```
baggingAndForest.ipynb 
                                                                                                                                                  三 댓글
                                                                                                                                                              🚜 공유 🌣
        파일 수정 보기 삽입 런타임 도구 도움말 모든 변경사항이 저장됨
       + 코드 + 텍스트
                                                                                                                                                  다시 연결 ▼
Q
       [ ] package version(R.version)
             [1] '4.2.0'
[ ] system("gdown --id 1puITTv_Y_2g9rKg7jSaT4XN41HuoFywr")
       [ ] system("gdown --id 1JCy0LLNnrauv8RVi0oZLfjRugEDe_Qc3")
       | hmegtrain <- read.csv("hmegN train.csv", header = T)</pre>
             str(hmeqtrain)
              $ ID : int 4952 5546 938 277 5204 5762 2354 2896 76 4700 ... $ BAD : int 1 1 1 1 1 1 1 1 1 1 ... $ LOAN : int 26100 35000 9100 5700 28100 44000 14200 16000 3900 24800 ... $ MORTDUE: num 73525 391000 17218 58400 61000 ...
              $ VALUE : num 89870 505000 36721 75000 99000 ...

$ REASON : chr "DebtCon" "DebtCon" "HomeImp" ...
                       : chr "Office" "ProfExe" "Other" "ProfExe" ...
              $ JOB
        [ ] hmeqtest <- read.csv("hmeqN_test.csv", header = T)</pre>
             str(hmeqtest)
              'data.frame': 378 obs. of 7 variables:
              $ ID : int 5632 675 3234 3537 3804 926 462 2229 4770 184 ...

$ BAD : int 1 1 1 1 1 1 1 1 1 ...

$ LOAN : int 38700 8000 17300 18600 20000 9000 6900 13700 25000 5000 ...
<>
               $ MORTDUE: num 119847 37871 73000 64248 60336 ...
$ VALUE : num 162365 89870 95000 82690 132430 ...
              REASON: chr "HomeImp" "HomeImp" "DebtCon" ...

$ JOB : chr "ProfExe" "ProfExe" "Other" "Mgr" ...
        결측값 관리
       [ ] colSums(is.na(hmeqtrain)); colSums(is.na(hmeqtest));
                      0 BAD:
                                                0 MORTDUE:
                                                                  0 VALUE:
                                                                                0 REASON:
                      O BAD:
                                  0 LOAN:
                                                0 MORTDUE:
                                                                  0 VALUE:
                                                                                O REASON:
                                                                                                O JOB:
       [ ] install.packages("reshape")
             library(reshape)
             Installing package into '/usr/local/lib/R/site-library'
             (as 'lib' is unspecified)
             also installing the dependencies 'Rcpp', 'plyr'
        변환 하여 넣어주는 작업
        [ ] hmeqtrain <- as.data.frame(hmeqtrain)</pre>
              hmeqtrain$BAD <- as.factor(hmeqtrain$BAD)</pre>
             hmeqtrain$LOAN <- as.numeric(hmeqtrain$LOAN)
             hmeqtrain$REASON <- as.factor(hmeqtrain$REASON)</pre>
             hmeqtrain$JOB <- as.factor(hmeqtrain$JOB)</pre>
             str(hmeqtrain)
             'data.frame': 2000 obs. of 7 variables:
              $ ID
                         : Factor w/ 2 levels "1", "2": 1 1 1 1 1 1 1 1 1 1 ...
               $ MORTDUE: num 73525 391000 17218 58400 61000 ...
               $ VALUE : num 89870 505000 36721 75000 99000 ...
               $ REASON : Factor w/ 3 levels "DebtCon", "HomeImp",..: 1 1 1 2 1 1 1 1 2 1 ... $ JOB : Factor w/ 7 levels "Mgr", "missing",..: 3 5 4 5 1 4 5 4 4 1 ...
        [ ] hmegtest <- as.data.frame(hmegtest)</pre>
             hmegtest$BAD <- as.factor(hmegtest$BAD)</pre>
             hmeqtest$LOAN <- as.numeric(hmeqtest$LOAN)</pre>
              hmeqtest$REASON <- as.factor(hmeqtest$REASON)</pre>
             hmeqtest$JOB <- as.factor(hmeqtest$JOB)</pre>
             str(hmeatest)
              'data.frame': 378 obs. of 7 variables:
                        : int 5632 675 3234 3537 3804 926 462 2229 4770 184 ...
: Factor w/ 2 levels "1","2": 1 1 1 1 1 1 1 1 1 1 ...
              $ ID
               $ BAD
```

```
: num 38700 8000 17300 18600 20000 9000 6900 13700 25000 5000 ...
        $ MORTDUE: num 119847 37871 73000 64248 60336 ...
        $ VALUE : num 162365 89870 95000 82690 132430 ...

$ REASON : Factor w/ 3 levels "DebtCon", "HomeImp",..: 2 2 1 1 1 1 2 1 1 1 ...

$ JOB : Factor w/ 7 levels "Mgr", "missing",..: 5 5 4 1 4 4 6 5 6 4 ...
  [ ] install.packages("ipred")
       library(ipred)
       library(rpart)
       Installing package into '/usr/local/lib/R/site-library'
       (as 'lib' is unspecified)
       also installing the dependencies 'listenv', 'parallelly', 'future', 'globals', 'future.apply', 'progressr', 'numDeriv', 'SQUAREM', 'lava',
- Bagging
  [ ] bagg.hmeq <- bagging(BAD~.-ID, #Y변수 - BAD | X변수 - ID를 뺀 모두
                             data=hmeqtrain,
                              nbag=1000, #bagging 을 1000번 해라
                              control=rpart.control(minsplit=10), #tree split을 일으킬때 최소한 10개는 들어가야 함
       bagg.hmeg
       Bagging classification trees with 1000 bootstrap replications
       Call: bagging.data.frame(formula = BAD ~ . - ID, data = hmeqtrain,
           nbag = 1000, control = rpart.control(minsplit = 10), coob = T)
       Out-of-bag estimate of misclassification error: 0.1565
  Y가 multinomial 이면, out of bag estimate를 계산 못한다는 메시지가 뜨는데, 그러면 coob=T 없애고 다시 돌려야 한다
  coob=T:bagging의 error rate -> 이걸 빼고 돌리면, out of bag error가 안나온다.
                                                                                                                                ↑ ↓ © 目 ‡ 🗓 📋 :
       install.packages("caret")
       library(caret)
       Installing package into '/usr/local/lib/R/site-library'
       (as 'lib' is unspecified)
       also installing the dependencies 'proxy', 'iterators', 'gower', 'hardhat', 'timeDate', 'e1071', 'foreach', 'ModelMetrics', 'pROC', 'recipes
       Loading required package: ggplot2
       Loading required package: lattice
       Warning message in system("timedatectl", intern = TRUE): "running command 'timedatectl' had status 1"
  [ ] bagg.predict <- predict(bagg.hmeq, hmeqtest, type="class")</pre>
       confusionMatrix(bagg.predict, hmeqtest$BAD)
       Confusion Matrix and Statistics
                 Reference
                2 52 184
                       Accuracy: 0.8492
                         95% CI: (0.8091, 0.8837)
           P-Value [Acc > NIR] : < 2.2e-16
                          Kappa : 0.6984
        Mcnemar's Test P-Value : 1.109e-09
                    Sensitivity: 0.7249
                   Specificity: 0.9735
                Pos Pred Value: 0.9648
                Neg Pred Value: 0.7797
                    Prevalence: 0.5000
                Detection Rate: 0.3624
          Detection Prevalence: 0.3757
             Balanced Accuracy: 0.8492
               'Positive' Class: 1
  correct classification rate = (137+184)/378 = 0.8492
```

