Homework 1

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Number 1

probability: 400 / 50,001 = 0.0080 .8% probability of a MI in high risk men not on statins odds: 400 / 49601 = 0.0081

Number 2

probability: 240 / 50,002 = 0.0048 .48% probability of a MI in high risk men on statins odds: 240 / 49762 = 0.0048

Number 3

probability: 0.008 - 0.0048 = 0.0032 odds: 0.0081 - 0.0048 = 0.0033

Number 4

There is a higher chance of MI in high risk men not on statins than on statins, so it seems that statins do help.

Number 5

low risk, not on statins: probability: 360 / 180,002 = 0.002 odds: 360 / 179642 = 0.002 low risk, on statins: probability: 30 / 20,001 = 0.0015 odds: 30 / 19971 = 0.0015

Number 6

The difference for low risk men is only 0.0005, so it seems to help but only a small amount.

Number 7

probability: (400 + 360) / (50,001 + 180,002) = 0.0033.33% probability of MI in men not on statins odds: (400 + 360) / (49601 + 179642) = 0.0033

Number 8

probability: (240 + 30) / (50,002 + 20,001) = 0.0039 .39% probability of MI in men on statins odds: (240 + 30) / (19971 + 49762) = 0.0039

Number 9

If we did not know about the risk status, we would probably conclude that status have no effect on the chance of MI, or maybe that they have a negative effect because the difference here is -0.0006.

Number 10

It seems that statins have an effect only on high risk men, but if we don't know the risk status then statins do not appear to have an effect on the chance of MI. There is a casual effect when we separate by risk status because low risk men are less likely to be taking statins and less likely to have MI just in general, so it doesn't make sense to lump them together with the high risk men.

Number 11

We could expect a similar phenomenon with allergy medication, as we discussed in class. If people don't already have allergies, they will likely not need to take allergy medication and they will be less likely to experience allergy symptoms regardless of whether they take medication or not.

Number 12

(see end of document)

Number 13

The coefficient for the Statins variable is -0.4812, which is significant at alpha < 0.001. This means that patients taking statins are less likely to have an MI, and patients at high risk are more likely to have MI.

```
##
## Call:
  glm(formula = MI ~ factor(RiskScore) + Statins, family = binomial(link = "logit"),
##
       data = MI, weights = N)
##
## Deviance Residuals:
##
                           3
                                              5
                                                                7
                                                                          8
         1
                  2
                                     4
                                                       6
##
   62.230
           -28.053
                      50.528
                              -22.106
                                         66.819
                                                 -27.002
                                                           20.022
                                                                     -7.079
##
## Coefficients:
##
                      Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                      -6.19902
                                  0.05092 -121.737
                      1.36624
                                             19.869
                                                    < 2e-16 ***
## factor(RiskScore)1
                                  0.06876
## Statins
                      -0.48120
                                  0.07587
                                             -6.342 2.26e-10 ***
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
##
  (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 13745
                                   degrees of freedom
                             on 7
## Residual deviance: 13346
                             on 5
                                   degrees of freedom
##
  AIC: 13352
##
## Number of Fisher Scoring iterations: 8
          (Intercept) factor(RiskScore)1
##
                                                     Statins
##
           -6.1990156
                               1.3662382
                                                  -0.4811988
```

Number 14

The coefficient for Statins is now -0.288, which is not significant (alpha > 1). The interaction term is also not significant (alpha > 1).

```
fit2 <- glm(MI ~ RiskScore*Statins, weights = N, data = MI, family = binomial(link = "logit"))
summary(fit2)
##
## Call:
## glm(formula = MI ~ RiskScore * Statins, family = binomial(link = "logit"),
       data = MI, weights = N)
##
## Deviance Residuals:
##
         1
                           3
                                    4
                                             5
                                                      6
                                                               7
                                                                        8
    62.150 -28.227
                      50.624 -21.883
                                        66.892 -26.819
                                                          19.752
##
## Coefficients:
##
                     Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                     -6.21262
                                 0.05276 -117.759
                                                    <2e-16 ***
## RiskScore
                      1.39232
                                 0.07282
                                           19.119
                                                    <2e-16 ***
## Statins
                     -0.28822
                                 0.19017
                                           -1.516
                                                     0.130
                                 0.20706
## RiskScore:Statins -0.22584
                                           -1.091
                                                     0.275
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 13745 on 7 degrees of freedom
##
## Residual deviance: 13345 on 4 degrees of freedom
## AIC: 13353
##
## Number of Fisher Scoring iterations: 8
```

Number 15

Given the models above, I would say that statins do have an effect, but the risk score is a more prominent predictor of whether a patient will have MI or not.