

svm_mejor

Laura Belmonte

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Probamos con el svm radial, sigmoidal y polinomial:

```
library(recipes)
```

```
## Loading required package: dplyr
```

```
##
```

```
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
```

```
##
```

```
## filter, lag
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
## intersect, setdiff, setequal, union
```

```
## Registered S3 method overwritten by 'future':
```

```
## method from
```

```
## all.equal.connection parallelly
```

```
##
```

```
## Attaching package: 'recipes'
```

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

```
##
```

```
## step
```

```
library(e1071)
```

```
library(ggplot2)
```

```
library(caret)
```

```
## Loading required package: lattice
```

```
library(pROC)
```

```
## Type 'citation("pROC")' for a citation.
```

```
##
## Attaching package: 'pROC'

## The following objects are masked from 'package:stats':
##
##      cov, smooth, var

library(dplyr)

load("~/Documents/GitHub/Mineria/DATA/dataaaaaaaaaaaaa.RData")
bd <- data_reducida

set.seed(123)

rec <- recipe(Exited ~ ., data = bd) %>%
  step_dummy(all_nominal_predictors(), -group, one_hot = TRUE)

bd <- prep(rec) %>% bake(new_data = NULL)

# División train/test

trainbase <- bd[bd$group == "train", ]
trainbase$group <- NULL

testbase <- bd[bd$group == "test", ]
testbase$group <- NULL

ind <- sample(1:nrow(trainbase), 0.7*nrow(trainbase))
train <- trainbase[ind, ]
test <- trainbase[-ind, ]

optimizar_kernel <- function(kernel_name, train_data, test_data,
                             thresholds = seq(0.05, 0.50, 0.01),
                             class_weights = c("0" = 1, "1" = 2)) {

  # -----
  # Grid por kernel
  # -----
  if (kernel_name == "radial") {
    grid <- expand.grid(
      cost = c(1, 5, 10, 20),
      gamma = c(0.001, 0.01, 0.05, 0.1)
    )
  }

  if (kernel_name == "polynomial") {
    grid <- expand.grid(
      cost = c(1, 5, 10),
      gamma = c(0.001, 0.01, 0.1),
      degree = c(2, 3, 4),
      coef0 = c(0, 1)
    )
  }
}
```

```

if (kernel_name == "sigmoid") {
  grid <- expand.grid(
    cost = c(1, 5, 10),
    gamma = c(0.001, 0.01, 0.1),
    coef0 = c(-1, 0, 1)
  )
}

mejor <- NULL
mejor_f1 <- -Inf

for (i in 1:nrow(grid)) {

  params <- grid[i, ]

  # Parámetros dinámicos
  args <- list(
    formula = Exited ~ .,
    data = train_data,
    kernel = kernel_name,
    cost = params$cost,
    gamma = params$gamma,
    class.weights = class_weights,
    probability = TRUE
  )

  if ("degree" %in% names(params)) args$degree <- params$degree
  if ("coef0" %in% names(params)) args$coef0 <- params$coef0

  modelo <- do.call(svm, args)

  # ---- Probabilidades TRAIN y TEST ----
  prob_train <- attr(predict(modelo, train_data, probability = TRUE), "probabilities")[, "1"]
  prob_test <- attr(predict(modelo, test_data, probability = TRUE), "probabilities")[, "1"]

  for (t in thresholds) {

    # Train
    pred_train <- ifelse(prob_train >= t, "1", "0")
    cm_train <- confusionMatrix(
      factor(pred_train),
      factor(train_data$Exited),
      positive = "1"
    )
    f1_train <- cm_train$byClass["F1"]

    # Test
    pred_test <- ifelse(prob_test >= t, "1", "0")
    cm_test <- confusionMatrix(
      factor(pred_test),
      factor(test_data$Exited),
      positive = "1"
    )
  }
}

```

```

f1_test <- cm_test$byClass["F1"]

# --- AUC ---
auc_val <- as.numeric(auc(test_data$Exited, prob_test, quiet = TRUE))

# -----
# Guardar mejor según F1_test
# -----
if (!is.na(f1_test) && f1_test > mejor_f1) {

  mejor_f1 <- f1_test

  mejor <- data.frame(
    Kernel = kernel_name,
    Cost = params$cost,
    Gamma = params$gamma,
    Degree = ifelse("degree" %in% names(params), params$degree, NA),
    Coef0 = ifelse("coef0" %in% names(params), params$coef0, NA),
    Threshold = t,
    F1_Train = f1_train,
    F1_Test = f1_test,
    Recall_Test = cm_test$byClass["Recall"],
    Precision_Test = cm_test$byClass["Precision"],
    Specificity_Test = cm_test$byClass["Specificity"],
    Accuracy_Test = cm_test$overall["Accuracy"],
    AUC = auc_val,
    stringsAsFactors = FALSE
  )
}

return(mejor)
}

#Optimización
kernels <- c("radial", "polynomial", "sigmoid")

resultados <- lapply(kernels, function(k) {
  optimizar_kernel(k, train, test)
})

final <- bind_rows(resultados) %>%
  arrange(desc(F1_Test))

print(final)

```

```

##           Kernel Cost Gamma Degree Coef0 Threshold  F1_Train  F1_Test
## F1...1 polynomial    1  0.10     4     0      0.17 0.5110861 0.4611006
## F1...2   radial     5  0.05    NA    NA      0.14 0.4952648 0.4569939
## F1...3   sigmoid    10  0.01    NA   -1      0.18 0.4962687 0.4514339
##           Recall_Test Precision_Test Specificity_Test Accuracy_Test      AUC
## F1...1   0.5758294      0.3844937      0.7681764      0.7295238 0.6998197

```

## F1...2	0.6232227	0.3607682	0.7222884	0.7023810	0.6898481
## F1...3	0.5781991	0.3702580	0.7526818	0.7176190	0.7008845