SPARSITY BLUES BRIAN A. FANNIN MARCH 21, 2018

OVERVIEW

- Where did this talk come from?
- Categorical vs continuous data
- Naive Bayes
- Decision trees
- Multiple Correspondance Analysis
- Let's model!



ORIGINS

- I gave a <u>talk</u> last year about APIs.
- As an afterthought, I tried to fit a model.
- The fits were challenging because the data was largely categorical.

THE DATA



BEFORE ANYONE GETS CARRIED AWAY...

From nflarrests.com:

Keep in mind there are 1700 NFL Players and their arrest rates are lower than the USA arrest rate.

Also: arrest != conviction

WHAT I TRIED TO MEASURE

I tried to measure whether a player would get a second arrest.

- Rate of 1st arrest requires player statistics for each season, which means a second source.
- I'm lazy. Let's check rate of second arrest.

JUST THE BASIC FACTS

- Number of players who've been arrested: 651
- Number of players w/more than one arrest: 146
- Probability of second arrest: 22.4%

So there is a small probability of having more than one arrest. Compare this to Bailey/Simon probability of second accident.

CATEGORICAL VS CONTINUOUS DATA

THERE ARE ONLY 2 KINDS OF DATA

- Continuous
- Categorical
 - Ordinal
 - Unordered
- OK, three would be a mixed distribution (zero-inflated, etc.)

Outcomes (for supervised learning) are either categorical or continuous (classification or regression).

CATEGORICAL DATA

- Gender
- Smoking
- Safe driver program
- Drug testing policy
- ...

Basically anything to which you could apply a schedule mod. And also:

- Class code
- Territory
- Zip code

And those are just the ones that might be in a rating manual.

CONTINUOUS OUTCOME

```
sims <- 1e3
tbl_linear <- tibble(
    x = runif(sims, 0, 10)
    , e = rnorm(sims, sd = 5)
) %>%
    mutate(
        y = 1.5 + 2 * x + e
    )
```



CATEGORICAL OUTCOME

- Logistic regression
- Support vector machine
- Tree methods

CATEGORICAL OUTCOME



CATEGORICAL PREDICTORS IN A LINEAR MODEL

```
set.seed(1234)
tbl_one_cat <- function(cat_label = 'a', sims = 1e3) {
    slope <- rnorm(1, 2, 2)
    intercept <- rnorm(1, 0, 10)
    tibble(
        x = runif(sims, 0, 10)
    , e = rnorm(sims, sd = 5)
    , category = rep(cat_label, sims)
) %>%
    mutate(
        y = intercept + slope * x + e
    )
}
tbl_cat <- map_dfr(letters[1:5], tbl_one_cat)</pre>
```

DIFFERENT INTERCEPTS



DIFFERENT SLOPES



OR BOTH



ISSUES

- Grouped data is looped data
- Handle this with credibility/hierarchical models
- What if we *only* have categorical predictors?

THE DESIGN MATRIX

categoryb:x	categorya:x	categorye	categoryd	categoryc	categoryb	categorya
С	8.6091538	0	0	0	0	1
С	6.4031061	0	0	0	0	1
C	0.0949576	0	0	0	0	1
C	2.3255051	0	0	0	0	1
C	6.6608376	0	0	0	0	1
C	5.1425114	0	0	0	0	1
C	6.9359129	0	0	0	0	1
C	5.4497484	0	0	0	0	1
C	2.8273358	0	0	0	0	1
C	9.2343348	0	0	0	0	1

Let's try some non-linear methods

NAIVE BAYES

BAYES

$$Pr(Y=y|X=x)=rac{Pr(Y=y)*Pr(X=x|Y=y)}{Pr(X=x)}$$

FIT

```
library(naivebayes)
```

```
fit_nb <- naive_bayes(
    formula = MultiArrest ~ PositionType
, data = tbl_players
)</pre>
```

```
## ========== Naive Bayes ===========
## Call:
## naive_bayes.formula(formula = MultiArrest ~ PositionType, data = tbl_players)
##
## A priori probabilities:
##
      FALSE
##
                 TRUE
## 0.7757296 0.2242704
##
## Tables:
##
## PositionType
                     FALSE
                                 TRUE
             D 0.534653465 0.541095890
##
             0 0.457425743 0.424657534
##
##
             S 0.007920792 0.034246575
```

