Classification Logistic Model

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# ISEFAR http://fermin.perso.math.cnrs.fr/

- k Nearest-Neighbors  $\checkmark$
- $\textcircled{O} \ {\rm K-Fold \ cross \ validation, \ Model \ selection \ \checkmark}$
- ${f 0}$  Generative Modeling (Naive Bayes, LDA, QDA)  $\checkmark$
- Logistic Modeling
- SVM
- Tree Based Methods (M. Zetlaoui course, Apprentissage)

• Direct modeling of Y|x.

The Binary logistic model  $(Y \in \{-1,1\})$ 

$$p_{+1}(\mathsf{x}) = rac{e^{eta^t arphi(\mathsf{x})}}{1+e^{eta^t arphi(\mathsf{x})}}$$

where  $\varphi(x)$  is a transformation of the individual **x** 

- In this model, one verifies that  $p_{+1}(\mathbf{x}) \ge p_{-1}(\mathbf{x}) \quad \Leftrightarrow \quad \beta^t \varphi(\mathbf{x}) \ge 0$
- True Y|x may not belong to this model  $\Rightarrow$  maximum likelihood of  $\beta$  only finds a good approximation!
- Binary Logistic classifier:

$$\widehat{f}_L(\mathbf{x}) = egin{cases} +1 & ext{if } \widehat{eta}^t arphi(\mathbf{x}) \geq 0 \ -1 & ext{otherwise} \end{cases}$$

where  $\widehat{\beta}$  is estimated by maximum likelihood.

#### Logistic Modeling

• Logistic model: approximation of  $\mathcal{B}(p_1(\mathbf{x}))$  by  $\mathcal{B}(h(\beta^t \varphi(\mathbf{x})))$  with  $h(t) = \frac{e^t}{1+e^t}$ .

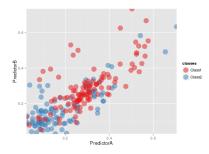
## Opposite of the log-likelihood formula

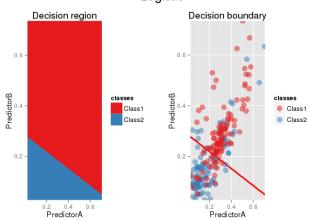
$$\begin{aligned} &-\frac{1}{n}\sum_{i=1}^{n}\left(\mathbf{1}_{y_{i}=1}\log(h(\beta^{t}\varphi(\mathbf{x})))+\mathbf{1}_{y_{i}=-1}\log(1-h(\beta^{t}\varphi(\mathbf{x})))\right)\\ &=-\frac{1}{n}\sum_{i=1}^{n}\left(\mathbf{1}_{y_{i}=1}\log\frac{e^{\beta^{t}\varphi(\mathbf{x})}}{1+e^{\beta^{t}\varphi(\mathbf{x})}}+\mathbf{1}_{y_{i}=-1}\log\frac{1}{1+e^{\beta^{t}\varphi(\mathbf{x})}}\right)\\ &=\frac{1}{n}\sum_{i=1}^{n}\log\left(1+e^{-y_{i}(\beta^{t}\varphi(\mathbf{x}))}\right)\end{aligned}$$

- Convex function in  $\beta$ !
- **Remark:** You can also use your favorite parametric model instead of the logistic one...

## Synthetic Dataset

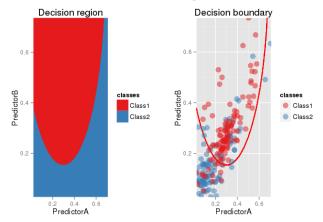
- Two features/covariates.
- Two classes.
- Dataset from *Applied Predictive Modeling*, M. Kuhn and K. Johnson, Springer
- Numerical experiments with R.





#### Logistic

### Quadratic Logistic







Given Y a variable to explain by d variables X<sub>1</sub>,..., X<sub>d</sub>, how to select (systematically) the most interesting subset of variables to do the prediction?

Variable selection

Find automatically a sub-group of variables to explain Y.

More generaly, given k models M<sub>1</sub>,..., M<sub>k</sub>, which one to use?

Model selection

Criterion to compare the performance of different models.

- Assume we have two competing models  $\mathcal{M}_p$  (with p parameters) and  $\mathcal{M}_q$  (with q parameters) such that  $\mathcal{M}_p \subset \mathcal{M}_q$ .
- Can we test if  $\mathcal{M}_p$  is sufficient?

