Homework 2

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Question 2: A Hospital Audit

Hospital Audits are important to determine the effectiveness of hospital operations from a objective standpoint. In this particular case, the goal is to determining the performance of radiologists using a statistical audit of their recent patient interactions - a crucial link between modern data-science and hospital operations. Two overall questions are posited:

- 1. First question: are some radiologists more clinically conservative than others in recalling patients, holding patient risk factors equal?
- 2. Second question: when the radiologists at this hospital interpret a mammogram to make a decision on whether to recall the patient, does the data suggest that they should be weighing some clinical risk factors more heavily than they currently are?

At the core of each question is reducing the number of false negatives - where a radiologist recommends a patient to conduct further tests and thereby allows a patient to begin immediately; and false positives - where a radiologist recommends further tests but ultimately turns out that there was no cancer. By introducing a statistical model, the goal is to augment the predictive capabilities of radiologist and offer a better standard of care for patients.

This audit is structured in four parts: first is a brief summary of the data and how it is structured, second is a demonstration and presentation of answering question one, third is a similar approach for question two, fourth is a conclusion of the audit's findings and recommendations for improvement of future radiologist performance or audit effectiveness.

Part One: Brief Summary of Data

```
##
           radiologist
                              cancer
                                                 recall
                                                                      age
##
    radiologist13:198
                         Min.
                                 :0.00000
                                            Min.
                                                    :0.0000
                                                               age4049
                                                                        :287
##
    radiologist34:197
                         1st Qu.:0.00000
                                            1st Qu.:0.0000
                                                               age5059
                                                                        :284
    radiologist66:198
                         Median :0.00000
                                            Median :0.0000
##
                                                               age6069
                                                                        :199
##
    radiologist89:197
                         Mean
                                 :0.03749
                                            Mean
                                                    :0.1499
                                                               age70plus:217
##
    radiologist95:197
                         3rd Qu.:0.00000
                                            3rd Qu.:0.0000
##
                         Max.
                                 :1.00000
                                            Max.
                                                    :1.0000
##
                                                                     density
                         symptoms
                                                    menopause
       history
##
           :0.0000
                              :0.00000
                                                          :321
                                                                 density1: 89
                                         postmenoHT
                                                                 density2:332
##
    1st Qu.:0.0000
                      1st Qu.:0.00000
                                         postmenoNoHT
                                                          :360
    Median :0.0000
                      Median :0.00000
                                         postmenounknown: 35
                                                                 density3:460
                                                                 density4:106
##
    Mean
           :0.1763
                      Mean
                              :0.04863
                                         premeno
                                                          :271
                      3rd Qu.:0.00000
##
    3rd Qu.:0.0000
                              :1.00000
    Max.
           :1.0000
                      Max.
```

The data of mammograms used in this audit were selected from a Hospital in Seattle, Washington. At this hospital, five radiologists were selected at random for the audit - where about 200 mammograms were randomly selected from the hospital for each. For a total of 987 mammograms covering 7 parameters:

- age: 40-49*, 50-59, 60-69, 70 and older
- family history of breast cancer: 0=No*, 1=Yes
- history of breast biopsy/surgery: 0=No*, 1=Yes

- breast cancer symptoms: 0=No*, 1=Yes
- menopause/hormone-therapy status: Pre-menopausal, Post-menopausal & no hormone replacement therapy (HT), Post-menopausal & HT*, Post-menopausal & unknown HT
- previous mammogram: 0=No*, 1=Yes
- breast density classification: 1=Almost entirely fatty, 2=Scattered fibroglandular tissue*, 3=Heterogeneously dense, 4=Extremely dense

Of these factors, two are of special interest: [recall] and [cancer]. In the abstract [recall] can be explained as the following: upon seeing the medical history of a patient, they can either recommend either one of two actions: recall for further screening or not. It is presumed that radiologists utilize all of the information available before they make a decision. This implies that there is a inherent correlative factor between recall and patient history. On the other hand [cancer] is whether or not a patient, whether through the recall screening process, or through another pathway of discovery - develops cancer within a 12 month window after seeing the radiologist.

Part Two: Clinical Conservativism

Without knowing how patients are assigned to radiologists, it is presumed that the relationship is random at best, and preferential at worst. With a random assignment, we can presume that each radiologist chosen for the audit would have seen, on average, the same makeup of patients that would necessitate a mammogram. A random assignment would entail a random drawing of cancer patients from the overall total cancer patient pool from the population. If preferential - meaning that a patient approaches a radiologist and requests care and upon the approval of the radiologist, we see an issue of sampling error within the audit data; as there is a bias introduced between patient selection and radiologist. Radiologist may either self-select for more difficult cases or easier based on preference and patients self-select based on their estimate of the reputation of the radiologist within the medical community.

