

R Notebook

Parametros:

Measure = Area under the curve
Columns = sampling, weight_space, ruspool
Performance = holdout_measure
Filter keys = imba.rate
Filter values = 0.03

```
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      :990  Mode :logical  
## classif.randomForest:990 FALSE:2376  
## classif.xgboost    :990  TRUE :594  
##                   NA's :0  
##  
##  
##  
##           measure      sampling      ruspool  
## Accuracy           : 0  ADASYN: 594  Mode :logical  
## Area under the curve :2970 FALSE :1782 FALSE:2376  
## F1 measure           : 0  SMOTE : 594  TRUE :594  
## G-mean              : 0                   NA's :0  
## Matthews correlation coefficient: 0  
##  
##  
## tuning_measure  holdout_measure  holdout_measure_residual  
## Min. :0.3023  Min. :0.0000  Min. :0.00057  
## 1st Qu.:0.8996  1st Qu.:0.8212  1st Qu.:0.67875  
## Median :0.9945  Median :0.9789  Median :0.87915  
## Mean :0.9169  Mean :0.8812  Mean :0.81326  
## 3rd Qu.:0.9999  3rd Qu.:0.9994  3rd Qu.:0.98229  
## Max. :1.0000  Max. :1.0000  Max. :1.00000  
## NA's :96      NA's :96      NA's :96  
## iteration_count      dataset      imba.rate  
## Min. :1      abalone : 45  Min. :0.03  
## 1st Qu.:1      adult : 45  1st Qu.:0.03  
## Median :2      annealing : 45  Median :0.03  
## Mean :2      arrhythmia : 45  Mean :0.03  
## 3rd Qu.:3      balance-scale: 45  3rd Qu.:0.03  
## Max. :3      bank : 45  Max. :0.03  
## NA's :96      (Other) :2700
```

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

```

# 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$performance)))

# 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] 198 5
```

```

# Renomeando a variavel
df = df_tec_wide_residual

summary(df)

```

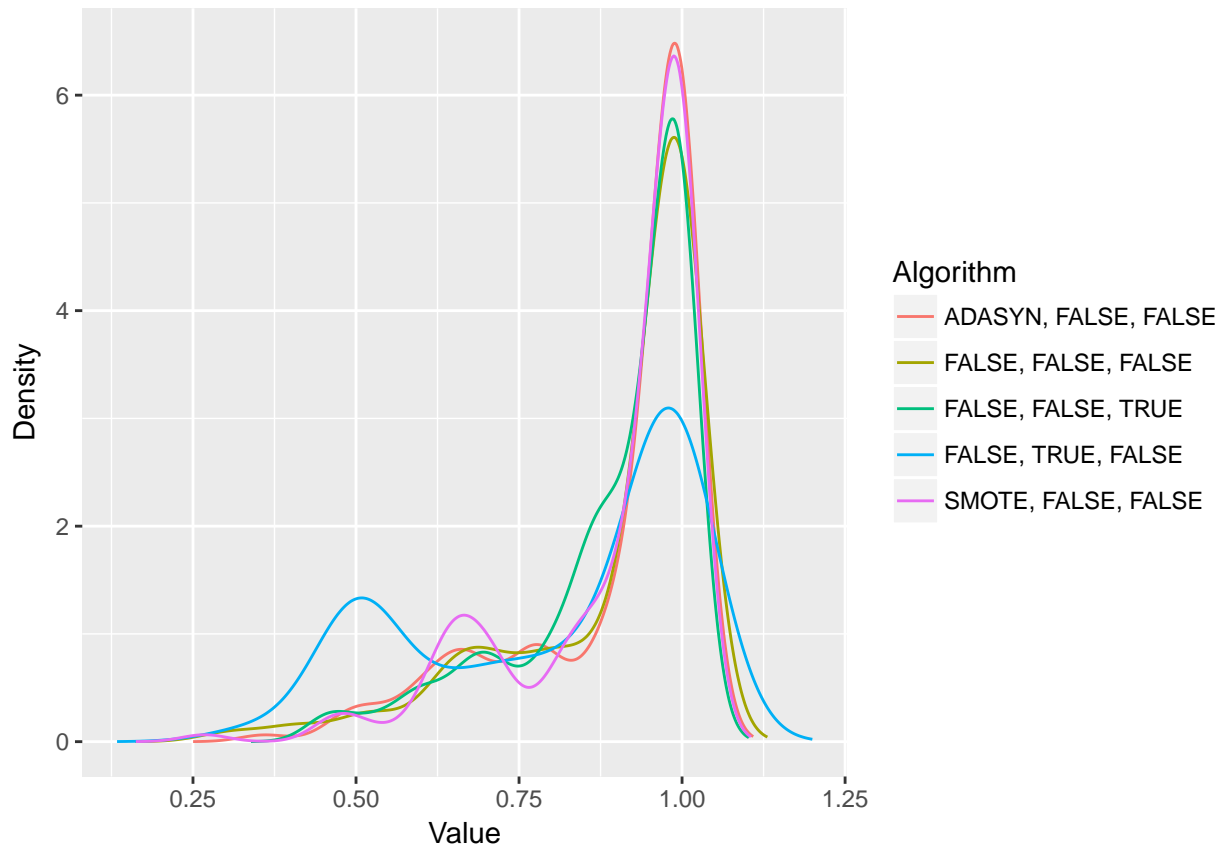
```

## ADASYN, FALSE, FALSE FALSE, FALSE, FALSE FALSE, FALSE, TRUE
## Min. :0.3593 Min. :0.2924 Min. :0.4413
## 1st Qu.:0.8462 1st Qu.:0.8252 1st Qu.:0.8593
## Median :0.9782 Median :0.9769 Median :0.9556
## Mean :0.8994 Mean :0.8965 Mean :0.8961
## 3rd Qu.:0.9985 3rd Qu.:0.9991 3rd Qu.:0.9967
## Max. :1.0000 Max. :1.0000 Max. :1.0000
## NA's :10 NA's :2 NA's :6
## FALSE, TRUE, FALSE SMOTE, FALSE, FALSE
## Min. :0.3333 Min. :0.2679
## 1st Qu.:0.6520 1st Qu.:0.8503
## Median :0.9258 Median :0.9693
## Mean :0.8189 Mean :0.8959
## 3rd Qu.:0.9957 3rd Qu.:0.9977
## Max. :1.0000 Max. :1.0000
## NA's :5 NA's :9

```

Fazendo teste de normalidade

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



Testando as diferencas

```
friedmanTest(df)
```

```
##
## Friedman's rank sum test
##
## data: df
## Friedman's chi-squared = 53.327, df = 4, p-value = 7.278e-11
```

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

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

Plotando grafico de Critical Difference

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

