

# Introduction to Computational Medicine I

## Project 2: Lecture 1

Objective: to construct predictive models to classify  $\{\text{sepsis}, \text{nonsepsis}\}$  from EHR and numerical waveforms

Notation:  $Y_i \in \{0, 1\}$  is label for patient  $i$  where sepsis = 1

$X_i = \begin{pmatrix} x_1^i \\ x_2^i \\ \vdots \\ x_m^i \end{pmatrix}$  is feature vector for patient  $i$  (eg. age, gender)

Problem: Given data set  $\{(X_i, Y_i)\}_{i=1}^N$  where  $Y_i$  are independent samples of a Bernoulli R.V., construct/estimate the PMF

$$\Pr(Y_i = y_0) = (p_i)^{y_0} (1 - p_i)^{1 - y_0} \quad y_0 = 0, 1$$

where  $p_i = f(X_i; \beta)$

Idea: We want to pick  $f$  and  $\beta$  such that  $\Pr(Y_1 = y_1, Y_2 = y_2, \dots, Y_N = y_N)$  is "high".  
so why not pick  $f$  &  $\beta$  to maximize this quantity?

(2)

In statistics, this is called the likelihood function. It is a function of parameters of a statistical model (PMF or PDF) given data. In our case,

$$L(f, \beta) = \Pr(Y_1=y_1, Y_2=y_2, \dots, Y_N=y_N; f; \beta)$$

it is not  
"Probability"  
or say, it is a  
nc of a parameter  
for a given  
observed  
outcome!

$$= \prod_{i=1}^N (1-p_i)^{y_i} p_i^{1-y_i}$$

$$= \prod_{i=1}^N (p_i)^{y_i} (1-p_i)^{1-y_i}$$

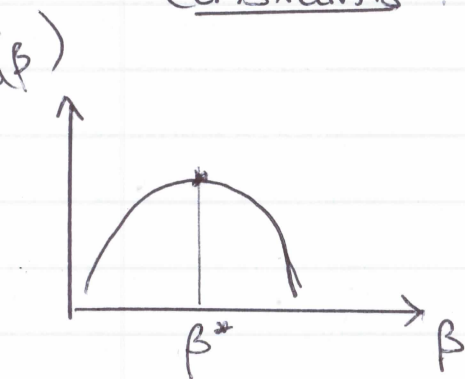
$$= \prod_{i=1}^N (f(x_i; \beta))^{y_i} (1-p_i)^{1-y_i}$$

Q: Why don't we fix  $f(\cdot)$ -parametrized class of functions?

Constraints : (a) need  $0 \leq f(x) \leq 1$

(ii) maximizing  $L$  should be doable

(iii)  $L$  should be concave in  $\beta$

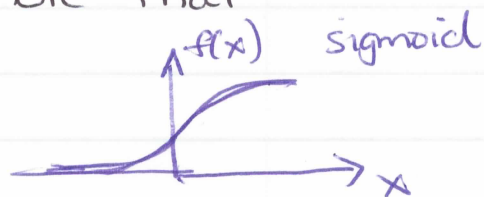


(3)

candidate:  $f(x) = \frac{e^{\beta^T x}}{1 + e^{\beta^T x}}$  satisfies (i)

Does it satisfy (iii)? First note that

$$\log\left(\frac{f(\cdot)}{1-f(\cdot)}\right) = \beta^T x$$



$$\begin{aligned}\log(f) &= \beta^T x - \log(1 + e^{\beta^T x}) \\ \log(1-f) &= -\log(1 + e^{\beta^T x})\end{aligned}$$

$$\Rightarrow \log(f) - \log(1-f) = \beta^T x \quad \checkmark \quad \text{Back to } L(\cdot)$$

$$L(\beta) = \prod_{i=1}^N (f)^{y_i} (1-f)^{1-y_i}$$

$$\Rightarrow \log L(\beta) \triangleq \ell(\beta) = \sum_{i=1}^N y_i \log(f) + (1-y_i) \log(1-f)$$

$$\Rightarrow \ell(\beta) = \sum_{i=1}^N y_i \log\left(\frac{f}{1-f}\right) + \log(1-f)$$

