

# Cross-Validation

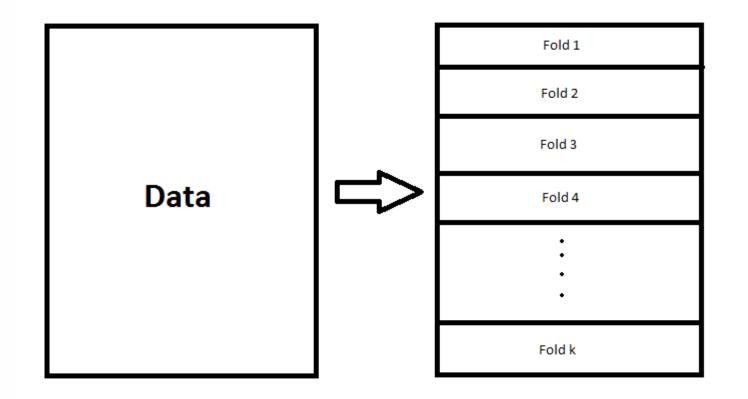
K-Folds



- In the cross-validation, so far we have partitioned the data into two parts or three parts
- In k-folds cross-validation, we divide the data into k equal (more or less) parts and then assuming each fold as test set average the error



#### K-Folds Cross-Validation



 The k-folds cross-validation process starts from dividing the data into k equal parts and that too randomly



# K-Folds

	Data	Model Buliding on Training Set and Then Applying the built model on Test set				
Partitions	Α	Α	Α	Α	Α	Α
	В	В	В	В	В	В
	С	С	С	С	С	С
	D	D	D	D	D	D
	E	E	E	E	E	E
			Test Set			
			Training Set			



### K-Folds Steps

- 1. Consider Fold 1 as a test set for the time being
- 2. With Fold 1 as test set, and remaining all other folds (data) as train set fit a model
- 3. Evaluate the model for the test set (Fold 1) and record the error in the evaluation
- 4. Consider Fold 2 as a test set
- 5. Repeat steps 2 and 3 for Fold 2
- 6. For each of the k folds, repeat 2 and 3
- 7. Aggregate the errors calculated in all the evaluations.



 K-folds can be created using function createFolds() from package caret

Syntax : createFolds(y , k , ...)

Where

y : response variable

k: folds to be created



#### Description

- a cross-section from 1987
- number of observations : 546
- country : Canada

#### • A dataframe containing :

- **price**: sale price of a house
- **lotsize**: the lot size of a property in square feet
- **bedrooms**: number of bedrooms
- **bathrms**: number of full bathrooms
- **stories**: number of stories excluding basement
- driveway: does the house has a driveway?
- recroom: does the house has a recreational room?
- fullbase: does the house has a full finished basement?
- gashw: does the house uses gas for hot water heating?
- airco: does the house has central air conditioning?
- garagepl : number of garage places
- prefarea: is the house located in the preferred neighbourhood of the city?



## R Program (Creating Folds)

```
library(caret)
trainIndex <- createFolds(y = Housing$price , k = 5)
fold1 <- Housing[trainIndex$Fold1,]
fold2 <- Housing[trainIndex$Fold2,]
fold3 <- Housing[trainIndex$Fold3,]
fold4 <- Housing[trainIndex$Fold4,]
fold5 <- Housing[trainIndex$Fold5,]

train1 <- Housing[-trainIndex$Fold1,]
train2 <- Housing[-trainIndex$Fold2,]
train3 <- Housing[-trainIndex$Fold3,]
train4 <- Housing[-trainIndex$Fold4,]
train5 <- Housing[-trainIndex$Fold5,]</pre>
```



### R Program (Evaluation)

```
library(ranger) ## Uses Fast Implementation of Random Forest

################# Fold 1 #############################

model.RF1 <- ranger(price~. , data = train1)
pred.RF1 <- predict(model.RF1 , data = fold1)

MAPE <- function(y, yhat) {
    mean(abs((y - yhat)/y))
}

error1 <- MAPE(fold1$price , pred.RF1$predictions)</pre>
```

• This model building and evaluation is repeated for all 5 folds



#### Aggregating the Error

```
> AvgError <- mean(c(error1,error2,error3,error4,error5))
> AvgError
[1] 0.1825631
> MaxError <- max(c(error1,error2,error3,error4,error5))
> MaxError
[1] 0.201473
```

- Errors on all the folds are averaged or maximum error can be considered.
- Hence, we get more promising estimate of the error.



#### Alternative – caret way

• In caret package, we have a train control function trainControl(), object of which needs to be passed to the train() function itself.

Syntax: trainControl( method, number, ...)

Where

method: resampling method; "boot", "cv", "LOOCV" etc are its possible options

number: number of folds to be created



## Using trainControl()

```
> library(caret)
> model.RF <- train(price~. , data = Housing, method="ranger",
                   trControl = trainControl(method = "cv", number = 5),
                   tuneGrid = expand.grid(mtry=c(2:5)) )
> model.RF
Random Forest
546 samples
11 predictor
No pre-processing
Resampling: Cross-Validated (5 fold)
Summary of sample sizes: 438, 436, 437, 438, 435
Resampling results across tuning parameters:
       RMSE
                 Rsquared
 mtry
       16087.38 0.6541124
       15986.18 0.6465407
       16045.53 0.6413658
       16238.65 0.6308672
RMSE was used to select the optimal model using the smallest value.
The final value used for the model was mtry = 3.
```