# R Notebook

## Parametros:

**Measure =** Accuracy
**Columns =** sampling, weight_space, ruspool
**Performance =** tuning_measure
**Filter keys =** NULL
**Filter values =** NULL

```
library("scmamp")
library(dplyr)
```

## Tratamento dos dados

Carregando data set compilado

```
ds = read.csv("/home/rodrigo/Dropbox/UNICAMP/IC/estudo_cost_learning/SummaryResults/summary_compilation_
summary(ds)
```

```
##                  learner      weight_space
##  classif.ksvm        :17100   Mode :logical
##  classif.randomForest:17100   FALSE:41040
##  classif.xgboost     :17100   TRUE :10260
##                               NA's :0
##
##
##
##                              measure        sampling      ruspool
##  Accuracy                        :10260   ADASYN:10260   Mode :logical
##  Area under the curve            :10260   FALSE :30780   FALSE:41040
##  F1 measure                      :10260   SMOTE :10260   TRUE :10260
##  G-mean                          :10260                  NA's :0
##  Matthews correlation coefficient:10260
##
##
##  tuning_measure    holdout_measure   holdout_measure_residual
##  Min.   :-0.1277   Min.   :-0.2120   Min.   :-0.4658
##  1st Qu.: 0.5924   1st Qu.: 0.3114   1st Qu.: 0.1648
##  Median : 0.9624   Median : 0.8193   Median : 0.5192
##  Mean   : 0.7570   Mean   : 0.6469   Mean   : 0.5099
##  3rd Qu.: 0.9965   3rd Qu.: 0.9879   3rd Qu.: 0.8636
##  Max.   : 1.0000   Max.   : 1.0000   Max.   : 1.0000
##  NA's   :1761      NA's   :1761      NA's   :1761
##  iteration_count                      dataset        imba.rate
##  Min.   :1       abalone                  :  900   Min.   :0.0010
##  1st Qu.:1       adult                    :  900   1st Qu.:0.0100
##  Median :2       bank                     :  900   Median :0.0300
##  Mean   :2       car                      :  900   Mean   :0.0286
##  3rd Qu.:3       cardiotocography-10clases:  900   3rd Qu.:0.0500
##  Max.   :3       cardiotocography-3clases :  900   Max.   :0.0500
```

```
##  NA's   :1761    (Other)                  :45900
```

Filtrando pela metrica

```
ds = filter(ds, measure == params$measure)
```

Filtrando o data set

```
if(params$filter_keys != 'NULL' && !is.null(params$filter_keys)){
  ds = filter_at(ds, .vars = params$filter_keys, .vars_predicate = any_vars(. == params$filter_values))
}

summary(ds)
```

```
##                   learner     weight_space
##  classif.ksvm         :3420   Mode :logical
##  classif.randomForest:3420   FALSE:8208
##  classif.xgboost      :3420   TRUE :2052
##                               NA's :0
##
##
##
##                               measure        sampling     ruspool
##  Accuracy                         :10260   ADASYN:2052   Mode :logical
##  Area under the curve             :    0   FALSE :6156   FALSE:8208
##  F1 measure                       :    0   SMOTE :2052   TRUE :2052
##  G-mean                           :    0                 NA's :0
##  Matthews correlation coefficient:    0
##
##
##  tuning_measure   holdout_measure  holdout_measure_residual
##  Min.   :0.0904   Min.   :0.0152   Min.   :0.0346
##  1st Qu.:0.9591   1st Qu.:0.9535   1st Qu.:0.3643
##  Median :0.9861   Median :0.9800   Median :0.7162
##  Mean   :0.9546   Mean   :0.9493   Mean   :0.6531
##  3rd Qu.:0.9959   3rd Qu.:0.9925   3rd Qu.:0.9406
##  Max.   :1.0000   Max.   :1.0000   Max.   :1.0000
##  NA's   :348      NA's   :348      NA's   :348
##  iteration_count                      dataset       imba.rate
##  Min.   :1       abalone                  : 180   Min.   :0.0010
##  1st Qu.:1       adult                    : 180   1st Qu.:0.0100
##  Median :2       bank                     : 180   Median :0.0300
##  Mean   :2       car                      : 180   Mean   :0.0286
##  3rd Qu.:3       cardiotocography-10clases: 180   3rd Qu.:0.0500
##  Max.   :3       cardiotocography-3clases : 180   Max.   :0.0500
##  NA's   :348     (Other)                  :9180
```

