MLR With Qualitative Predictors

An input variable with c classes will be respresented via c-1 binary input variables (this is effectively one-hot encoding). When all of said inputs are 0, this represents the *baseline* case, which we can choose arbitrarily. Then, each class is represented by making one of the variables 1 and all the others 0 (one is "hot").

With this scheme, we effectively end up with c regression models. The pseudo-variables going to 0 causes certain terms to drop out and the pseudo-variable going to 1 causes a term to be added to the intercept.

If we have 2 inputs, X1, X2 where X2 is a pseudo-variable plot indicating one of two classes, we can plot Y against X1 and leave the categorical variables off of the x-axis and instead just distinguishing between classes via a color, we can observe how a change in class affects the linear relationship.

```
df <- read.csv("../datasets/insurance.csv")
model_sum <- lm(month ~ size + factor(type), df)
summary(model_sum)</pre>
```

```
##
## Call:
## lm(formula = month ~ size + factor(type), data = df)
##
## Residuals:
##
      Min
                1Q Median
                                3Q
                                       Max
##
   -5.6915 -1.7036 -0.4385
                           1.9210
                                   6.3406
##
## Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                 33.874069
                             1.813858
                                     18.675 9.15e-13 ***
## size
                -0.101742
                             0.008891 -11.443 2.07e-09 ***
## factor(type)1 8.055469
                            1.459106
                                       5.521 3.74e-05 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 3.221 on 17 degrees of freedom
## Multiple R-squared: 0.8951, Adjusted R-squared: 0.8827
## F-statistic: 72.5 on 2 and 17 DF, p-value: 4.765e-09
```

This model has an absence of any interaction effect between X_1 and X_2 because we have not modeled it in. The change in mean adoption time between the two classes is called the main effect (β_2) . We can add in an X_1X_2 to the model and observe its change:

```
df <- read.csv("../datasets/insurance.csv")
model_prod <- lm(month ~ size + factor(type) + size * factor(type), df)
summary(model_prod)</pre>
```

```
##
## Call:
## lm(formula = month ~ size + factor(type) + size * factor(type),
##
       data = df
##
## Residuals:
##
       Min
                1Q Median
                                 3Q
                                        Max
  -5.7144 -1.7064 -0.4557
                           1.9311
                                    6.3259
##
## Coefficients:
##
                        Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                      33.8383695 2.4406498 13.864 2.47e-10 ***
```

```
## size
                     -0.1015306 0.0130525
                                           -7.779 7.97e-07 ***
                                             2.225
## factor(type)1
                      8.1312501
                                3.6540517
                                                     0.0408 *
## size:factor(type)1 -0.0004171
                                0.0183312
                                            -0.023
                                                     0.9821
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.32 on 16 degrees of freedom
## Multiple R-squared: 0.8951, Adjusted R-squared: 0.8754
## F-statistic: 45.49 on 3 and 16 DF, p-value: 4.675e-08
```

The model is similar and the estimate of the interaction effect is very close to zero. We can test the hypothesis that it is zero using the results of the table, dividing the estimate by its stderr to get t^* and comparing to $t(1-\alpha/2, n-p)$. Equivalently, we can also perform a GLT where the reduced model is the original and the full is the new one:

```
anova(model_sum, model_prod)
```

```
## Analysis of Variance Table
##
## Model 1: month ~ size + factor(type)
## Model 2: month ~ size + factor(type) + size * factor(type)
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 17 176.39
## 2 16 176.38 1 0.0057084 5e-04 0.9821
```

So the term is insignificant.

Here, $\beta_0 + \beta_1 X_1$ describes the linear model on the baseline category. β_1 describes the linear impact of X_1 on Y, β_2 describes the main effect of the difference in categories (associated with X_2 , not X_1), and β_3 describes the interaction effect between X_1 and X_2 , which is associated with X_1 .

So the model for the other category is $(\beta_0 + \beta_2) + (\beta_1 + \beta_3)X_1$.

We might also call β_3 β_{12} and will do this when there are multiple interaction terms to keep track of.

Main effects describe a difference in *intercept* between categories, while interaction effects describe a difference in *linear impact* between categories.

Note that we do not have interaction effects between pseudo-variables.

To test various hypotheses, we need to translate them into equations of the parameters. See L17.11-14 for examples. We can then translate these into reduced and full models. If we are *given* information, then this is incorporated into the full model (see L17.14).

L20

MLR Diagnostics

See Ch10 p.384

Added-Variable Plots

See p.384 / L20

Considers the marginal effect of a predictor variable being added in after all others are considered. Achieved by plotting the residuals of Y others against $X_i|others$.

Hat Matrix Review

L20.10 (hat formula, residuals, variance)

Hat matrix is effectively a transformation that when applied to Y yields \hat{Y}

Studentized Residuals

L20.11

These are just residuals divided by stderr; measures the number of stderrs of the deviations. Detects Y outliers.

Studentized Deleted Residual

L20.12

Also Y-outliers. Can also be caused by non-normality or heteroskedasticity of errors.

Bonferroni Procedure L20.13

Hat Matrix & Leverage

Leverage of the *i*th X value is the *i*th diagonal of the hat matrix, h_{ii} . The larger it is, the smaller the variance of the residual.

Extreme values of h can indicate X outliers.

See comments on L20.16

DFFITS

p.400 / L20.18

Identifies influence on a single fitted value.

Cook's Distance

p.402 / L20.19

Influence on all fitted values.

DFBETAS

p.404 / L20.20

Influence of the *i*th case on each regression coefficient.

influencePlot()

Shows DFFITS / Cook's in size of circle, vertical lines drawn at 2x and 3x average hat value, horizontal lines at -2, 0, 2 on the studentized residual scale.

Multicollinearity

VIF

p.408 / L20.28

Variance inflation factor measures how much the variance of the estimated value of the regression coefficient of a certain input variable is increased. Also related to \mathbb{R}^2 and tolerance.

Advanced Remedial Measures

WLS

p.421 / L21

Addresses heteroskedasticity. Downside is that MSE has no clear interpretation in the context of the problem so it cannot be used to compare models.

t- and F- tests assume normal error with constant variance, but if we're using WLS, that might be an issue, so we might want to look into Bootstrapping.

Ridge Regression

p.431

Addresses multicollinearity issues. Shrinks estimators by adding a size penalty so that they become biased but closer on average to the true value (useful when there is a very wide spread on them).

Same note on bootstrapping as above applies here.

Robust Regression

p.437

Addresses influential cases. Similar to WLS in that b oth use a weight function to adjust influence of observation and both can handle unequal variance. If primary issue is heteroskedasticity, use WLS, but robust also uses weight function to trim influence of outliers/influential cases. They can be used together in some cases.

Bootstrapping

p.458

Allows approximate estimation of CI in WLS, Robust, Ridge regression and correct intervals when errors are strongly non-normal.

One-Way ANOVA Factor Effects Model

HW7 / L22 / p.733

Two-Way

HW7 / L23 / p.812

See especially 849 for intervals on factor effects when factors do not interact, 856 when they do.

843 for F-tests.

841 for ANOVA table.