Regardless of assignment, the primary method of which we rank the clinical conservationism is to create a model that is trained on each of the radiologists' and then test the model on data from both the radiologist and other patients not seen by the radiologist in question. The goals behind this approach are twofold: one is to recreate a evaluation profile of the radiologist through a linear model of determining whether or not a patient should be recalled, two to determine whether or not a patient who is recalled or not develops cancer within a 12 month time frame.

The table below depicts the Root Mean Squared Error (RMSE) of each radiologist's model tested on a small sub-sample of the radiologist's test data and other radiologists' testing data.

```
##
                                                   lm2
                          lm1
                                    lm1.w
                                                            lm2.w
                  0.365743440 0.363424033
## radiologist13
                                           0.43912457 0.41722775
## radiologist34
                  0.285635286 0.388601868
                                           0.34351880 0.43011474
## radiologist66
                  0.392460315 0.369352289
                                           0.50307706 0.43786460
## radiologist89
                  0.408167944 0.400619339
                                           0.47049508 0.45182382
## radiologist95
                  0.349184207 0.382951933
                                           0.42744528 0.51249380
## SuperRad
                  0.361069858 0.354034509
                                           0.37239727 0.34937231
## Rad13.compare
                  0.004673582 0.009389524
                                           0.06672730 0.06785544
## Rad34.compare -0.075434573 0.034567359
                                          -0.02887846 0.08074243
## Rad66.compare
                  0.031390457 0.015317780
                                           0.13067979 0.08849229
## Rad89.compare
                  0.047098086 0.046584830
                                           0.09809781 0.10245151
## Rad95.compare -0.011885651 0.028917424
                                           0.05504802 0.16312149
```

Example:

• radiologist13: we have a the same linear model, lm1 = glm(recall ~ .-cancer, data=brca_train, maxit = maxit), trained to 20% of radiologist13's sample data as well as the whole mammogram data - excluding radiologist13's.

- SuperRad: is a model trained on a 20% random sample of the whole data set and tested on the remainder of the whole data set. This pseudo-radiologist serves as the benchmark for comparing radiologists to an artificial standard if one radiologist had access and saw all of the patients from the data set.
- Rad13.compare: is determined by subtracting the model RMSE result of *radiologist13 by Super-Rad. A positive value means that a model trained on radiologist13's training data did worse once it was tested on out of sample testing data and vice-versa.

```
##
                          lm1
                                    lm1.w
                                                   lm2
                                                            lm2.w
                  0.365743440 0.363424033
## radiologist13
                                            0.43912457 0.41722775
## radiologist34
                  0.285635286 0.388601868
                                            0.34351880 0.43011474
## radiologist66
                  0.392460315 0.369352289
                                            0.50307706 0.43786460
## radiologist89
                  0.408167944 0.400619339
                                            0.47049508 0.45182382
## radiologist95
                  0.349184207 0.382951933
                                            0.42744528 0.51249380
## SuperRad
                  0.361069858 0.354034509
                                            0.37239727 0.34937231
## Rad13.compare
                  0.004673582 0.009389524
                                            0.06672730 0.06785544
                                          -0.02887846 0.08074243
## Rad34.compare -0.075434573 0.034567359
## Rad66.compare
                  0.031390457 0.015317780
                                            0.13067979 0.08849229
## Rad89.compare
                  0.047098086 0.046584830
                                            0.09809781 0.10245151
## Rad95.compare -0.011885651 0.028917424
                                            0.05504802 0.16312149
```

Just by viewing the table, it can be clearly discerned under lm1 that on average, radiologists 13, 66, and 89 had worse performance than the benchmark SuperRad when looking at the RadXX.compare values for each radiologist; while 34 and 95 had better performance. But when we examine the results of each radiologists model tested on the global data set, we find that on average, all radiologists were worse off. However lm1 is a linear regression involving non-interacting variables from the data set. If we were to examine lm2 <<glm(recall ~ (.-cancer)^2,data=brca_train,maxit = maxit) where we interact every variable with itself and another we find different results. Radiologist 95's model performance flips and becomes worse with 95's within-sample data. But once tested on the global data set, all radiologists' models performed worse than the benchmark. The takeaway from this analysis demonstrates that human radiologists, on average, are not as effective in determining whether or not a patient should be recalled than a statistical model. Although this might increase the number of false positives and false negatives, the overall increase in cancer detection would allow immediate treatment for true positives who otherwise would have gone undiagnosed. As for whether or not this behavior can be determined to be clinically conservative, meaning that radiologist will opt to recall a patient even if the clinical factors do not signal a need to recall, the distinction is minimal at best and hard to determine as all of the radiologists selected in the audit perform marginally better or worse than the benchmark.

