The task Examples Data exploration Model selection

### Some Project Issues

Antanas Verikas antanas.verikas@hh.se

IDE, Halmstad University

2013

#### The task

Construct a regression model

$$\widehat{y}(\mathbf{x}) = f(\mathbf{x}, \mathbf{w}) \tag{1}$$

Construct a classifier

$$\widehat{y}(\mathbf{x}) = \widehat{p}(c_i|\mathbf{x}) \tag{2}$$

#### Given

A training set

$$\mathbf{Z}_{\text{Train}} = \{\mathbf{x}(n), \mathbf{y}(n)\}_{n=1,\dots,N_{\text{Train}}}$$
(3)

- An outline for the project and report
- MATLAB
- Answers to some questions

### Not given

A test set

$$\mathbf{Z}_{\text{Test}} = \{\mathbf{x}(n), \mathbf{y}(n)\}_{n=1,\dots,N_{\text{Test}}} \tag{4}$$

- The answers to the problems.
- Infinite time.

#### Meat fat content

Estimate the fat content of a meat sample on the basis of its near infrared (NIR) absorbance spectrum.

- Regression problem,  $\hat{y}(\mathbf{x}) = f(\mathbf{x}, \mathbf{w})$
- $\mathbf{x}(n) = \mathbf{a} \ 100$  channel absorbance spectrum (D = 100).
- y(n) = measured contents of moisture, fat and protein.
- $N_{\text{Train}} = 172, N_{\text{Test}} = 43.$

Aspects: variable transforms (PCA),
variable selection,
linear and nonlinear models,
model selection,
estimating generalization performance...

### Thyroid diagnosis

Tell if a person is normal, hypothyroid, or hyperthyroid, on the basis of medical test results.

- Classification problem,  $\widehat{y}(\mathbf{x}) = \widehat{p}(c_i|\mathbf{x})$
- $\mathbf{x}(n) = 21$  variables (results from different tests), some are continuous, others are discrete (D = 21).
- $\mathbf{y}(n) = \text{class name (normal, hypothyroid, hyperthyroid)}.$
- $N_{\text{Train}} = 5000, N_{\text{Test}} = 2200.$

Aspects: variable transforms, variable selection, linear and nonlinear models, model selection, committees of models...

## Data exploration

- Scatter plots.
- Correlation coefficients.
- Histograms.
- Fisher index.
- Outlier detection.
- Data transformations:
  - Standardization,
  - PCA,
  - Box-Cox transforms.

### The 1st and the 10th components

Reveals if there are any simple relationships (But misses interactions)

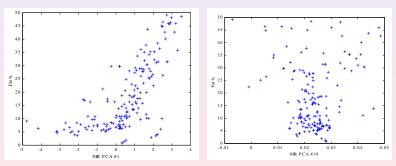


Figure: The 1st NIR principal component vs. the fat % in the meat (*left*). The 10th NIR principal component shows no similar relationship.

### PCA #1-9 vs. fat %

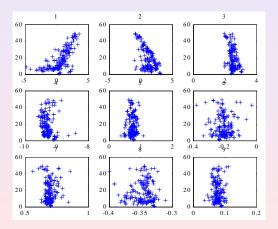


Figure: Scatter plots for PCA #1-9 vs. fat %

### The 1st and the 10th components

Pearson's:  $r_{xy} = \sigma_{xy}/\sqrt{\sigma_{xx}\sigma_{yy}} \in [-1,1]$ . Spearman's:  $(s_{xy})$  uses ranks instead of values.

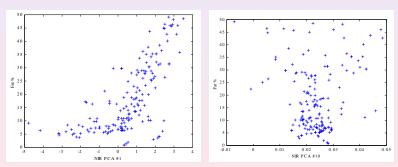


Figure: Left:  $r_{xy} = 0.68$ ,  $s_{xy} = 0.70$ . Right:  $r_{xy} = 0.04$ ,  $s_{xy} = -0.01$ .

