# STAT 775: Machine Learning HW 01

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## Abstract

Exploration of K-Nearest Neighbor (k-NN), Linear Regression, and associated topics.

## 1 Exercise 2.1 (2 points)

Suppose each of K-classes has an associated target  $t_k$ , which is a vector of all zeros, except a one in the  $k^{th}$  position. Show that classifying to the largest element of  $\hat{y}$  amounts to choosing the closest target,  $\min_k ||t_k - \hat{y}||$ , if the elements of  $\hat{y}$  sum to one.

### Solution:

Let us consider the difference between the distance between  $\hat{y}$  and the target  $t_m$  corresponding to  $\hat{y}$ 's greatest member value,  $\hat{y}_m$ , call it  $\hat{y}_m$ , and the distance between  $\hat{y}$  and any other target  $t_n$ .

First, let us consider  $\hat{y}_m$  and  $\hat{y}_l$ . By definition,  $\hat{y}_m \geq \hat{y}_l$  for any l. Second, since the signs of these differences will be the same, let us just consider both distances to be positive. This will let us consider distance values  $d: d \in [0,1]$ , and, given the monotonicity of x and  $x^2$  in the range  $[0,\infty)$ , we can safely consider the Euclidean distance squared ( $||x||^2$ ), in place of normal Euclidean distance.

Next, we compare the difference of the distances described above:

$$\begin{aligned} \|\hat{y} - t_m\|^2 - \|\hat{y} - t_n\|^2 &= \\ \sqrt{(\hat{y}_m - 1)^2 + (\hat{y}_n - 0)^2 + \sum_{i,i \neq m,n} (\hat{y}_i - 0)^2}^2 - \sqrt{(\hat{y}_m - 0)^2 + (\hat{y}_n - 1)^2 + \sum_{i,i \neq m,n} (\hat{y}_i - 0)^2}^2 \\ &= (\hat{y}_m - 1)^2 + \hat{y}_n^2 + \sum_{i,i \neq m,n} (\hat{y}_i - 0)^2 - \hat{y}_m^2 - (\hat{y}_n - 1)^2 - \sum_{i,i \neq m,n} (\hat{y}_i - 0)^2 \\ &= (\hat{y}_m - 1)^2 + \hat{y}_n^2 - \hat{y}_m^2 - (\hat{y}_n - 1)^2 \\ &= \hat{y}_m^2 - 2\hat{y}_m + 1 - \hat{y}_m^2 - \hat{y}_n^2 + 2\hat{y}_n - 1 + \hat{y}_n^2 \\ &= -2\hat{y}_m + 2\hat{y}_n \\ &= 2(\hat{y}_n - \hat{y}_m) \\ &\leq 0 \text{ since } \hat{y}_m \geq \hat{y}_l \end{aligned}$$

Thus, the distance between  $\hat{y}$  and  $t_m$  must be less than or equal to that of the distance between  $\hat{y}$  and  $t_n$ . Therefore, classifying to the largest element of  $\hat{y}$  amounts to choosing the closest target,  $\min_k ||t_k - \hat{y}||$ .

## 2 Exercise 2.3 (4 points)

Derive equation (2.24).  $(d(p, N) = (1 - \frac{1}{2}^{1/N})^{1/p})$ 

Solution:

This question refers to our consideration of a p-dimensional unit "ball" centered about the origin and the median distance from the origin to the closest of any of N points uniformly distributed in the ball. For the sake of alleviating confusion, I believe that "uniformly distributed" means that the points are distributed where each coordinate of the point follows a U(-1,1) distribution, NOT that the points are distributed to be spaced evenly from one another.

Let us denote the median distance from the origin to the closest of any of the N points as d (short for d(p, N)). Let us call any of the N points as  $a_i$ . For simplicity, let the notation  $\|\cdot\|$  mean the Euclidean distance between a point and the origin.

