Exercise 2



Before we vary the decision threshold, lets see if we can find the default 50% threshold in our model.

To do that, let's look at the posterior class probabilities and count how many of them are higher than 50%. These are the observations that are classified YES.

We plot these probabilties together with true class values and compare it with the plot p1 from exercise 1 – just to see if we are right.

The plots also help us to see whats going on after we will have changed the threshold in the next step.

Varying the threshold

Try for yourself!

Inspect the code and plot, and try to interpret it!

Data Science Dr. Gwendolin Wilke 27

Exercies 2



Now we change the threshold from 50% to 20% - we have to do this manually:

- First we find all the observations with posterior probability of >=0.2, and see how many we have here.
- We then manually reclassify them: We create a new variable lda.reclassified, that we use instead of lda.pred\$class. (lda.pred\$class is where the predict function had stored the class predictions in exercise 1.)
- We set it to "No" for all observations except the ones with posterior >=0.2.
- We then do the same plots as before, but with the manually reclassified

```
# Imposing a lower threshold for Yes
sum(lda.pred$posterior[,2] >= 0.2) # how many observations are classified Yes with a 20% threshold
# reclassify
    lda.reclassified <- rep("No", length(lda.class))
    lda.reclassified[lda.pred$posterior[,2] >= 0.2] <- "Yes"
# Plotting the new classification
    lda.reclassified.df <- data.frame(balance=test.data$balance,lda.reclassified=lda.reclassified) # make a data frame for plotting
    p3 <- ggplot() + geom_point(data = lda.reclassified.df, aes(x=balance, y=lda.reclassified, col=test.data$default), size=5)
    lda.prob.df <- data.frame(balance=test.data$balance,lda.prob=lda.pred$posterior[,2]) # make a data frame for plotting
    p4 <- ggplot() + geom_point(data = lda.prob.df, aes(x=balance, y=lda.prob, col=test.data$default), size=5) +
        geom_hline(yintercept = 0) + geom_hline(yintercept = 1) + geom_hline(yintercept = 0.2, linetype="dashed") + ylim(0,1)
    grid.arrange(p3, p4, nrow = 1)
    par(mfrow=c(1,1))</pre>
```

Try for yourself and inspect the plots!

Particularly, compare the plots produced now with the plots produced with the original class predictions of step 1.

Data Science Dr. Gwendolin Wilke 28