### PCA #1-9 vs. fat %

$$r_{xy} = \begin{pmatrix} 0.6798 & -0.5763 & -0.3944 \\ -0.1439 & 0.0586 & -0.0233 \\ 0.0062 & 0.0289 & -0.0187 \end{pmatrix}$$

$$s_{xy} = \begin{pmatrix} 0.6996 & -0.4689 & -0.3276 \\ -0.2620 & 0.0806 & -0.0363 \\ 0.0172 & 0.0079 & -0.0750 \end{pmatrix}$$

If  $|r_{xy}|, |s_{xy}| > 1.96/\sqrt{N}$ , the correlation is significant at the 95% level.

$$N = 172 \Rightarrow 1.96/\sqrt{N} = 0.1494$$

# Histograms (1)

Useful in classification problems to detect class specific distribution.

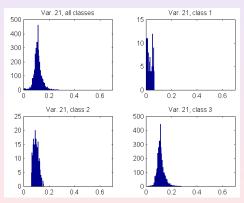


Figure: Class-specific distributions.

# Histograms (2)

Detect non-normal distributions and test transformations.

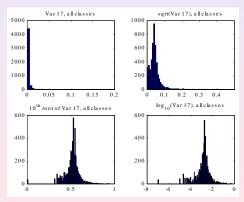


Figure: Affect of different transformations.

#### Standardization

Useful in classification problems to detect class specific distribution.

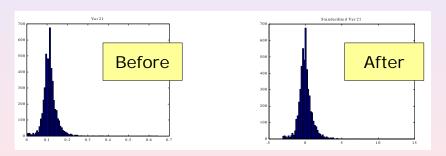


Figure: Affect of standardization.

### Whitening

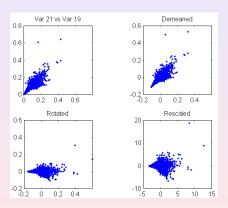
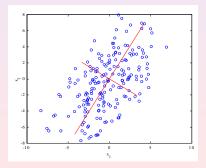


Figure: Top-left: **X**, Top-right:  $\mathbf{X} - \mathbf{1}\mu^T$ , Bottom-left:  $(\mathbf{X} - \mathbf{1}\mu^T)\mathbf{Q}$ , Bottom-right:  $(\mathbf{X} - \mathbf{1}\mu^T)\mathbf{Q}\mathbf{\Lambda}^{-1/2}$ .

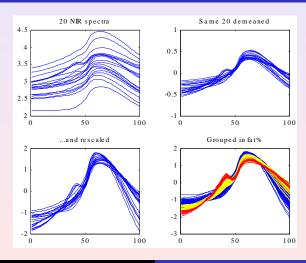
### Principal components

Express the data in the new basis  $\mathbf{Q}$ , with eigenvectors of the data covariance matrix  $\mathbf{\Sigma}$  as basis vectors.

$$\mathbf{\Sigma}\mathbf{q}_i = \lambda_i \mathbf{q}_i \tag{5}$$

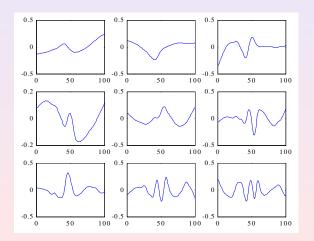


# PCA example: NIR spectra



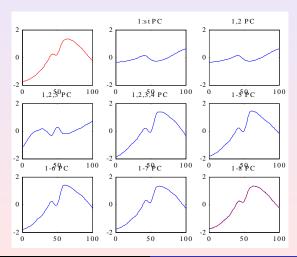
Scatter plots
Correlation coefficients
Histograms
Standardization
Whitening
Principal components

### NIR: The 9 leading eigenvectors



$\lambda_i$
2.4308
0.9372
0.0489
0.0256
0.0108
0.0023
0.0014
0.0002
0.0001

#### NIR reconstruction with PCA



### Comparing regression models

- Standard t-test, if E or log(E) are (approximately) normally distributed.
- Non-parametric Wilcoxon test, if E or log(E) are not normally distributed.
- Paired t—test, if we test models A and B on the same data, and residuals e(n) are normally distributed.

