

Eric.Rmd

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```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.1 --
```

```
## v ggplot2 3.3.3      v purrr  0.3.4
## v tibble  3.1.4      v dplyr  1.0.7
## v tidyr   1.1.3      v stringr 1.4.0
## v readr   2.0.1      v forcats 0.5.1
```

```
## -- Conflicts ----- tidyverse_conflicts() --
```

```
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()     masks stats::lag()
```

```
library(mgcv)
```

```
## Loading required package: nlme
```

```
##
```

```
## Attaching package: 'nlme'
```

```
## The following object is masked from 'package:dplyr':
```

```
##
```

```
## collapse
```

```
## This is mgcv 1.8-31. For overview type 'help("mgcv-package")'.
```

```
test_data <- read.csv("data-test.csv")
train_data <- read.csv("data-train.csv")
```

```
train_data <- train_data %>%
  mutate(Fr = as.ordered(Fr)) %>%
  mutate(Re = as.ordered(Re))
unique(train_data$Re)
```

```
## [1] 224 90 398
```

```
## Levels: 90 < 224 < 398
```

```
lm1 <- lm(R_moment_1 ~ St + Re + Fr, data = train_data)
gam1 <- gam(R_moment_1 ~ s(St) + Re + Fr, data = train_data)

gam2 <- gam(R_moment_1 ~ s(St, by = Fr) + Re + Fr, data = train_data)

gam3 <- gam(R_moment_1 ~ s(St, by = Re) + Re + Fr, data = train_data)
```

10-folds Cross validation skeleton code

```
# https://stats.stackexchange.com/questions/61090/how-to-split-a-data-set-to-do-10-fold-cross-validation
set.seed(21)
# Randomly shuffle training data before splitting into 10 folds
shuffled_train <- train_data[sample(nrow(train_data)),]

# Create 10 folds
folds <- cut(seq(1,nrow(train_data)),breaks=10,labels=FALSE)
```

CV for GAM & R-Moment1

```
# error
rmse.cv.gam.inter <- rep(0, 10)
rmse.cv.gam.simple <- rep(0, 10)
# Cross validation: Use gam2 for example
for(i in 1:10){
  #Segment your data by fold using the which() function
  testIndexes <- which(folds==i,arr.ind=TRUE)
  testData <- shuffled_train[testIndexes, ]
  y.test <- testData$R_moment_1
  trainData <- shuffled_train[-testIndexes, ]

  #Use the test and train data
  #1. gam model with interaction
  gam_cv_inter <- gam(R_moment_1 ~ s(St, by = Fr) + Re + Fr, data = trainData)
  pred_gam_inter <- predict.gam(gam_cv_inter, testData, type='response')
  rmse.cv.gam.inter[i] = mean((pred_gam_inter - y.test)^2)

  #2. gam model without interaction
  gam_cv_simple <- gam(R_moment_1 ~ St + Re + Fr, data = trainData)
  pred_gam_simple <- predict.gam(gam_cv_simple, testData, type='response')
  rmse.cv.gam.simple[i] = mean((pred_gam_simple - y.test)^2)
}

print(mean(rmse.cv.gam.inter))
```

```
## [1] 0.0002905743
```

```
print(mean(rmse.cv.gam.simple))
```

```
## [1] 0.0002545483
```

CV for GAM & R-Moment2

```

# error
rmse.cv.gam.inter.2 <- rep(0, 10)
rmse.cv.gam.simple.2 <- rep(0, 10)
# Cross validation: Use gam2 for example
for(i in 1:10){
  #Segment your data by fold using the which() function
  testIndexes <- which(folds==i,arr.ind=TRUE)
  testData <- shuffled_train[testIndexes, ]
  y.test <- testData$R_moment_4
  trainData <- shuffled_train[-testIndexes, ]

  #Use the test and train data
  #1. gam model with interaction
  gam_cv_inter_2 <- gam(R_moment_4 ~ s(St, by = Fr) + Re + Fr, data = trainData)
  pred_gam_inter_2 <- predict.gam(gam_cv_inter_2, testData, type='response')
  rmse.cv.gam.inter.2[i] = mean((pred_gam_inter_2 - y.test)^2)

  #2. gam model without interaction
  gam_cv_simple_2 <- gam(R_moment_4 ~ St + Re + Fr, data = trainData)
  pred_gam_simple_2 <- predict.gam(gam_cv_simple_2, testData, type='response')
  rmse.cv.gam.simple.2[i] = mean((pred_gam_simple_2 - y.test)^2)
}

print(mean(rmse.cv.gam.inter.2))

```

```
## [1] 2.341012e+20
```

```
print(mean(rmse.cv.gam.simple.2))
```

```
## [1] 2.30681e+20
```