## Comparing speed of packages for computing ROC curves

Toby Dylan Hocking

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#### 1 Introduction

The goal of this vignette is to compare how long it takes to compute the ROC curves and AUC using different R packages: WeightedROC, ROCR, pROC, glmnet.

#### 2 Data

The data set I will analyze is the spam data set discussed in the book "The Elements of Statistical Learning" by Hastie, Tibshirani, and Friedman. The book is available to read for free as a downloadable PDF at

http://statweb.stanford.edu/~tibs/ElemStatLearn/

The data set is available in the R package ElemStatLearn:

```
> library(ElemStatLearn)
> data(spam)
> is.label <- names(spam) == "spam"</pre>
> X <- as.matrix(spam[,!is.label])</pre>
> y <- spam[,is.label]</pre>
> set.seed(1)
> train.i <- sample(nrow(spam), nrow(spam)/2)</pre>
> sets <-
    list(train=list(features=X[train.i, ], label=y[train.i]),
         test=list(features=X[-train.i, ], label=y[-train.i]))
> str(sets)
List of 2
 $ train:List of 2
  ..$ features: num [1:2300, 1:57] 0 0 0 0 0 0 0 0 0 0 ...
  ... - attr(*, "dimnames")=List of 2
  ....$ : chr [1:2300] "1222" "1712" "2635" "4176" ...
  ....$ : chr [1:57] "A.1" "A.2" "A.3" "A.4" ...
              : Factor w/ 2 levels "email", "spam": 2 2 1 1 2 1 1 1 1 2 ...
  ..$ label
 $ test :List of 2
  ..$ features: num [1:2301, 1:57] 0 0 0.06 0 0 0 0 0 0 0.05 ...
  ...- attr(*, "dimnames")=List of 2
  ....$ : chr [1:2301] "1" "7" "10" "12" ...
  ....$ : chr [1:57] "A.1" "A.2" "A.3" "A.4" ...
              : Factor w/ 2 levels "email", "spam": 2 2 2 2 2 2 2 2 2 2 ...
  ..$ label
```

I divided the data set into half train, half test. I will fit a model on the training set and see if it works on the test set.

### 3 Model fitting

Below, I fit an L1-regularized logistic regression model to the spam training set.

```
> library(glmnet)
> system.time({
+  fit <- cv.glmnet(sets$train$features, sets$train$label, family="binomial")
+ })
  user system elapsed
16.572  0.012  16.640</pre>
```

On my Intel i7 2.8GHz CPU, it took about 10 seconds to fit the model.

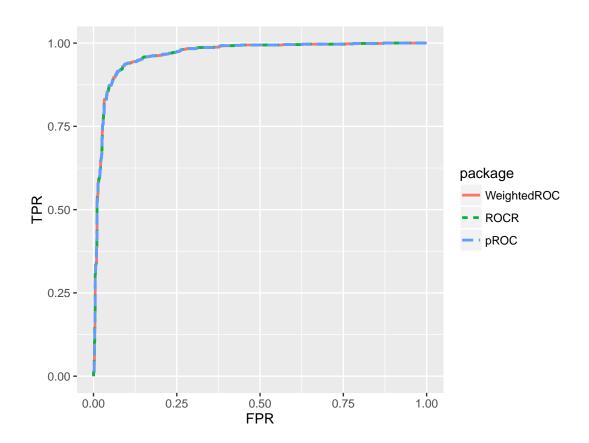
### 4 Timing test ROC curve computation

ROC analysis is useful for evaluating binary classifiers. Below we compute the ROC curves using WeightedROC, pROC, and ROCR.

```
> library(WeightedROC)
> library(ROCR)
> library(pROC)
> set <- sets$test
> guess <- predict(fit, set$features)</pre>
> if(require(microbenchmark)){
    microbenchmark(WeightedROC={
      wroc <- WeightedROC(guess, set$label)</pre>
    }, ROCR={
      pred <- prediction(guess, set$label)</pre>
      perf <- performance(pred, "tpr", "fpr")</pre>
    }, pROC={
      proc <- roc(set$label, guess, algorithm=2)</pre>
Unit: milliseconds
                                                                       max neval
        expr
                                                median
                    min
                               lq
                                        mean
                                                              uq
WeightedROC 3.214201
                        3.430741 4.563544 3.924361 5.35343 13.49265
                                                                             100
                                                                             100
        ROCR 6.835659 7.247196 9.522626 10.115177 11.20842 16.45128
        pROC 27.939451 28.895652 39.996864 41.944814 49.70992 55.11324
cld
a
 b
```