# - Radom Forest

```
[ ] install.packages("randomForest")
    require(randomForest)
    Installing package into '/usr/local/lib/R/site-library'
(as 'lib' is unspecified)
    Loading required package: randomForest
     randomForest 4.7-1.1
     Type rfNews() to see new features/changes/bug fixes.
     Attaching package: 'randomForest'
     The following object is masked from 'package:ggplot2':
         margin
[ ] rf.hmeq <- randomForest(BAD~.-ID,
                          data=hmegtrain,
                           importance=TRUE,
                           ntree=1000,
                          mtry=2)
     rf.hmea
     Call:
     randomForest(formula = BAD ~ . - ID, data = hmeqtrain, importance = TRUE,
Type of random forest: classification
                                                                                      ntree = 1000, mtry = 2)
                          Number of trees: 1000
     No. of variables tried at each split: 2
     Confusion matrix:
                  0.198
     1 802 198
     2 45 955
                     0.045
입력변수(컬럼)가 5개(root 5 => mtry를 2로 쓴 것이다)
[ ] rf.predict <- predict(rf.hmeq, hmeqtest, type = "class")</pre>
     summary(rf.predict)
[ ] confusionMatrix(rf.predict, hmeqtest$BAD)
     Confusion Matrix and Statistics
               Reference
     Prediction 1 2
1 159 19
              2 30 170
                       Kappa : 0.7407
      Mcnemar's Test P-Value : 0.1531
                 Sensitivity: 0.8413
                Specificity: 0.8995
              Pos Pred Value: 0.8933
              Neg Pred Value: 0.8500
                 Prevalence : 0.5000
              Detection Rate: 0.4206
        Detection Prevalence: 0.4709
           Balanced Accuracy: 0.8704
```

