Applied Data Science II - Homework 5

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Libraries

```
library(tidyverse)
## -- Attaching packages ------ tidyverse 1.3.1 --
## v ggplot2 3.3.5 v purrr
                                0.3.4
## v tibble 3.1.6 v dplyr 1.0.7
## v tidyr 1.1.4 v stringr 1.4.0
## v readr 2.1.1 v forcats 0.5.1
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
library(MASS)
##
## Attaching package: 'MASS'
## The following object is masked from 'package:dplyr':
##
##
       select
library(nnet)
library(caret)
## Loading required package: lattice
##
## Attaching package: 'caret'
```

```
## The following object is masked from 'package:purrr':
##
##
       lift
library(glmnet)
## Loading required package: Matrix
##
## Attaching package: 'Matrix'
## The following objects are masked from 'package:tidyr':
##
##
       expand, pack, unpack
## Loaded glmnet 4.1-3
1. ISLR 5.4, Question 5
a)
library(ISLR2)
##
## Attaching package: 'ISLR2'
## The following object is masked from 'package:MASS':
##
##
       Boston
default <- Default
head(default)
     default student
##
                      balance
                                   income
## 1
          No
                  No 729.5265 44361.625
## 2
          No
                 Yes 817.1804 12106.135
## 3
                  No 1073.5492 31767.139
          No
                  No 529.2506 35704.494
## 4
          No
## 5
          No
                  No
                     785.6559 38463.496
## 6
                 Yes 919.5885 7491.559
          No
logit <- glm(default ~ income + balance, data = default, family = binomial(link="logit"))</pre>
summary(logit)
```

```
##
## Call:
## glm(formula = default ~ income + balance, family = binomial(link = "logit"),
       data = default)
##
## Deviance Residuals:
      Min
                      Median
                                           Max
## -2.4725 -0.1444 -0.0574 -0.0211
                                        3.7245
## Coefficients:
                 Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -1.154e+01 4.348e-01 -26.545 < 2e-16 ***
                2.081e-05 4.985e-06
                                      4.174 2.99e-05 ***
## income
## balance
                5.647e-03 2.274e-04 24.836 < 2e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 2920.6 on 9999 degrees of freedom
## Residual deviance: 1579.0 on 9997 degrees of freedom
## AIC: 1585
## Number of Fisher Scoring iterations: 8
b)
set.seed(1)
# split the dataset into training and testing sets
training_samples <- default$default %>%
  createDataPartition(p = 0.8, list = FALSE)
train_data <- default[training_samples, ]</pre>
test_data <- default[-training_samples, ]</pre>
# check it worked properly
dim(train_data); dim(test_data)
## [1] 8001
## [1] 1999
# Fit a logistic model
logit <- glm(default ~ income + balance, data = train_data, family = binomial(link="logit"))</pre>
# summary(logit)
```

```
# Make predictions for the full model
logit_pred <- predict(logit, newdata=test_data, "response")</pre>
logit_predicted_classes <- as.factor(ifelse(logit_pred > 0.5, "Yes", "No"))
# Let's make a table
logit_table <- table(test_data$default, logit_predicted_classes)</pre>
caret::confusionMatrix(logit_table)
## Confusion Matrix and Statistics
##
##
        logit_predicted_classes
##
           No Yes
##
     No 1930
                 3
##
     Yes
           51
                15
##
##
                  Accuracy: 0.973
##
                    95% CI: (0.9649, 0.9796)
##
       No Information Rate: 0.991
       P-Value [Acc > NIR] : 1
##
##
##
                     Kappa: 0.3479
##
   Mcnemar's Test P-Value: 1.596e-10
##
##
##
               Sensitivity: 0.9743
##
               Specificity: 0.8333
            Pos Pred Value: 0.9984
##
            Neg Pred Value: 0.2273
##
##
                Prevalence: 0.9910
##
            Detection Rate: 0.9655
      Detection Prevalence: 0.9670
##
##
         Balanced Accuracy: 0.9038
##
          'Positive' Class : No
##
##
c)
0.3 split:
set.seed(1)
# split the dataset into training and testing sets
training_samples <- default$default %>%
  createDataPartition(p = 0.3, list = FALSE)
train_data <- default[training_samples, ]</pre>
test_data <- default[-training_samples, ]</pre>
```

```
# Fit a logistic model
logit <- glm(default ~ income + balance, data = train_data, family = binomial(link="logit"))</pre>
# summary(logit)
# Make predictions for the full model
logit_pred <- predict(logit, newdata=test_data, "response")</pre>
logit_predicted_classes <- as.factor(ifelse(logit_pred > 0.5, "Yes", "No"))
# Let's make a table
logit_table <- table(test_data$default, logit_predicted_classes)</pre>
caret::confusionMatrix(logit_table)
## Confusion Matrix and Statistics
##
##
        logit_predicted_classes
##
           No Yes
     No 6737
##
                29
     Yes 151
##
                82
##
##
                  Accuracy : 0.9743
##
                    95% CI: (0.9703, 0.9779)
##
       No Information Rate: 0.9841
##
       P-Value [Acc > NIR] : 1
##
##
                     Kappa: 0.4653
##
   Mcnemar's Test P-Value : <2e-16
##
##
##
               Sensitivity: 0.9781
               Specificity: 0.7387
##
##
            Pos Pred Value: 0.9957
            Neg Pred Value: 0.3519
##
##
                Prevalence: 0.9841
##
            Detection Rate: 0.9626
##
      Detection Prevalence: 0.9667
##
         Balanced Accuracy: 0.8584
##
##
          'Positive' Class : No
##
0.5 split:
set.seed(1)
# split the dataset into training and testing sets
training_samples <- default$default %>%
```

```
createDataPartition(p = 0.5, list = FALSE)
train_data <- default[training_samples, ]</pre>
test_data <- default[-training_samples, ]</pre>
# Fit a logistic model
logit <- glm(default ~ income + balance, data = train_data, family = binomial(link="logit"))</pre>
# summary(logit)
# Make predictions for the full model
logit_pred <- predict(logit, newdata=test_data, "response")</pre>
logit_predicted_classes <- as.factor(ifelse(logit_pred > 0.5, "Yes", "No"))
# Let's make a table
logit_table <- table(test_data$default, logit_predicted_classes)</pre>
caret::confusionMatrix(logit_table)
## Confusion Matrix and Statistics
##
##
        logit_predicted_classes
##
           No Yes
     No 4817
                16
##
     Yes 126
##
                40
##
##
                  Accuracy : 0.9716
##
                    95% CI: (0.9666, 0.976)
       No Information Rate: 0.9888
##
##
       P-Value [Acc > NIR] : 1
##
##
                     Kappa: 0.3495
##
   Mcnemar's Test P-Value : <2e-16
##
##
               Sensitivity: 0.9745
##
##
               Specificity: 0.7143
##
            Pos Pred Value: 0.9967
##
            Neg Pred Value: 0.2410
                Prevalence: 0.9888
##
##
            Detection Rate: 0.9636
##
      Detection Prevalence: 0.9668
##
         Balanced Accuracy: 0.8444
##
##
          'Positive' Class : No
##
```

