### 4.4 Generative Models for Classification

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#### The Need for Alternatives

#### Why not just use logistic regression?

- If there is a lot of separation between the classes, logistic regression models are surprisingly unstable.
  - ▶ (Coefficient estimates can vary significantly given the same data generating process.)
- If the distribution of the predictors X is approx. normal and the sample size is small, these alternatives may be more accurate than logistic regression.
- ▶ The methods in this section have more natural extensions to three or more classes.

#### Idea

- ▶ Model the distribution of the predictors *X* separately for each response class.
- ▶ Use Bayes' theorem to work these into estimates for P(Y = k | X = x).

#### The Setup

Suppose Y can take on K distinct, unordered values.

- Let  $\pi_k$  represent the overall probability that a randomly chosen observation comes from the kth class.
  - $\triangleright$  Generally estimated as the proportion of training observations belonging to class k.
- Let  $f_k(x) = P(X|Y = k)$  denote the density function of X for an observation from the kth class.
  - ▶ So  $f_k(x)$  should be relatively large if there is a high probability that an observation from the kth class has  $X \approx x$ .
- ► Then Bayes' Theorem states

$$p_k(x) = P(Y = k | X = x) = \frac{\pi_k f_k(x)}{\sum_{l=i}^{K} \pi_l f_l(x)}$$

This is the *posterior probability* that an observation belongs to the kth class, given X = x.

Goal: estimate  $f_k(x)$  to approximate the Bayes' classifier  $p_k(x)$ .

We will classify an observation into the category for which  $p_k(x) = P(Y = k | X = x)$  is greatest.

Assume  $f_k(x)$  is normally distributed (Gaussian):

$$f(x) = \frac{1}{\sqrt{2\pi\sigma_k^2}} \exp\left[-\frac{(x-\mu_k)^2}{2\sigma_k^2}\right]$$

where  $\mu_k$  and  $\sigma_k^2$  are the mean and standard deviation parameters for the kth class.

Also assume  $\sigma_1^2 = \sigma_2^2 = \cdots = \sigma_K^2 = \sigma^2$  (shared variance term for all classes).

Combining the Bayes' Theorem set up with these assumptions, we get

$$p_k(x) = \frac{\frac{\pi_k}{\sqrt{2\pi\sigma^2}} \exp\left[-\frac{1}{2\sigma^2}(x - \mu_k)^2\right]}{\sum_{l=i}^K \frac{\pi_l}{\sqrt{2\pi\sigma^2}} \exp\left[-\frac{1}{2\sigma^2}(x - \mu_l)^2\right]}$$

which looks a mess, but it can be shown this is equivalent to assigning the observation to the class for which

$$\delta_k(x) = x \left(\frac{\mu_k}{\sigma^2}\right) - \frac{\mu_k^2}{2\sigma^2} + \log(\pi_k)$$

is largest.

$$\delta_k(x) = x \left(\frac{\mu_k}{\sigma^2}\right) - \frac{\mu_k^2}{2\sigma^2} + \log(\pi_k)$$

If K=2 and  $\pi_1=\pi_2$ , this classifier assigns an observation to

- ightharpoonup class 1 if  $2x(\mu_1 \mu_2) > \mu_1 \mu_2$ .
- class 2 otherwise.

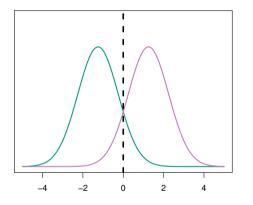
The Bayes' decision boundary is the point for which  $\delta_1=\delta_2$ , which in this setting is

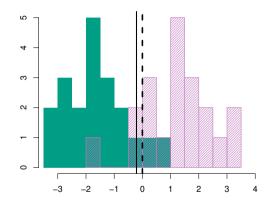
$$x = \frac{\mu_1^2 - \mu_2^2}{2(\mu_1 - \mu_2)} = \frac{\mu_1 + \mu_2}{2}$$

### Example

- Consider predictors generated from two normal distributions where  $mu_1 = -1.25$ ,  $\mu_2 = 1.25$ , and  $\sigma_1 = \sigma_2 = 1$ .
- Assume an observation is equally likely to come from either class, i.e.,  $\pi_1 = \pi_2 = 0.5$ .
- Then the (known) Bayes' classifier assigns an observation to class 1 if x < 0 and class 2 otherwise.

# Example





- ▶ 20 observations drawn from each class.
- ▶ LDA decision boundary shown as solid vertical line.

In practice, we must estimate  $\mu_1, \ldots, \mu_K$ ,  $\pi_1, \ldots, \pi_K$ , and  $\sigma$ .

$$\hat{\mu}_k = \frac{1}{n_k} \sum_{i:y_i = k} x_i$$

$$\hat{\sigma}^2 = \frac{1}{n - K} \sum_{k=1}^K \sum_{i:y_i = k} (x_i - \hat{\mu}_k)^2$$

$$\hat{\pi}_k = \frac{n_k}{n}$$

Where n is the number of training observations and  $n_k$  is the number of training observations in the kth class.

 $\hat{\sigma}^2$  is a weighted average of sample variances across the K classes.

Assign an observation X = x to the class for which

$$\delta_k(x) = x \left(\frac{\hat{\mu}_k}{\hat{\sigma}^2}\right) - \frac{\hat{\mu}_k^2}{2\hat{\sigma}^2} + \log(\hat{\pi}_k)$$

is largest.

# Example: Using Penguin Body Mass to Predict Species

```
data(penguins, package = "palmerpenguins")
mod1 <- lda(species ~ body_mass_g, penguins)</pre>
predval <- predict(mod1)$class</pre>
species <- penguins$species[!is.na(penguins$species) & !is.na(penguins$bod
table(predval, species)
##
              species
## predval Adelie Chinstrap Gentoo
##
    Adelie
                  140
                             64
                                    14
## Chinstrap
##
     Gentoo
               11
                                   109
mean(predval == species)
## [1] 0.7280702
```