Part Three: Weighing Different Clinical Risk Factors

We first approach this question by developing four linear models that attempts to predict cancer rates based on the parameters available in the data set.

```
    lm3 <<- glm(cancer ~ recall,data=brca_train,maxit = maxit)</li>
    lm4 <<- glm(cancer ~ recall + history,data=brca_train,maxit = maxit)</li>
    lm5 <<- glm(cancer ~ .,data=brca_train,maxit = maxit)</li>
    lm6 <<- glm(cancer ~ (.)^2,data=brca_train,maxit = maxit)</li>
```

Because the goal of this question is to determine whether or not radiologists are effectively utilizing all of a patient's clinical data to determine whether or not to recall a patient, we first examine **lm3** and **lm4**. Both are linear models designed to find the partial effect of whether or not a patient was recalled and if they developed cancer within the next 12 months. However the distinction is that **lm3** only has recall as its x variable while **lm4** has both recall and family history.

```
summary(1m3)
```

```
##
## Call:
## glm(formula = cancer ~ recall, data = brca_train, maxit = maxit)
## Deviance Residuals:
##
       Min
                   1Q
                        Median
                                       3Q
                                                Max
## -0.15385 -0.01783 -0.01783 -0.01783
                                            0.98217
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 0.017831
                          0.007138
                                     2.498
                                             0.0127 *
              0.136016
                          0.018547
                                     7.334 5.57e-13 ***
## recall
##
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.03428528)
##
##
       Null deviance: 28.861
                             on 789
                                      degrees of freedom
## Residual deviance: 27.017 on 788 degrees of freedom
## AIC: -418.78
##
## Number of Fisher Scoring iterations: 2
summary(lm5)
##
## Call:
## glm(formula = cancer ~ . - recall, data = brca_train, maxit = maxit)
##
## Deviance Residuals:
##
        Min
                   1Q
                        Median
                                       3Q
                                                Max
## -0.13967 -0.05402 -0.03076 -0.01538
                                            0.99279
##
## Coefficients:
##
                             Estimate Std. Error t value Pr(>|t|)
                                        0.037904 -0.347 0.72859
## (Intercept)
                            -0.013158
                                        0.021682 -0.242
## radiologistradiologist34 -0.005238
                                                          0.80918
                                                 -0.429
## radiologistradiologist66 -0.009295
                                        0.021687
                                                          0.66835
## radiologistradiologist89 -0.009161
                                       0.022178 -0.413
                                                          0.67968
## radiologistradiologist95 -0.009552
                                       0.021770 - 0.439
                                                          0.66095
## ageage5059
                             0.032176
                                       0.022905
                                                   1.405
                                                          0.16049
## ageage6069
                             0.025219
                                       0.027285
                                                   0.924
                                                          0.35563
## ageage70plus
                             0.068629
                                       0.027004
                                                   2.541
                                                          0.01123 *
## history
                             0.014234
                                       0.018145
                                                   0.784
                                                          0.43302
## symptoms
                             0.003980
                                       0.030258
                                                   0.132
                                                          0.89537
                                       0.016983 -0.860
## menopausepostmenoNoHT
                           -0.014608
                                                          0.38995
## menopausepostmenounknown -0.006012
                                        0.039544 -0.152
                                                          0.87920
                                                   0.682
## menopausepremeno
                             0.016505
                                        0.024185
                                                          0.49516
## densitydensity2
                             0.013153
                                        0.026395
                                                   0.498
                                                          0.61840
                             0.026385
                                        0.026551
                                                   0.994
                                                          0.32065
## densitydensity3
                             0.089434
                                        0.033242
                                                   2.690 0.00729 **
## densitydensity4
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 0.03639461)
```

```
##
## Null deviance: 28.861 on 789 degrees of freedom
## Residual deviance: 28.169 on 774 degrees of freedom
## AIC: -357.78
##
## Number of Fisher Scoring iterations: 2
```