### Example (p=2 and q=4)

- Models:
  - $\mathcal{M}_p$  : logit $p_{\beta}^{(p)}(x) = \beta_0 + \beta_1 X_1 + \beta_2 X_2$
  - $\mathcal{M}_q$ : logit $p_{\beta}^{(q)}(x) = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \beta_4 X_4$

Test

$$H_0: \beta_3 = \beta_4 = 0$$
 against  $H_1: \beta_3 \neq 0$  or  $\beta_4 \neq 0$ 

#### Deviance (residual deviance sous R)

- Log-likelihood  $\mathcal{M}_p$ :  $\mathcal{L}_p = \log(L_p(\hat{\beta}, \mathcal{D}_n))$
- Log-likelihood of model  $\mathcal{M}_q$ :  $\mathcal{L}_q = \log(L_q(\hat{\beta}, \mathcal{D}_n))$
- Deviance between the two models:

$$\mathcal{D}_{q-p} = 2(\mathcal{L}_q - \mathcal{L}_p) = 2(\log(\mathcal{L}_q(\hat{\beta}, \mathcal{D}_n)) - \log(\mathcal{L}_p(\hat{\beta}, \mathcal{D}_n)))$$

Asymptotically under  $H_0$  :  $(\mathcal{D}_{q-p}) \sim \chi^2(q-p)$ 

Under R : If W and V are two objects obtained with glm such that W is a submodel of V, the command

```
anova(W,V,test="Chisq")
```

performs this test.

- Let  $\mathcal{M}$  be a generic logistic model and denote p its number of parameters.
- Let  $\hat{\beta}$  be the ML estimate in this model  $\mathcal{M}$ .
- The AIC and BIC consist in minimizing

$$-2 imes \log(L(\hat{eta}, \mathcal{D}_n)) + \kappa(n) imes p$$

over all models.

- Different choices for the factor κ(n):
  - AIC :  $\kappa(n) = 2$ .
  - BIC :  $\kappa(n) = \log n$ .
- The BIC criterion leads to the selection of a model with a smaller dimension than AIC.





### Prediction for a new individual:

- A new individual x<sub>new</sub> appears and we want to predict if he has the disease (y<sub>new</sub> = 1) or not (y<sub>new</sub> = 0).
- $\bullet$  We have estimated the logistic coefficients  $\widehat{\beta}$  so that

$$\mathbb{P}\left\{Y=1|X\right\}\approx\frac{e^{\widehat{\beta}^{T}\mathbf{X}}}{1+e^{\widehat{\beta}^{T}\mathbf{X}}}.$$
(1)

• Any ideas to predict y<sub>new</sub>?

Prediction (threshold = 0.5)

Then,

- If  $\widehat{\mathbb{P}}(y = \text{Yes}|X) > 0.5$ , we predict y = Yes;
- If  $\widehat{\mathbb{P}}(y = \text{Yes}|X) \leq 0.5$ , we predict y = No.

### Confusion Matrix : Cross table of the prediction vs the truth

##	pred		
##	CHD	No	Yes
##	No	45	12
##	Yes	14	29

**Prediction error** : (14 + 12)/100 = 0.26



Score

$$\mathsf{Accuracy} = \frac{\mathsf{TP} + \mathsf{TN}}{\mathsf{TP} + \mathsf{TN} + \mathsf{FP} + \mathsf{FN}}$$

#### Scores

$$\mathsf{Accuracy} = \frac{\mathsf{TP} + \mathsf{TN}}{\mathsf{TP} + \mathsf{TN} + \mathsf{FP} + \mathsf{FN}}$$

 $\left. \begin{array}{c} \mbox{True Positive Rate} \\ \mbox{Sensitivity} \\ \mbox{Recall} \end{array} \right\} = \frac{\mbox{TP}}{\mbox{\#(real P)}} = \frac{\mbox{TP}}{\mbox{TP} + \mbox{FN}}$ 

To label digits:

- True label  $y_i \in \{0, ..., 9\}$ ,
- Predicted label  $\widehat{y_i} \in \{0, \dots, 9\}$ ,
- Confusion matrix is thus of size  $10 \times 10$ .

#### Confusion matrix

