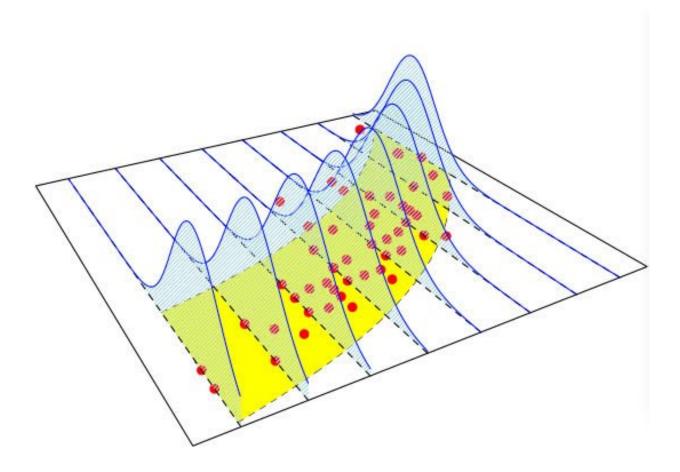
# Week 3 - AYU - Pod

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GLM in R

```
library(tidyverse)
d <- read_csv("data/TermLife.csv")
d <- d[d$FACE>0, ]
modelMLR <- glm(FACE ~ EDUCATION+NUMHH+INCOME, data=d)
summary(modelMLR)
##</pre>
```

```
## Call:
## glm(formula = FACE ~ EDUCATION + NUMHH + INCOME, data = d)
##
## Deviance Residuals:
##
                        Median
                                       3Q
       Min
                   1Q
                                                Max
            -651472
                       -339712
                                  -31468
                                           13039540
##
  -2655152
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.773e+06 6.107e+05 -2.904 0.003987 **
## EDUCATION
               1.463e+05 3.849e+04
                                       3.801 0.000178 ***
## NUMHH
               1.098e+05 6.552e+04
                                       1.675 0.095001 .
## INCOME
               3.392e-01 1.201e-01
                                       2.825 0.005077 **
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 2.547961e+12)
##
      Null deviance: 7.6816e+14 on 274 degrees of freedom
##
## Residual deviance: 6.9050e+14 on 271 degrees of freedom
## AIC: 8642.1
## Number of Fisher Scoring iterations: 2
```

#### Logistic Regression

We will use the [Wisconsin Hospital Data] again for this example. In the data, our response variable is the total charge, which is a numeric variable, so we cannot use logistic regression for this variable. We will create a binary variable from the total charge. Instead of the exact charge, we are interested in the charge is small (less than the median) or large (more than the median). Create a variable TOTCHG2 that takes the below value

- small if TOTCHG is smaller than the average of TOTCHG
- large otherwise

```
library(tidyverse)
d <- read_csv('data/frees/HospitalCosts.csv')
d$TOTCHG2 = ifelse(d$TOTCHG > median(d$TOTCHG), 1, 0)
```

Now that TOTCHG2 is binary, we can regress it using the logistic regression.

```
model <- glm(TOTCHG2 ~ AGE + factor(GENDER) + LOS + factor(RACE) + APRDRG, data=d, family = binomial(
summary(model)</pre>
```

```
## Deviance Residuals:
                    Median
      Min
                10
                                  30
                                          Max
                    0.0000
## -2.9326 -0.5560
                              0.5374
                                       3.5948
##
## Coefficients:
                    Estimate Std. Error z value Pr(>|z|)
                   3.067e-01 7.824e-01
## (Intercept)
                                        0.392 0.69508
## AGE
                   1.608e-01 2.626e-02
                                          6.123 9.18e-10 ***
## factor(GENDER)1 -8.395e-01 2.822e-01 -2.975 0.00293 **
## LOS
                   2.636e+00 2.603e-01 10.127 < 2e-16 ***
## factor(RACE)2
                  -1.253e+00 1.466e+00
                                        -0.855 0.39266
## factor(RACE)3
                 1.194e+01 1.455e+03
                                          0.008 0.99345
## factor(RACE)4 -4.095e-01 1.718e+00
                                        -0.238 0.81156
## factor(RACE)5
                 -4.230e+00 1.501e+00 -2.818 0.00484 **
                 -1.494e+01 1.020e+03 -0.015 0.98831
## factor(RACE)6
## APRDRG
                  -1.020e-02 1.421e-03 -7.181 6.90e-13 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 691.76 on 498 degrees of freedom
## Residual deviance: 355.38 on 489 degrees of freedom
     (1 observation deleted due to missingness)
## AIC: 375.38
## Number of Fisher Scoring iterations: 14
Prediction
Find the probability that a person get charged a large amount.
predict(model, list(AGE = 15, GENDER = 1, LOS = 1, RACE = 1, APRDRG = 600), type = 'response')
          1
## 0.1672514
# Checking the accuracy of the model
predicted_value = ifelse(predict(model, d , type = 'response')>=.5, 1, 0)
true_value = d$TOTCHG2
library(caret)
confusion_matrix = confusionMatrix(data=factor(predicted_value), reference = factor(true_value))
```

