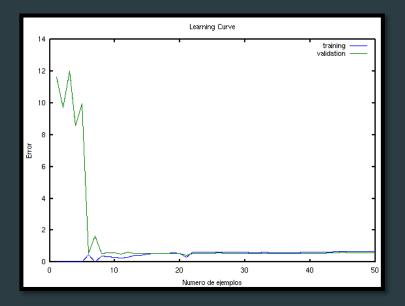


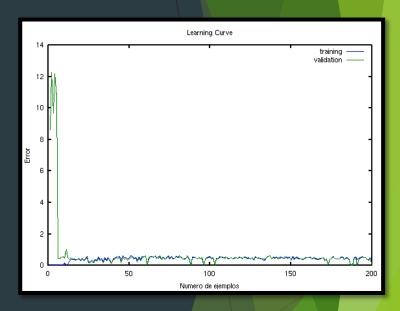
Lowest hidden units

Highest hidden units

- Adding polynomial features reduces high bias.
- ▶ Pit it all together
 - ▶ Combine more hidden units/layers with more features.
 - Degrees 2, 3, 4 and 5.
 - ▶ Hidden Units 2, 4, 10, 50 and 100.

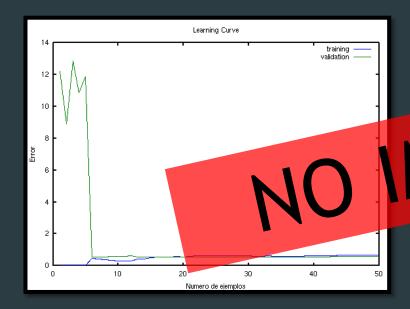


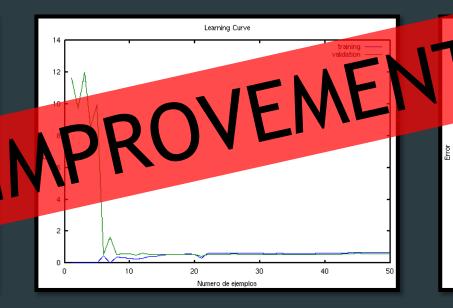




The same result as before.

More training sets





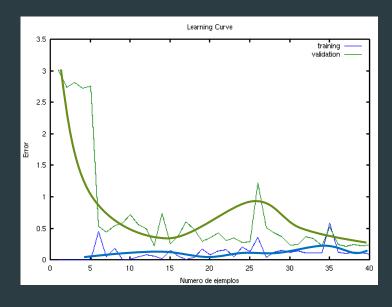
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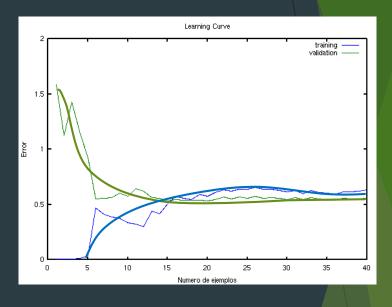
Learning Curve

Numero de ejemplos

- Only thing left is to add the regularization parameter lambda.
- Remember:
 - λ ↑ fixes high bias Our Problem
 - $\triangleright \lambda \downarrow$ fixes high variance



Lowest lambda



Highest lambda

A big difference between regularizated and non-regularizated.

Conclusions

- Regularization is far the most important thing to be aware of.
- Is positive to use polynomial adding (only if it worths).
- For every algorythm implemented, Error check is a must do.

Which is better?



What does the future brings?

- **SVM**
 - Best C Parameter.
 - ▶ Which kernel is the better.
 - Check what percentage of data makes the best model.
- Diagnostic
 - ▶ Using the same algorythms as before.
- Prognostic
 - With all the data learning, to prognose if a patient would be regresive or no regresive.
- Maybe more...