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FuzzyART: An R Package for ART-based Clustering.

Louis Steinmeister* Donald C. Wunsch II*

2021-06-09

Abstract

Adaptive Resonance Theory (ART) was introduced by Steven Grossberg as a theory of human cognitive information processing (Grossberg 1976, 1980). Extending the capabilities of the ART 1 model, which can learn to categorize patterns in binary data, fuzzy ART as described in (Carpenter, Grossberg, and Rosen 1991) has become one of the most commenly used Adaptive Resonance Theory models (Brito da Silva, Elnabarawy, and Wunsch 2019). By incorporating fuzzy set theroy operators, fuzzy ART is capable of learning from binaray and bounded real valued data. Its advantage over other unsupervised learning algorithms lies in the flexibility of the learning rule. If a given input feature does not resemble a known category satisfactorily, as determined by the vigilance test, a new category is initialized. Hence, the total number of categories (or clusters) is not determined a-priori, like k-means, but chosen in accordance with the data and the context of already learnt representations. This vignette explores the use of the fuzzy ART implementation as provided by the FuzzyART R package.

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Packacke Usage

Feel free to use this package as specified by the license (MIT). However, please consider citing this work in any publication that this package may contribute to.

To install the package run

```
devtools::install_gitlab(repo = "acil-group/rFuzzyART", host = "git.mst.edu")
```

and run

```
library(FuzzyART)
```

to load the package for use.

Training

Before the fuzzy ART model can be trained, one needs to determine a minimum set of parameters: - rho: Vigilance parameter in (0,1). - alpha: Choice parameter alpha > 0. Can be viewed as a regularization

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parameter penalizing large weights. - beta: Learning rate in (0,1).

Input Pre-processing

Before training, it is important to remember scaling the inputs to lie in the d-dimensional unit hypercube $([0,1]^d)$, where d is the dimension of the inputs. In other words, each input variable needs to be normalized to the interval [0,1]. This can easily be done with the normalize() function.

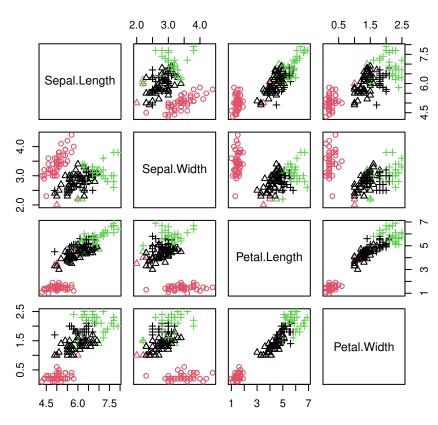
```
library(FuzzyART)
library(mclust)
#> Warning: package 'mclust' was built under R version 4.1.0
#> Package 'mclust' version 5.4.7
#> Type 'citation("mclust")' for citing this R package in publications.
print("Original data:")
#> [1] "Original data:"
summary(iris)
                     Sepal.Width
#>
     Sepal.Length
                                     Petal.Length
                                                     Petal.Width
#>
   Min.
           :4.300
                    Min.
                          :2.000
                                    Min.
                                         :1.000
                                                    Min.
                                                           :0.100
#>
   1st Qu.:5.100
                    1st Qu.:2.800
                                    1st Qu.:1.600
                                                    1st Qu.:0.300
#>
   Median :5.800
                   Median :3.000
                                    Median :4.350
                                                    Median :1.300
#>
  Mean
           :5.843
                   Mean
                         :3.057
                                    Mean
                                          :3.758
                                                    Mean
                                                          :1.199
#>
   3rd Qu.:6.400
                    3rd Qu.:3.300
                                    3rd Qu.:5.100
                                                    3rd Qu.:1.800
#>
   Max.
           :7.900
                    Max.
                           :4.400
                                    Max.
                                           :6.900
                                                    Max.
                                                           :2.500
#>
          Species
#>
              :50
   setosa
   versicolor:50
#>
   virginica :50
#>
#>
#>
#>
print("Normalized data:")
#> [1] "Normalized data:"
iris.normalized = normalize(df = subset(iris,select = -Species))
summary(iris.normalized)
#>
     Sepal.Length
                      Sepal.Width
                                      Petal.Length
                                                        Petal.Width
#>
  Min.
           :0.0000
                     Min.
                           :0.0000
                                      Min.
                                             :0.0000
                                                       Min.
                                                              :0.00000
   1st Qu.:0.2222
                     1st Qu.:0.3333
                                      1st Qu.:0.1017
                                                       1st Qu.:0.08333
#>
#>
   Median :0.4167
                     Median :0.4167
                                      Median :0.5678
                                                       Median :0.50000
                            :0.4406
           :0.4287
#> Mean
                     Mean
                                      Mean
                                             :0.4675
                                                       Mean
                                                              :0.45806
#>
  3rd Qu.:0.5833
                     3rd Qu.:0.5417
                                      3rd Qu.:0.6949
                                                       3rd Qu.:0.70833
                    Max. :1.0000
                                                       Max. :1.00000
#>
  Max. :1.0000
                                      Max. :1.0000
```

Training the ART model

In our Iris example we shall use parameters close to the ones used in (Hoa and Bui 2012); that is alpha \approx 0.8, beta \approx 0.1, rho \approx 0.5. For the wine dataset, the parameters as specified in (Elnabarawy, Tauritz, and Wunsch 2017) appear superior. We will demonstrate to power of this implementation on a number of popular datasets.

