566 Project Model Fitting, Selection, and Diagnostics

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0. Preliminaries

```
# setwd("C:/Users/Kaelan/Desktop/stat-566-causal-inference/code")
# access to stepAIC() function
library(MASS)
# make dummy variables
library(fastDummies)
# access variance inflation factor function
library(alr4)
## Loading required package: car
## Loading required package: carData
## Loading required package: effects
## Registered S3 methods overwritten by 'lme4':
    method
    cooks.distance.influence.merMod car
    influence.merMod
    dfbeta.influence.merMod
    dfbetas.influence.merMod
                                     car
## lattice theme set by effectsTheme()
## See ?effectsTheme for details.
# make correlation plot
library(corrplot)
## corrplot 0.88 loaded
source('566_exploratoryDataAnalysis.R')
## Attaching package: 'plyr'
## The following object is masked from 'package:ggpubr':
##
       mutate
##
# convert variables to factors
for (i in 1:k) {
 dat[, i] <- as.factor(dat[, i])</pre>
```

1. Fitting our Model (Logistic Regression)

```
# full model
glm.full <- glm(h1n1_vaccine ~ ., family = binomial(link = 'logit'), data = dat)</pre>
```

2. Variable Selection using Bayesian Information Criterion (BIC)

```
best.BIC <- stepAIC(glm.full, direction = "both", trace = FALSE, k = log(n))
best.BIC.variables <- attr(best.BIC$terms, "term.labels")
dat.best <- dat[c("h1n1_vaccine", best.BIC.variables)]</pre>
```

3. Model Diagnostics - Analyzing Multicollinearity Issues

correlation plots
corrplot(cor(dat.best.dummy))