Computando as médias das iteracoes

```
ds = group_by(ds, learner , weight_space , measure , sampling , ruspool , dataset , imba.rate)
ds = summarise(ds, tuning_measure = mean(tuning_measure), holdout_measure = mean(holdout_measure),
               holdout_measure_residual = mean(holdout_measure_residual))

ds = as.data.frame(ds)
```

Criando dataframe

```r
# Dividindo o ds em n, um para cada técnica
splited_df = ds %>% group_by_at(.vars = params$columns) %>% do(vals = as.data.frame(.)) %>% select(vals)

# Juntando cada uma das partes horizontalmente em um data set
df_tec_wide = do.call("cbind", splited_df)

# Renomeando duplicacao de nomes
colnames(df_tec_wide) = make.unique(colnames(df_tec_wide))

# Selecionando apenas as medidas da performance escolhida
df_tec_wide_residual = select(df_tec_wide, matches(paste("^", params$performance, "$|", params$performa

# Renomeando colunas
new_names = NULL
for(i in (1:length(splited_df))){
  id = toString(sapply(splited_df[[i]][1, params$columns], as.character))
  new_names = c(new_names, id)
}
colnames(df_tec_wide_residual) = new_names

# Verificando a dimensao do df
dim(df_tec_wide_residual)
```

```
## [1] 684   5
```