# Comparing regression models: Example (1)

- Estimate mean and variance for the generalization error (or log error) using K-fold cross validation. log(E) is often more normal than E.
- Compare means using t—test (or Wilcoxon test):

$$|\mu_A - \mu_B| > 1.96 \frac{\sigma_{AB}}{\sqrt{K}}$$

$E_{\mathtt{Test,A}}$	$E_{\mathtt{Test,b}}$
0.6488	3.3016
0.1891	0.6465
1.1335	5.0245
1.3333	12.473
0.3178	1.2315
3.2901	5.8008
3.2843	8.6193
0.9631	0.4997
1.3872	0.5822
1.1908	4.3542

# Comparing regression models: Example (2)

$$\mu_{A} = \frac{1}{10} \sum_{k=1}^{10} \log(E_{k,A}) = 0.0013$$

$$\mu_{B} = \frac{1}{10} \sum_{k=1}^{10} \log(E_{k,B}) = 0.9253$$

$$\sigma_{A} = \sqrt{\frac{1}{9} \sum_{k=1}^{10} [\log(E_{k,A}) - \mu_{A}]^{2}} = 0.9034$$

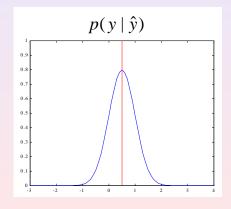
$$\sigma_{B} = \sqrt{\frac{1}{9} \sum_{k=1}^{10} [\log(E_{k,B}) - \mu_{B}]^{2}} = 1.1898$$

# Comparing regression models: Example (3)

$$\begin{array}{rcl} \sigma_{AB} & = & \sqrt{\sigma_A^2 + \sigma_B^2} = 1.4940 \\ |\mu_A - \mu_B| & = & 0.9240 < 1.96 \frac{\sigma_{AB}}{\sqrt{10}} = 0.9260 \\ & \Rightarrow & \text{Not significant difference!} \end{array}$$

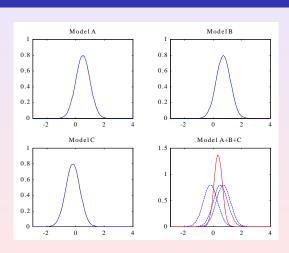
## Interpreting the output: Regression

$$\begin{array}{lcl} \widehat{y}(\mathbf{x}) & = & \langle y(\mathbf{x}) \rangle_{\varepsilon} \\ \sigma_{\varepsilon}^{2} & \approx & \mathtt{MSE}_{\mathtt{Test}} \\ y(\mathbf{x}) & = & \widehat{y}(\mathbf{x}) \pm 1.96 \sqrt{\mathtt{MSE}_{\mathtt{Test}}} \end{array}$$



### Combining models

$$\widehat{y}_{\text{com}} = \frac{\sum_{k} \widehat{y}_{k} / \text{MSE}_{k}}{\sum_{k} 1 / \text{MSE}_{k}}$$



## Comparing classification models

- Use K-fold cross-validation method.
- Use McNemar's test for paired testing

$$z = \frac{|n_A - n_B| - 1}{\sqrt{n_A + n_b}}$$

 $n_A$  = mistakes made by A but not B and vice versa.

If z > 1.96 then we have a significant difference at the 95% confidence level.

#### Error bars

Error bars on the classification error:

$$R = \widehat{R} \pm 1.96 \sqrt{\frac{\widehat{R}(1-\widehat{R})}{N}}$$

 $\widehat{R}=$  estimate of the classification error on out-of-sample test data.