## glm(formula = TOTCHG2 ~ AGE + factor(GENDER) + LOS + factor(RACE) +
## APRDRG, family = binomial(link = "logit"), data = d)

## ## Call:

confusion\_matrix

##

## Confusion Matrix and Statistics

```
##
             Reference
              0 1
## Prediction
##
            0 236 37
            1 13 213
##
##
##
                  Accuracy: 0.8998
##
                    95% CI: (0.87, 0.9247)
##
       No Information Rate: 0.501
##
       P-Value [Acc > NIR] : < 2.2e-16
##
##
                     Kappa: 0.7996
##
   Mcnemar's Test P-Value: 0.001143
##
##
##
               Sensitivity: 0.9478
##
               Specificity: 0.8520
##
            Pos Pred Value: 0.8645
##
            Neg Pred Value: 0.9425
                Prevalence: 0.4990
##
##
            Detection Rate: 0.4729
##
     Detection Prevalence: 0.5471
##
         Balanced Accuracy: 0.8999
##
##
          'Positive' Class: 0
##
```

#### Poisson Regression

```
p = read_csv('data/poisson_sim.csv')
p$prog <- factor(p$prog, levels=1:3, labels=c("General", "Academic",
                                                      "Vocational"))
summary(m1 <- glm(num_awards ~ prog + math, family="poisson", data=p))</pre>
##
## Call:
## glm(formula = num_awards ~ prog + math, family = "poisson", data = p)
## Deviance Residuals:
##
       Min
                 1Q
                     Median
                                   3Q
                                           Max
## -2.2043 -0.8436 -0.5106
                               0.2558
                                        2.6796
##
## Coefficients:
                  Estimate Std. Error z value Pr(>|z|)
##
                  -5.24712
                              0.65845 -7.969 1.60e-15 ***
## (Intercept)
## progAcademic
                   1.08386
                              0.35825
                                        3.025 0.00248 **
## progVocational 0.36981
                              0.44107
                                        0.838 0.40179
## math
                   0.07015
                              0.01060
                                        6.619 3.63e-11 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
```

```
Null deviance: 287.67 on 199 degrees of freedom
## Residual deviance: 189.45 on 196 degrees of freedom
## AIC: 373.5
##
## Number of Fisher Scoring iterations: 6
d <- read_csv('data/poisson_sim.csv')</pre>
model = glm(num_awards ~ factor(prog) + math, data = d, family = 'poisson')
summary(model)
##
## Call:
## glm(formula = num_awards ~ factor(prog) + math, family = "poisson",
##
      data = d)
##
## Deviance Residuals:
      Min
                1Q
                    Median
                                  3Q
                                          Max
## -2.2043 -0.8436 -0.5106 0.2558
                                       2.6796
## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -5.24712 0.65845 -7.969 1.60e-15 ***
                                      3.025 0.00248 **
## factor(prog)2 1.08386
                            0.35825
                            0.44107
## factor(prog)3 0.36981
                                      0.838 0.40179
                            0.01060
## math
                 0.07015
                                      6.619 3.63e-11 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for poisson family taken to be 1)
##
      Null deviance: 287.67 on 199 degrees of freedom
## Residual deviance: 189.45 on 196 degrees of freedom
## AIC: 373.5
## Number of Fisher Scoring iterations: 6
# Coefficients
exp(coef(model))
##
     (Intercept) factor(prog)2 factor(prog)3
                                                     math
     0.00526263
                   2.95606545
                                 1.44745846
                                               1.07267164
# Goodness-of-fit test
gof.pvalue = 1 - pchisq(model$deviance, model$df.residual)
gof.pvalue
```

## [1] 0.6182274

### Questions

- 1. With your group find datasets that suitable for logistic regression to
- Specify the response variable and the input variables to build logistic regression.
- Compute the confusion matrix and report the accuracy of the model
- 2. With your group find datasets that suitable for poisson regression to
- Specify the response variable and the input variables to build poisson regression.
- Evaluate the quality of the poisson model.