0.75 split

```
set.seed(1)
# split the dataset into training and testing sets
training_samples <- default$default %>%
  createDataPartition(p = 0.75, list = FALSE)
train_data <- default[training_samples, ]</pre>
test_data <- default[-training_samples, ]</pre>
# Fit a logistic model
logit <- glm(default ~ income + balance, data = train_data, family = binomial(link="logit"))</pre>
# summary(logit)
# Make predictions for the full model
logit_pred <- predict(logit, newdata=test_data, "response")</pre>
logit_predicted_classes <- as.factor(ifelse(logit_pred > 0.5, "Yes", "No"))
# Let's make a table
logit_table <- table(test_data$default, logit_predicted_classes)</pre>
caret::confusionMatrix(logit_table)
## Confusion Matrix and Statistics
##
##
        logit_predicted_classes
##
           No Yes
     No 2412
                 4
##
##
     Yes
           58
                25
##
##
                  Accuracy : 0.9752
##
                    95% CI: (0.9683, 0.9809)
       No Information Rate: 0.9884
##
       P-Value [Acc > NIR] : 1
##
##
##
                     Kappa : 0.4367
##
##
   Mcnemar's Test P-Value: 1.685e-11
##
               Sensitivity: 0.9765
##
               Specificity: 0.8621
##
##
            Pos Pred Value: 0.9983
##
            Neg Pred Value: 0.3012
                Prevalence: 0.9884
##
##
            Detection Rate: 0.9652
##
      Detection Prevalence: 0.9668
##
         Balanced Accuracy: 0.9193
##
##
          'Positive' Class : No
##
```

d)

##

```
set.seed(1)
# split the dataset into training and testing sets
training_samples <- default$default %>%
  createDataPartition(p = 0.75, list = FALSE)
train_data <- default[training_samples, ]</pre>
test_data <- default[-training_samples, ]</pre>
# Fit a logistic model
logit <- glm(default ~ income + balance + as.factor(student), data = train_data, family = binor
# summary(logit)
# Make predictions for the full model
logit_pred <- predict(logit, newdata=test_data, "response")</pre>
logit_predicted_classes <- as.factor(ifelse(logit_pred > 0.5, "Yes", "No"))
# Let's make a table
logit_table <- table(test_data$default, logit_predicted_classes)</pre>
caret::confusionMatrix(logit_table)
## Confusion Matrix and Statistics
##
##
        logit_predicted_classes
           No Yes
##
##
     No 2412
                 4
##
     Yes
           59
                24
##
##
                  Accuracy : 0.9748
                    95% CI: (0.9679, 0.9806)
##
##
       No Information Rate: 0.9888
       P-Value [Acc > NIR] : 1
##
##
##
                     Kappa : 0.4228
##
##
   Mcnemar's Test P-Value: 1.022e-11
##
               Sensitivity: 0.9761
##
##
               Specificity: 0.8571
##
            Pos Pred Value: 0.9983
##
            Neg Pred Value: 0.2892
##
                Prevalence: 0.9888
            Detection Rate: 0.9652
##
##
      Detection Prevalence: 0.9668
##
         Balanced Accuracy: 0.9166
```

```
## 'Positive' Class : No
##
```

