R Formula Interface

and Model/Design Matrices SYS 6018 | Spring 2023 Rfmla.pdf

```
#-- Required Packages
library(splines)
library(tidyverse)
```

1 Raw input data

The raw input data is often in the form of a data frame (or tibble). For example,

```
#-- Raw Input Data
# cat is categorical with 3 levels: A,B,C
# num is numerical
# y is numerical response variable

Z = tibble(cat=c('A','A','B','B','C','C'), num=1:6, y=rnorm(6))

Z
#> # A tibble: 6 x 3
#> cat num y
#> <chr> <int> <dbl>
#> 1 A 1 -1.01
#> 2 A 2 -0.209
#> 3 B 3 0.262
#> 4 B 4 0.374
#> 5 C 5 -0.666
#> 6 C 6 0.914
```

has three columns, cat is categorical data, num which is numerical data, and y which is the outcome variable.

2 Formula in models

The formula interface in R allows you to make transformations of the input data frame automatically. For example, categorical (or factor) columns will generate the appropriate dummy variables.

The default behavior is to convert categorical data to a *factor* and drop the first level.

The formula interface is easy to use:

```
#- numerical data only
lm(y~num, data=Z)$coef
#> (Intercept)
#> -0.8908
                 0.2386
#- transformations
lm(y~log(num), data=Z)$coef
#> (Intercept) log(num)
#> -0.8509 0.7254
#- use I() to make custom functions
lm(y\sim I(3*num), data=Z)$coef
#> (Intercept) I(3 * num)
#> -0.89075 0.07954
#- we have already seen poly()
lm(y~poly(num, degree = 3), data=Z)$coef
#> (Intercept) poly(num, degree = 3)1 poly(num, degree = 3)2
#> -0.05555 -0.23329
             -0.05555
                            0.99826 -0.23329
#>
#> poly(num, degree = 3)3
       0.92124
#- how about B-splines
library(splines)
lm(y~bs(num), data=Z)$coef
#> (Intercept) bs(num)1 bs(num)2 bs(num)3
#> -1.1227 3.3290 -0.8128 1.8798
#- two predictors
lm(y~cat + num, data=Z)$coef
#> (Intercept) catB catC num
#> -1.8536 -0.7332 -2.5869 0.8299
lm(y~cat + num - 1, data=Z)$coef
#> catA catB catC num
#> -1.8536 -2.5868 -4.4405 0.8299
#- a:b stands for interactions
lm(y~cat + num + cat:num, data=Z)$coef
#> (Intercept) catB catC num catB:num
#> -1.8066 1.7349 -6.7583 0.7986 -0.6873
                                        num catB:num catC:num
                                                              0.7812
#- use . to represent everything in data
lm(y~., data=Z)$coef
#> (Intercept) catB catC num
#> -1.8536 -0.7332 -2.5869 0.8299
lm(y-. - num, data=Z) $coef # use . to include all, then remove some
#> -0.6087 0.9266 0.7328
```

2.1 model.matrix()

Behind the scenes, lm() is calling the function model.matrix() to construct the *model matrix* (also known as a *design matrix*). The model matrix is the real valued X matrix used for calculating the coefficients. You have to pass a formula object into model.matrix().

```
fmla = formula(y~num+cat)
model.matrix(fmla, data=Z)
#> (Intercept) num catB catC
```

```
#> 1
      1 1 0
                       0
#> 2
          1 2 0
#> 3
          1 3 1
                       0
#> 4
           1 4 1
                      0
#> 5
           1 5 0 1
      1 6 0 1
#> attr(, "assign")
#> [1] 0 1 2 2
#> attr(, "contrasts")
#> attr(, "contrasts") $cat
#> [1] "contr.treatment"
fmla = formula(y~num+cat-1) # remove intercept
model.matrix(fmla, data=Z)
#> num catA catB catC
#> 1 1 0 0
#> 2 2 1 0 0
#> 3 3 0 1 0
#> 4 4 0 1 0
#> 5 5 0 0 1
#> 6 6 0 0
#> attr(, "assign")
#> [1] 1 2 2 2
#> attr(, "contrasts")
#> attr(, "contrasts")$cat
#> [1] "contr.treatment"
```

Or, if you are good with data manipulation construct the model matrix manually.

```
library(dplyr)
Ζ 응>응
 transmute(
  intercept = 1,
  x1 = num,
  x2 = num^2,
  x3 = ifelse(cat == 'B', 1, 0), x4 = ifelse(cat == 'C', 1, 0)
 ) 응>응
as.matrix()
#> intercept x1 x2 x3 x4
1 2 4 0 0
#> [2,]
#> [3,]
             1 3 9 1 0
#> [4,]
#> [5,]
             1 4 16 1
             1 5 25 0 1
#> [6,] 1 6 36 0 1
```

Some functions (e.g., glmnet) do not take formulas so you will have to pass in the model matrix X directly. Another word of caution, some functions (again like glmnet) add the intercept automatically so you should not include a columns of ones.

The function lm.fit() fits a linear model from a model matrix:

2.2 Comparison

It is always good to compare the approaches just to make sure there are no mistakes.

```
fmla = formula(y~num+cat + I(num^2) + sqrt(num))
#- lm()
beta.lm = lm(fmla, data=Z)$coef

#- lm.fit()
X = model.matrix(fmla, data=Z)
beta.lmfit = lm.fit(X, Z$y)$coef

#- direct matrix operations
beta.eq = solve(t(X) %*% X) %*% t(X) %*% Z$y
# solve(crossprod(X), crossprod(X, Z$y)) # Alternative
#- output
tibble(beta.lm, beta.lmfit, beta.eq)
```

beta.lm	beta.lmfit	beta.eq
-14.1974	-14.1974	-14.1974
-10.9976	-10.9976	-10.9976
0.5629	0.5629	0.5629
-1.2136	-1.2136	-1.2136
0.6875	0.6875	0.6875
23.4996	23.4996	23.4996