CAN WE WORK THAT OUT MANUALLY?

```
prior_y <- sum(tbl_players$MultiArrest) / nrow(tbl_players)
prob_x <- sum(tbl_players$PositionType == 'D') / nrow(tbl_players)
tbl_cond <- tbl_players %>% filter(MultiArrest)
prob_x_cond <- sum(tbl_cond$PositionType == 'D') / nrow(tbl_cond)
prior_y * prob_x_cond / prob_x
## [1] 0.226361
predict(fit_nb, type = 'prob')[1, 'TRUE']
## TRUE
## 0.226361
prior_y
## [1] 0.2242704</pre>
```

TWO CATEGORIES

One:

$$Pr(Y = y|X = x) = \frac{Pr(Y = y) * Pr(X = x|Y = y)}{Pr(X = x)}$$

Two:

$$Pr(Y = y | X = x, Z = z)$$

$$= \frac{Pr(Y = y) * Pr(X = x | Y = y) * Pr(Z = z | Y = y)}{Pr(X = x) * Pr(Z = z)}$$

HOW ABOUT A LOT OF CATEGORIES?

HOW DO OUR PLAYERS LOOK?



NAIVE BAYES

- Often used in text processing
- Great for a sparse matrix
- It is 'naive' because we assume independence between categories

A DECISION TREE

CHARACTERISTICS OF A DECISION TREE

- Divides a sample into regions/subsets
- The 'prediction' is a function (usually the mean) of some value within each category
- Membership is assessed by computing some measure of fit. If a split improves the criteria, then it is made.
- Forward only, 'greedy'
- Number of levels and other criteria control the size and shape of the tree

MEASURES OF FIT

For regression:

• Construct regions which minimize residual sum of squares

For classification:

• Construct regions which maximize homogeneity

LINEAR FIT

```
library(tree)
fit_tree <- tree::tree(formula = y ~ x, data = tbl_linear)</pre>
summary(fit_tree)
##
## Regression tree:
## tree::tree(formula = y ~ x, data = tbl_linear)
## Number of terminal nodes: 5
## Residual mean deviance: 22.64 = 22530 / 995
## Distribution of residuals:
                            Median
                                        Mean 3rd Qu.
##
        Min.
               1st Qu.
                                                              Max.
## -17.290000 -3.194000 0.000456 0.000000 3.007000 20.740000
```



CATEGORICAL FIT

a	b	output
red	black	1
red	white	1
red	black	0
blue	white	0
blue	black	0

TWO MEASURES OF HOMOGENEITY

$$Gini = \sum p * (1 - p)$$

 $Entropy = -\sum p * log(p)$
MEASURE TOTAL ENTROPY

```
entropy <- function(y) {
   tbl <- tibble(y) %>%
    group_by(y) %>%
    summarise(prob = n()) %>%
    mutate(
        prob = prob / sum(prob)
        , ent = -prob * log(prob))
   tbl$ent %>% sum()
}
```

MEASURE ENTROPY POST-SPLIT

```
entropy_post <- function(tbl, out_col, split_col) {
   split_col <- enquo(split_col)
   out_col <- enquo(out_col)

   tbl %>%
    group_by(!! split_col) %>%
    summarise(
       ent = entropy(!! out_col)
      , group_pct = n() / nrow(tbl)
   ) %>%
    ungroup() %>%
   summarise(
       ent_post = sum(ent * group_pct)
   ) %>%
   pull(ent_post)
}
```

WHICH COLUMN WORKS BETTER ON OUR TOY DATA?

```
entropy(tbl_toy$output)
## [1] 0.6730117
```

```
tbl_toy %>%
    entropy_post(output, a)
## [1] 0.3819085
```

```
tbl_toy %>%
    entropy_post(output, b)
## [1] 0.6591674
```

a	b	output	
red	black	1	
red	white	1	
red	black	0	
blue	white	0	
blue	black	0	

POTENTIAL NODE SPLITS

```
entropy(tbl_players$MultiArrestNum)
## [1] 0.5322599
tbl_players %>%
    entropy_post(MultiArrestNum, PositionType)
## [1] 0.528498
```

```
tbl_players %>%
    entropy_post(MultiArrestNum, Season)
## [1] 0.5092358
```

```
tbl_players %>%
    entropy_post(MultiArrestNum, ArrestSeasonState)
## [1] 0.5313111
```

WHAT SPLITS?

```
library(rpart)
fit_tree <- tree(</pre>
    data = tbl_players
  , formula = MultiArrestFactor ~ PositionType + Season + ArrestSeasonState)
summary(fit_tree)
##
## Classification tree:
## tree(formula = MultiArrestFactor ~ PositionType + Season + ArrestSeasonState,
       data = tbl_players)
##
## Variables actually used in tree construction:
## [1] "Season"
                     "PositionType"
## Number of terminal nodes: 3
## Residual mean deviance: 1.021 = 661.6 / 648
## Misclassification error rate: 0.2197 = 143 / 651
```

PLOT THE TREE

plot(fit_tree)
text(fit_tree, pretty = 0)



NOTE

- 1. Full disclosure: I used both rpart and tree for the fit. For reasons that I've not yet debugged, rpart gave me no nodes.
- 2. A package's insistence on using factors may cause you to lose your mind.