The true membership of individual observations is indicated by the symbol while the color corresponds to the category membership according to our trained fuzzy ART model. The Rand Index measures the similarity between two sets of clustering partitions. In this case, we benchmark the performance of the unsupervised fuzzy ART model against the ground truth, the labels associated with each observation.

Iris

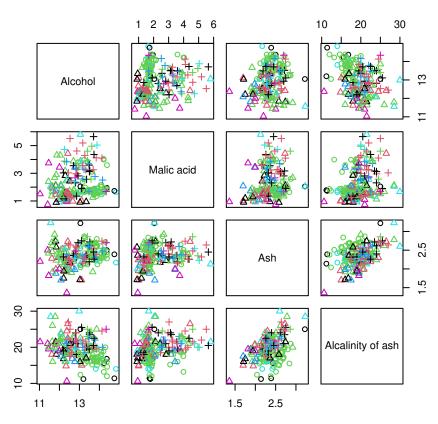


Dataset: Iris -- Rand Index: 0.61

Wine

Note that for a better presentation, we will only be displaying the first four features of the wine dataset. However, all features were used during training.

```
# load the wine dataset
wine.address <- "http://archive.ics.uci.edu/ml/machine-learning-databases/wine/wine.data"
wine <- read.csv(wine.address,header = FALSE)
wine.colnames = c("Label", "Alcohol", "Malic acid", "Ash", "Alcalinity of ash", "Magnesium",
"Total phenols", "Flavanoids", "Nonflavanoid phenols", "Proanthocyanins",</pre>
```



Dataset : Wine -- Rand Index: 0.14

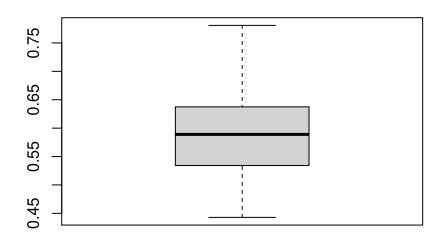
Evaluation of package performance

The examples as shown here certainly have further potential for fine tuning. Nevertheless, in reference to results achieved in (Illetskova et al. 2019), the achievable performance as demonstrated here appear to be on par with the baseline algorithm (as described in the cited work). This can be verified via:

```
#Iris
inputs = subset(iris,select = -Species)
labels.true = as.numeric(unlist(iris$Species))
```

```
normalized_inputs = normalize(df = inputs)
test_iris = function(seed)
ſ
  mod = FuzzyART_train(normalized_inputs,alpha = .8,rho = .5,
                         beta = .12,max_epochs = 1000,max_clusters =20,
                         eps = 10^-8, random_seed = seed, show_status = FALSE,
                         beta_decay = .9)
  return(adjustedRandIndex(mod$Labels,labels.true))
}
res_iris = sapply(X = 1:50,FUN = test_iris)
print(paste0("Mean: ", mean(res_iris)))
print(paste0("StD: ",sqrt(var(res_iris))))
boxplot(res iris, main = "Boxplot of Rand Index -- Iris")
#Wine
inputs = subset(wine, select = -Label)
labels.true = wine$Label
normalized_inputs = normalize(df = inputs)
test_wine = function(seed)
{
  mod = FuzzyART_train(normalized_inputs,alpha = .8679,rho = .375,
                        beta = .9797,max_epochs = 2000,max_clusters =20,
                         eps = 10^-8, random_seed = seed, show_status = FALSE,
                        beta decay = .9)
  return(adjustedRandIndex(mod$Labels,labels.true))
}
res_wine = sapply(X = 1:50,FUN = test_wine)
print(paste0("Mean: ", mean(res_wine)))
print(paste0("StD: ",sqrt(var(res_wine))))
boxplot(res_wine, main = "Boxplot of Rand Index -- Wine")
#> [1] "Mean: 0.588421981887455"
#> [1] "StD: 0.0825407082831491"
#> Registered S3 method overwritten by 'GGally':
#>
    method from
#>
     +.gg
            ggplot2
#> Registered S3 method overwritten by 'sets':
#> method
                   from
#>
    print.element ggplot2
<code>#> Warning: replacing previous import 'GGally::%>%'</code> by <code>'sets::%>%'</code> when loading
#> 'bootcluster'
```

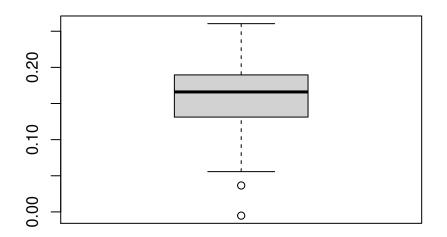
Boxplot of Rand Index -- Iris



#> [1] "Mean: 0.157224686186454"

#> [1] "StD: 0.0490131046535643"

Boxplot of Rand Index -- Wine



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