It is clear that WeightedROC is the fastest computation (on my computer, median 1.8 milliseconds), followed by ROCR (7.2 milliseconds), and finally pROC (26.4 milliseconds). However, all of the ROC computations are much faster than the model fitting (10 seconds). Below, we plot the ROC curves.

```
+ data=roc.curves, size=1)+
+ coord_equal()
> print(rocPlot)
>
```



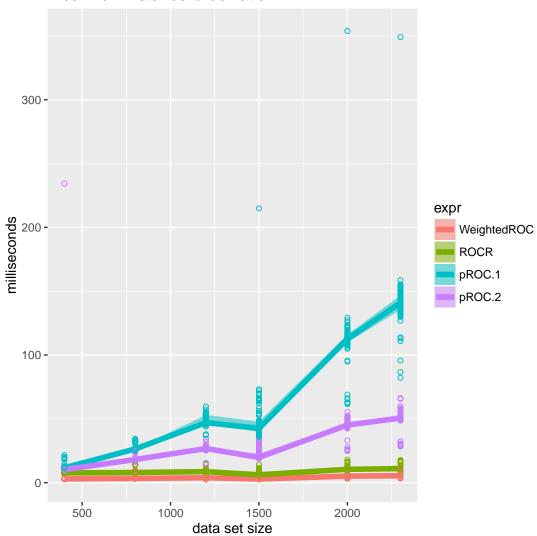
# 5 Scaling

In this section I was interested in seeing if there are any differences between algorithms as the number of data points changes.

```
> if(require(microbenchmark)){
+    stats.by.size.expr <- list()
+    ms.by.size <- list()
+    for(size in c(400, 800, 1200, 1500, 2000, 2300)){
+       indices <- seq(1, length(set$label), l=size)
+      y <- set$label[indices]
+      y.hat <- guess[indices]
+      this.size <- microbenchmark(WeightedROC={
            wroc <- WeightedROC(y.hat, y)
+      }, ROCR={</pre>
```

```
pred <- prediction(y.hat, y)</pre>
        perf <- performance(pred, "tpr", "fpr")</pre>
      }, pROC.1={
        proc <- roc(y, y.hat, algorithm=1)</pre>
      }, pROC.2={
        proc <- roc(y, y.hat, algorithm=2)</pre>
      })
      this.size$milliseconds <- this.size$time/1e6
      ms.by.size[[paste(size)]] <- data.frame(size, this.size)</pre>
      this.by.expr <- split(this.size, this.size$expr)</pre>
      for(expr in names(this.by.expr)){
        stats <- with(this.by.expr[[expr]], {</pre>
          data.frame(median=median(milliseconds),
+
                      q25=quantile(milliseconds, 0.25),
                      q75=quantile(milliseconds, 0.75))
+
        })
+
        stats.by.size.expr[[paste(size, expr)]] <- data.frame(size, expr, stats)</pre>
+
      }
    }
+
    ms <- do.call(rbind, ms.by.size)</pre>
    algo.stats <- do.call(rbind, stats.by.size.expr)</pre>
    timePlot <- ggplot()+</pre>
      geom_ribbon(aes(size, ymin=q25, ymax=q75, fill=expr),
                   data=algo.stats, alpha=1/2)+
      geom_line(aes(size, median, color=expr), data=algo.stats, size=2)+
      geom_point(aes(size, milliseconds, color=expr), data=ms, pch=1)+
      ylab("milliseconds")+
      xlab("data set size")+
      ggtitle("mean +/- 1 standard deviation")
    print(timePlot)
```





The figure above shows that for the spam data, the data set size does not affect the speed ordering of the algorithms for ROC curve computation. In all cases, WeightedROC is fastest, followed by ROCR, then pROC.2, then pROC.1. It makes sense that pROC.2 is faster than pROC.1, since pROC.2 uses the cumsum function, but pROC.1 does not.