BAGGING/RANDOM FORESTS

- Avoid overfit by bootstrapping
- Fit hundreds of resampled trees
- Take the average of results
- We don't get that sweet tree plot

RANDOM FOREST

```
library(randomForest)
fit_forest <- randomForest(
    formula = MultiArrestFactor ~ PositionType + Season + ArrestSeasonState
   , data = tbl_players
)</pre>
```

VARIABLE IMPORTANCE

varImpPlot(fit_forest)



MULTIPLE CORRESPONDENCE ANALYSIS

WHAT IS MCA?

- PCA, but for categories
- CA, but for multiple variables

WHY MCA?

- Dimensionality reduction
- Could also consider (hierarchical) cluster analysis
- Others?

HOW DOES IT WORK?

- Candidly, I can't easily explain it.
- Creates a "complete disjunctive table", i.e. a "one hot encoding" table
- This creates points in a high-dimensional space
- Synthesizes new dimensions which capture the most variance between the points

COMPLETE DISJUNCTIVE TABLE

id	metro	region	
1	urban	north	
2	urban	south	
3	rural	east	
4	urban	north	

CDT, OR "ONE-HOT ENCODING"

```
tbl_toy_mca_one_hot <- tbl_toy_mca %>%
gather(category, value, -id) %>%
unite(cdt, -id) %>%
mutate(count = 1L) %>%
tidyr::spread(cdt, count, fill = 0L)
```

tbl_toy_mca_one_hot %>% knitr::kable()

id	metro_rural	metro_urban	region_east	region_north	region_south
1	0	1	0	1	0
2	0	1	0	0	1
3	1	0	1	0	0
4	0	1	0	1	0

EXTRACT DATA FOR PROCESSING

```
tbl_cats <- tbl_players %>%
ungroup() %>%
select(
    CrimeCategory, ArrestSeasonState, Conference
   , Division, DayOfWeek, Outcome, Position, PositionType
   , Season) %>%
mutate_if(is.character, as.factor)
library(FactoMineR)
fit_mca <- MCA(tbl_cats, graph = FALSE)</pre>
```

VISUALIZE IN THE REDUCED DIMENSIONS



MCA: CATEGORICAL -> CONTINUOUS

Call: glm(formula = MultiArrestNum ~ 0 + dim_1 + dim_2, family = binomial(), data = tbl_players)

Deviance Residuals: Min 1Q Median 3Q Max -1.815 -1.182 -1.158 -1.129 1.236

Coefficients: Estimate Std. Error z value Pr(>|z|) dim_1 0.27532 0.16245 1.695 0.0901 . dim_2 -0.05509 0.13790 -0.399 0.6895

```
- Signif. codes: 0 '' 0.001 " 0.01 " 0.05 '.' 0.1 " 1
```

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 902.48 on 651 degrees of freedom

Residual deviance: 898.83 on 649 degrees of freedom AIC: 902.83

Number of Fisher Scoring iterations: 4

LET'S MODEL!

HOW WE'LL MODEL

- 1. Pick a performance measure
- 2. Setup cross-validation
- 3. Train some models
- 4. Measure performance

OUR PERFORMANCE MEASURE

Misclassification rate

Other options:

- True positive rate
- False positive rate
- Other confusion matrix metrics
- Area under the curve (AUC): A number close to 1 is good

MEASURES

```
misclass <- function(tbl_test, fit_obj) {
   tbl_test <- tbl_test %>%
    mutate(
        pred = predict(fit_obj, type = 'class', newdata = tbl_test)
        , misclass = pred != MultiArrestFactor
        )
      sum(tbl_test$misclass) / nrow(tbl_test)
}
```

N-FOLD CROSS VALIDATION

library(modelr)
set.seed(1234)
tbl_folds <- crossv_kfold(tbl_players, k = 10)</pre>

TBL_FOLDS

```
tbl_folds %>% head()
## # A tibble: 6 x 3
    train
##
                                   .id
                    test
    <list>
                    <list>
                                   <chr>
##
## 1 <S3: resample> <S3: resample> 01
## 2 <S3: resample> <S3: resample> 02
## 3 <S3: resample> <S3: resample> 03
## 4 <S3: resample> <S3: resample> 04
## 5 <S3: resample> <S3: resample> 05
## 6 <S3: resample> <S3: resample> 06
```

WHAT'S IN TBL_FOLDS?