Adding the dummy variable to the model did not lead to a reduction in test error rate.

Session Info

```
sessionInfo()
```

```
## R version 4.1.2 (2021-11-01)
## Platform: x86_64-apple-darwin17.0 (64-bit)
## Running under: macOS Big Sur 10.16
##
## Matrix products: default
           /Library/Frameworks/R.framework/Versions/4.1/Resources/lib/libRblas.0.dylib
## BLAS:
## LAPACK: /Library/Frameworks/R.framework/Versions/4.1/Resources/lib/libRlapack.dylib
##
## locale:
## [1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/c/en_US.UTF-8/en_US.UTF-8
## attached base packages:
## [1] stats
                 graphics grDevices utils
                                                datasets methods
                                                                    base
##
## other attached packages:
##
  [1] ISLR2_1.3-1
                        glmnet_4.1-3
                                         Matrix_1.4-0
                                                         caret_6.0-90
    [5] lattice_0.20-45 nnet_7.3-17
                                         MASS_7.3-55
                                                         forcats_0.5.1
   [9] stringr_1.4.0
                                                         readr_2.1.1
                        dplyr_1.0.7
                                         purrr_0.3.4
## [13] tidyr_1.1.4
                        tibble_3.1.6
                                         ggplot2_3.3.5
                                                         tidyverse_1.3.1
##
## loaded via a namespace (and not attached):
   [1] nlme_3.1-155
##
                             fs_1.5.2
                                                   lubridate_1.8.0
## [4] httr 1.4.2
                             tools 4.1.2
                                                   backports 1.4.1
## [7] utf8_1.2.2
                             R6_2.5.1
                                                   rpart_4.1.16
## [10] DBI_1.1.2
                             colorspace_2.0-2
                                                   withr_2.4.3
## [13] tidyselect_1.1.1
                             compiler_4.1.2
                                                   cli_3.1.1
## [16] rvest_1.0.2
                                                   xm12_1.3.3
                             formatR_1.11
## [19] scales_1.1.1
                             proxy_0.4-26
                                                   digest_0.6.29
## [22] rmarkdown_2.11
                             pkgconfig_2.0.3
                                                   htmltools_0.5.2
## [25] parallelly_1.30.0
                             dbplyr_2.1.1
                                                   fastmap_1.1.0
## [28] rlang_1.0.0
                             readxl_1.3.1
                                                   rstudioapi_0.13
## [31] shape_1.4.6
                             generics_0.1.1
                                                   jsonlite_1.7.3
## [34] ModelMetrics_1.2.2.2 magrittr_2.0.2
                                                   Rcpp_1.0.8
## [37] munsell_0.5.0
                             fansi_1.0.2
                                                   lifecycle_1.0.1
## [40] stringi_1.7.6
                             pROC_1.18.0
                                                   yaml_2.2.2
## [43] plyr_1.8.6
                             recipes_0.1.17
                                                   grid_4.1.2
```

##	[49]	parallel_4.1.2 haven_2.4.3 knitr_1.37	listenv_0.8.0 splines_4.1.2 pillar_1.6.5	<pre>crayon_1.4.2 hms_1.1.1 future.apply_1.8.1</pre>
		reshape2_1.4.4	codetools_0.2-18	stats4_4.1.2
##	[58]	reprex_2.0.1	glue_1.6.1	evaluate_0.14
##	[61]	data.table_1.14.2	modelr_0.1.8	vctrs_0.3.8
##	[64]	tzdb_0.2.0	foreach_1.5.1	cellranger_1.1.0
##	[67]	gtable_0.3.0	future_1.23.0	assertthat_0.2.1
##	[70]	xfun_0.29	gower_0.2.2	prodlim_2019.11.13
##	[73]	broom_0.7.12	e1071_1.7-9	class_7.3-20
##	[76]	survival_3.2-13	timeDate_3043.102	iterators_1.0.13
##	[79]	lava_1.6.10	globals_0.14.0	ellipsis_0.3.2
##	[82]	ipred_0.9-12	-	- · ·