By itself, we can see that the **recall** variable has a very significant (p-value close to 0) and large effect on whether or not a patient develops cancer. This makes sense because upon evaluating a patient, a radiologist will then determine whether or not the patient will be recalled and receive additional testing. Based on their experience and education, they will want to find the factors that most likely contributes to cancer. At the same time however, we also see significant (in terms of p-value and magnitude) effects from **age**, **menopause/hormone-therapy status**, and **breast density classification**. In light of these factors, a series of model efficacy tests were conducted to determine the effectiveness of different models.

```
##
                                      lm3.w
                           1m3
                                                       1m4
                                                                  lm4.w
## radiologist13
                  0.326750727
                                0.337005247
                                              0.3332733946
                                                            0.336809528
  radiologist34
                  0.213122379
                                0.311894469
                                              0.2255037862
                                                            0.319724743
## radiologist66
                  0.377001486
                                0.335069096
                                              0.3791503679
                                                            0.337329619
## radiologist89
                  0.376251046
                                0.320967727
                                              0.3945406249
                                                            0.360075362
  radiologist95
                  0.296198425
                                0.319017089
                                              0.3011243104
                                                            0.331615728
## SuperRad
                  0.332031569
                                0.329502308
                                              0.3330038087
                                                            0.330490614
## Rad13.compare -0.005280841
                                0.007502939
                                              0.0002695859
                                                            0.006318914
## Rad34.compare -0.118909190
                               -0.017607838
                                            -0.1075000225
                                                           -0.010765870
## Rad66.compare
                  0.044969917
                                0.005566788
                                              0.0461465593
                                                            0.006839005
## Rad89.compare
                  0.044219477 -0.008534581
                                              0.0615368162
                                                            0.029584748
  Rad95.compare -0.035833144
                               -0.010485219
                                            -0.0318794982
                                                            0.001125114
##
                           lm5
                                      lm5.w
                                                      lm6
                                                               lm6.w
## radiologist13
                  0.373740388
                                0.378109275
                                              0.406027074 0.40226902
## radiologist34
                  0.279916211
                                0.397728621
                                              0.311721104 0.41455655
## radiologist66
                  0.406538005
                                0.362819638
                                              0.416143141 0.38860249
## radiologist89
                  0.432300265
                                0.396129977
                                              0.474467845 0.45781915
  radiologist95
                  0.349830246
                                0.387427297
                                              0.386435267 0.43368063
## SuperRad
                  0.377678967
                                0.374250986
                                              0.380752134 0.37497029
## Rad13.compare -0.003938579
                                0.003858289
                                              0.025274940 0.02729873
## Rad34.compare -0.097762756
                                0.023477635
                                             -0.069031030 0.03958626
                  0.028859038
## Rad66.compare
                               -0.011431347
                                              0.035391007 0.01363220
## Rad89.compare
                  0.054621298
                                0.021878992
                                              0.093715711 0.08284886
## Rad95.compare -0.027848721
                                0.013176311
                                             0.005683133 0.05871034
```

Looking across **SuperRad** we see that the RMSE of each model remains fairly consistent throughout the different implementation and test of each model - except when we exclude **recall** in models **lm5** and **lm6**. The exclusion of **recall** has a meaningful impact models' ability to guess the cancer rate for each patient. Given this puzzling outcome, the next step would be to examine **lm5.w** and **lm6.w** where we take models that exclude **recall** - after all, as recall determinations occur after a radiologist sees a patient and not before, we cannot use it to predict cancer; and see which radiologist model performs the best. Iteration terms seems to be resulting in higher RMSE in the predictive model than by itself. Given the summary results from earlier regarding the significance of some variables over others, it can be concluded that radiologists weigh **age**, **menopause/hormone-therapy status**, and **breast density classification** as indicators of cancer than other factors excluding recall.

Part Four: Conclusion

Ultimately, it can be determined that human radiologists may appear to be more conservative than a statistical model, but the underlying analysis claims otherwise - the difference is small in nature and not of sufficient

significance to sacrifice patient care for a more effective diagnosing mechanism. The number of false positives and false negatives remain small in comparison when the model changes from one to another.

```
## [1] "lm3 Confusion Table"
##
      yhat
## y
              1
         0
     0 163
             27
         3
##
     1
              4
## [1] "lm4 Confusion Table"
      yhat
## y
         0
              1
##
     0 162
             28
     1
         3
## [1] "lm5 Confusion Table"
##
      yhat
## y
              1
##
     0 184
              6
         7
##
              0
## [1] "lm6 Confusion Table"
##
      yhat
## y
         0
              1
##
             26
     0 164
         5
```

- Pair-wise guesses and actual cancer results.
- (0,0) means that a patient did not have caner and was not recalled.
- (1,0) means that a patient had cancer but was not recalled.
- (0,1) means that a patient did not have cancer but was recalled.
- (1,1) means that a patient had cancer and was successfully recalled.