- Each row in the tibble holds:
- a training resample object
- a test resample object
- an id

A resample object is a list which contains a data frame and a vector of row indices.

```
tbl_folds$train[[1]] %>% class()
## [1] "resample"
```

ASSESS ONE FOLD

```
assess_fold <- function(obj_train, obj_test, method, the_formula) {
   tbl_train <- obj_train %>% as.data.frame()
   tbl_test <- obj_test %>% as.data.frame()
   fit <- do.call(
        method
        , args = list(formula = the_formula, data = tbl_train))
   misclass(tbl_test, fit)
}
one_fold_misclass <- assess_fold(
      tbl_folds$train[[1]]
      , tbl_folds$test[[1]]
      , tree::tree
      , as.formula('MultiArrestFactor ~ PositionType + Season'))</pre>
```

ASSESS ALL FOLDS

```
cross_validate <- function(formula, tbl_folds, method) {</pre>
  map2_dbl(
    tbl_folds$train
  , tbl_folds$test
  , assess_fold
  , method
  , formula
  ) %>% mean()
}
misclasses <- cross_validate(</pre>
    as.formula('MultiArrestFactor ~ PositionType + Season')
  , tbl_folds
  , tree::tree
)
misclasses <- cross_validate(</pre>
    as.formula('MultiArrestFactor ~ PositionType + Season')
  , tbl_folds
  , naive_bayes
)
```

MAKE FORMULAS

```
make_formula <- function(predictors, target, intercept = TRUE) {
   str_predictors <- paste(predictors, collapse = '+')
   if (intercept) {
     str_formula <- paste(target, '~ 1 + ')
   } else {
     str_formula <- paste(target, '~')
   }
   str_formula <- paste(str_formula, str_predictors)
   as.formula(str_formula)
}</pre>
```

A FEW FORMULAS

OUR MODELS TIBBLE

formula
MultiArrestFactor ~ PositionType + Season
MultiArrestFactor ~ PositionType + Season + DayOfWeek
MultiArrestFactor ~ PositionType + Season + DayOfWeek
MultiArrestFactor ~ PositionType + Season + DayOfWeek + Conference
MultiArrestFactor ~ PositionType + Season + DayOfWeek + Conference + Division
MultiArrestFactor ~ PositionType + Season + DayOfWeek + Conference + Division + TeamCity

ASSESS ALL FOLDS, ALL FORMULAS, ALL MODELS

```
tbl_models <- tbl_models %>%
mutate(
    misclass_tree = map_dbl(formula, cross_validate, tbl_folds, tree::tree)
    , misclass_nb = map_dbl(formula, cross_validate, tbl_folds, naive_bayes)
)
```

formula	misclass_tree	misclass_nb
MultiArrestFactor ~ PositionType + Season	0.224289	0.2273427
MultiArrestFactor ~ PositionType + Season + DayOfWeek	0.224289	0.2304196
MultiArrestFactor ~ PositionType + Season + DayOfWeek	0.224289	0.2304196
MultiArrestFactor ~ PositionType + Season + DayOfWeek + Conference	0.224289	0.2319580
MultiArrestFactor ~ PositionType + Season + DayOfWeek + Conference + Division	0.224289	0.2319580
MultiArrestFactor ~ PositionType + Season + DayOfWeek + Conference + Division + TeamCity	0.224289	0.2550350

CONCLUSION

WHAT DID WE LEARN CHARLIE BROWN?

- Categorical data is ubiquitous, but tricky to model
- Non-linear approaches like tree-based methods and Naive Bayes look at categorical differently
- MCA can address "curse of dimensionality" with categorical data
- Let's all keep doing this! Fitting categorical data is hard. Research is light.
Slides may be found here:

http://pirategrunt.com/sparsity_blues/#/

All of the code - even stuff you didn't see - is on GitHub

https://github.com/pirategrunt

THANK YOU!



REFERENCES

- <u>http://www.gastonsanchez.com/visually-enforced/how-to/2012/10/13/MCA-in-R/</u>
- <u>http://rpubs.com/dgrtwo/cv-modelr</u>
- <u>https://drsimonj.svbtle.com/k-fold-cross-validation-with-modelr-and-broom</u>
- <u>http://www.casact.org/pubs/forum/09wforum/flynn_francis.pdf</u>