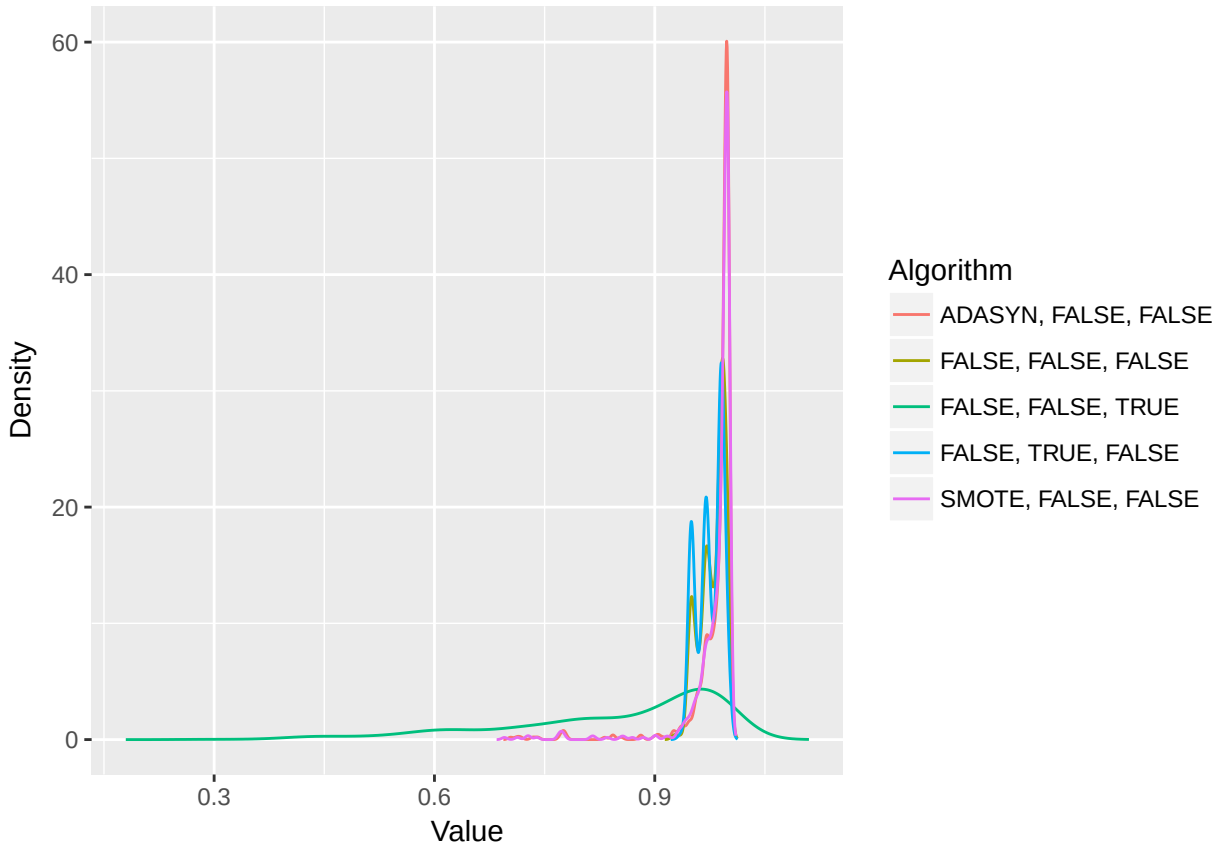
```r
# Renomeando a variavel
df = df_tec_wide_residual

summary(df)
```

```
##  ADASYN, FALSE, FALSE FALSE, FALSE, FALSE FALSE, FALSE, TRUE
##  Min.   :0.7035       Min.   :0.9264      Min.   :0.2896
##  1st Qu.:0.9814       1st Qu.:0.9694      1st Qu.:0.7775
##  Median :0.9948       Median :0.9875      Median :0.9095
##  Mean   :0.9832       Mean   :0.9799      Mean   :0.8529
##  3rd Qu.:0.9988       3rd Qu.:0.9931      3rd Qu.:0.9695
##  Max.   :1.0000       Max.   :1.0000      Max.   :1.0000
##  NA's   :43           NA's   :11          NA's   :14
##  FALSE, TRUE, FALSE SMOTE, FALSE, FALSE
##  Min.   :0.9346     Min.   :0.6952
##  1st Qu.:0.9642     1st Qu.:0.9809
##  Median :0.9794     Median :0.9947
##  Mean   :0.9762     Mean   :0.9829
##  3rd Qu.:0.9900     3rd Qu.:0.9992
##  Max.   :1.0000     Max.   :1.0000
##  NA's   :13         NA's   :35
```

## Fazendo teste de normalidade

```r
plotDensities(data = na.omit(df))
```




## Testando as diferencas

```
friedmanTest(df)
```

```
##
##  Friedman's rank sum test
##
## data:  df
## Friedman's chi-squared = 1312.3, df = 4, p-value < 2.2e-16
```

## Testando as diferencas par a par

```
test <- nemenyiTest (df, alpha=0.05)
abs(test$diff.matrix) > test$statistic
```

```
##      ADASYN, FALSE, FALSE FALSE, FALSE, FALSE FALSE, FALSE, TRUE
## [1,]                FALSE                TRUE               TRUE
## [2,]                 TRUE               FALSE               TRUE
## [3,]                 TRUE                TRUE              FALSE
## [4,]                 TRUE                TRUE               TRUE
## [5,]                FALSE                TRUE               TRUE
##      FALSE, TRUE, FALSE SMOTE, FALSE, FALSE
## [1,]               TRUE               FALSE
```

```
## [2,]              TRUE              TRUE
## [3,]              TRUE              TRUE
## [4,]             FALSE              TRUE
## [5,]              TRUE             FALSE
```

## Plotando grafico de Critical Diference

```
result = tryCatch({
    plotCD(df, alpha=0.05, cex = 0.35)
}, error = function(e) {})
```