#### [ ] Importante(II:mmcq)

#### A matrix: 5 × 4 of type dbl

	1	2	MeanDecreaseAccuracy	MeanDecreaseGini
LOAN	201.684073	295.22017	312.26298	556.18749
MORTDUE	19.744909	83.24400	93.12678	158.98536
VALUE	-11.782749	87.85284	80.72041	169.84688
REASON	9.476457	49.51785	47.09872	24.20300
JOB	23.855353	97.68437	100.68613	72.23654

LOAN 은 중요한 변수 (312) REASON 은 중요하지 않은 변수 (100)

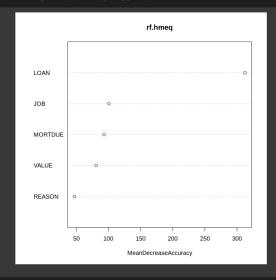
# [ ] importance(rf.hmeq, type=1)

A matrix: 5 × 1 of type dbl

#### MeanDecreaseAccuracy

LOAN	312.26298
MORTDUE	93.12678
VALUE	80.72041
REASON	47.09872
JOB	100.68613

# [ ] varImpPlot(rf.hmeq, type=1)



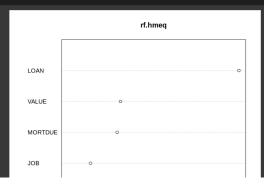
## [ ] importance(rf.hmeq, type=2)

A matrix:  $5 \times 1$  of type dbl

## MeanDecreaseGini

LOAN	556.18749
MORTDUE	158.98536
VALUE	169.84688
REASON	24.20300
JOB	72.23654

# [ ] varImpPlot(rf.hmeq, type=2)



```
REASON
                             MeanDecreaseGini
```

## Boosting - gbm package

```
categorical, numeric 둘다 쓸 수 있는 알고리즘
[ ] install.packages("gbm")
    require(qbm)
    Installing package into '/usr/local/lib/R/site-library'
    (as 'lib' is unspecified)
    Loading required package: gbm
    Loaded gbm 2.1.8
[ ] boost.hmeq <- gbm(BAD~.,-ID,</pre>
                       data=hmeqtrain,
                       distribution="multinomial", # 안쓰면 오류남
                       n.trees=1000,
                       shrinkage=0.01,
                       interaction.depth = 4)
    Warning message:
    "Setting `distribution = "multinomial"` is ill-advised as it is currently broken. It exists only for backwards compatibility. Use at your of
[ ] boost.predict <- predict.gbm(object=boost.hmeq,</pre>
                                  newdata=hmeqtest,
                                  n.trees=1000,
                                  type="response")
[ ] print(boost.predict)
      [1,] 0.99896035 0.0010396536
      [2,] 0.82277840 0.1772216027
       [3,] 0.99892768 0.0010723184
       [4,] 0.99894289 0.0010571094
       [5,] 0.99903882 0.0009611808
       [9,] 0.99853801 0.0014619949
      [11,] 0.99904089 0.0009591101
      [12,] 0.99901384 0.0009861571
      [13,] 0.69254755 0.3074524466
      [16,] 0.16383326 0.8361667428
      [19,] 0.99899695 0.0010030488
     [20,] 0.99855856 0.0014414408
      [21,] 0.99896249 0.0010375073
      [22,] 0.12394018 0.8760598154
      [23,] 0.99889786 0.0011021387
      [24,] 0.99897335 0.0010266484
      [26,] 0.99895963 0.0010403723
      [28,] 0.99899318 0.0010068163
      [29,] 0.99899271 0.0010072917
      [30,] 0.99901384 0.0009861571
      [31,] 0.79038312 0.2096168840
      [32,] 0.99899695 0.0010030488
      [33,] 0.63893614 0.3610638612
      [34,] 0.99899271 0.0010072917
      [36,] 0.99899695 0.0010030488
      [38,] 0.99840736 0.0015926358
      [39,] 0.99897730 0.0010226965
      [40,] 0.99890508 0.0010949153
      [41,] 0.86944124 0.1305587644
      [42,] 0.80029213 0.1997078680
```

[ ] result = data.frame(hmeqtest\$BAD, value) #실제값 print(result)

```
hmeqtest.BAD value
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24
25
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52
53
```

[ ] with(result, table(hmeqtest.BAD, value))

value hmeqtest.BAD 1 2 1 160 29 2 15 174