$$\ell(\beta) = \sum_{i=1}^N y_i \beta^T x + \log(1-f(\beta))$$


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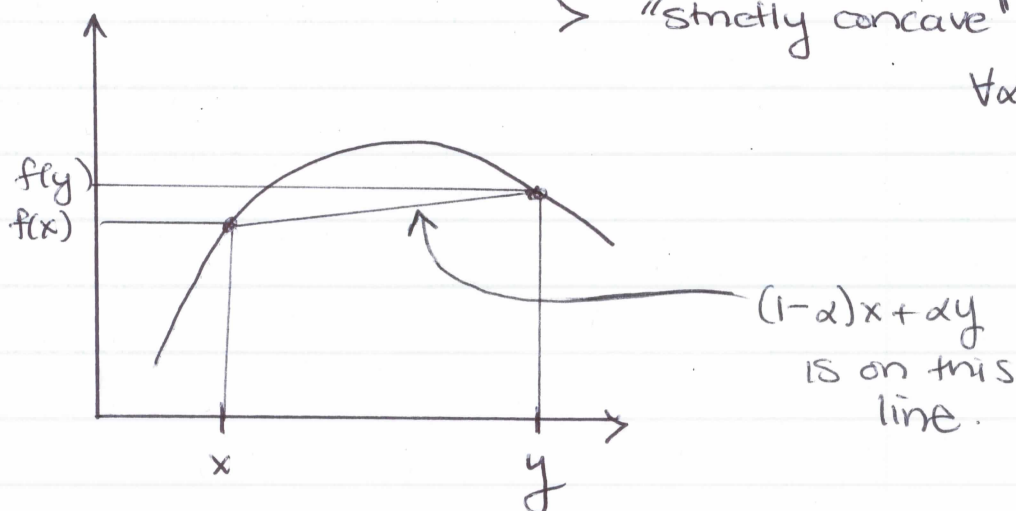
(4)

Concave functions:  $f$  is concave if

$$f((1-\alpha)x + \alpha y) \geq (1-\alpha)f(x) + \alpha f(y)$$

> "strictly concave"

$\forall \alpha \in (0,1)$



Fact: if  $f$  is differentiable, then

$$f \text{ concave} \stackrel{\text{iff}}{\iff} f'' \leq 0 \quad \forall x$$

So let's use this fact to check if  $\ell(\beta)$  is concave...

$$\ell'(\beta) \triangleq \frac{\partial \ell}{\partial \beta_j} = \sum_{i=1}^N y_i x_j - \frac{x_j e^{\beta^T x}}{(1 + e^{\beta^T x})} \quad \forall j$$

$$\frac{\partial^2 \ell}{\partial \beta_j^2} = -x_j^2 \frac{e^{\beta^T x}}{(1 + e^{\beta^T x})^2} \leq 0 \quad \forall x_j$$

$\Rightarrow \ell(\beta)$  is concave with our choice of  $f(\cdot)$  !!

Turns out it is very efficient to compute  $\beta^*$  where

$$\beta^* = \underset{\beta}{\operatorname{argmax}} \ell(\beta)$$

Note: We will use "glmfit" in MATLAB.

OK - so now we know how to construct

$$\hat{p}_i = \frac{e^{\beta^{*T} x_i}}{1 + e^{\beta^{*T} x_i}} \quad i=1, 2, \dots, N$$

Q: How do we use  $\hat{p}_i$  to classify patient  $i$ ?

A: if  $p_i > \theta$  then decide/classify as  
sepsis

else classify as nonsepsis

A simple threshold rule!!

Q: How can we evaluate our classifier?

(6)

		Predicted	
		S	NS
Actual	S	TP	FN
	NS	FP	TN

TP = true positive

TN = true negative

FP = false positive

FN = false negative

$$\text{Sensitivity} \triangleq \text{TPR} = \frac{\text{TP}}{\text{TP} + \text{FN}}$$

$$\text{Specificity} \triangleq \text{TNR} = \frac{\text{TN}}{\text{FP} + \text{TN}}$$