By the definition of the median, half of the  $x_i$ s should be further from the origin than d. Since the points are distributed indepently, we can say that:

$$\frac{1}{2} = \prod_{i=1}^{N} P(\|a_i\| > d)$$

We can also say that the likelihood of a point being randomly assigned to a certain region of the ball is the same as the fraction of the volume of the whole ball that region occupies. So we can estimate the probability that a point is further from the origin than d (via its complement) by:

$$P(||a_i|| > d) = 1 - P(||a_i|| \le d)$$

$$= 1 - \frac{kd^p}{k(1)^p}$$

$$= 1 - \frac{kd^p}{k}$$

$$= 1 - d^p$$

where k is the constant used in determining the volume of hypershperes (for p=2 (a circle),  $k=\pi$ ; for p=3 (a sphere),  $k=\frac{4}{3}\pi$ ; and so on...). Also, recall that the ball is a unit sphere with a radius of 1.

Since  $P(||a_i|| > d) = 1 - d^p$ , we have:

$$\frac{1}{2} = \prod_{i=1}^{N} 1 - d^{p}$$

and solving for d, we get:

$$\prod_{i=1}^{N} 1 - d^p = \frac{1}{2}$$

$$(1 - d^p)^N = \frac{1}{2}$$

$$1 - d^p = (\frac{1}{2})^{\frac{1}{N}}$$

$$d^p = 1 - (\frac{1}{2})^{\frac{1}{N}}$$

$$d(p, N) = (1 - (\frac{1}{2})^{\frac{1}{N}})^{\frac{1}{p}}$$

And so the derivation is complete.

# 3 Exercise 2.4 (4 points)

The edge effect problem discussed on page 23 is not peculiar to uniform sampling from bounded domains. Consider inputs drawn from a spherical multinormal distribution  $X \sim N(0, \mathbf{I}_p)$ . The squared distance from any sample point to the origin has a  $\chi_p^2$  distribution with mean p. Consider a prediction point  $x_0$  drawn from this distribution, and let  $a = \frac{x_0}{\|x_0\|}$  be an associated unit vector. Let  $z_i = a^T x_i$  be the projection of each of the training points on this direction. Show that the  $z_i$  are distributed N(0,1) with expected squared distance from the origin 1, while the target point has expected squared distance p from the origin. Hence for p = 10, a randomly drawn test point is about 3.1 standard deviations from the origin, while all the training points are on average one standard deviation along direction a. So most prediction points see themselves as lying on the edge of the training set.

### Solution:

Since  $z_i$  are the projections of the  $x_i$  onto a, the  $z_i$  are linear combinations of variables that follow an  $N(0, \mathbf{1})$  distribution, so the  $z_i$  themselves are normal with 0 mean and a variance of

$$Var(z_i) = ||a^T||^2 Var(x_i) = 1 * Var(x_i) = 1$$

since a is a unit vector and the  $x_i$  have variance of 1 by the problem statement.

# 4 Exercise 2.8 (10 points)

Compare the classification performance of linear regression and k-nearest neighbor classification on the zipcode data. In particular, consider only the 2's and 3's, and k = 1, 3, 5, 7 and 15. Show both the training and test error for each choice. The zipcode data are available from the book website www-stat.stanford.edu/ElemStatLearn.

### Solution:

It would appear that k-NN is the better classifier. It only performed marginally worse than linear regression on the training data (which is not of practical significance anyway), but on the novel test data, k-NN consistently performed better than the regression - and by a wider margin.

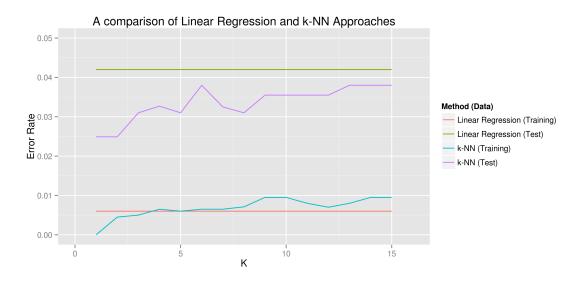


Figure 1: A graphical comparison of the errors of the Linear Regression and the k-NN classification methods.

The R code used is as follows:

```
##
# ESL Exercise 2.8
##
########################
# some initial setup #
#######################
working.directory = "~/Desktop/STAT775/HW01/Exercise_02_08"
training.file.name = "zip.test"
test.file.name = "zip.train"
setwd(working.directory)
require(ggplot2)
require(reshape)
####################
# Data Preparation #
####################
read.and.filter.data <- function(file.name) {</pre>
```

```
#
  # Args:
  # Returns:
  data = read.table(
    file.name,
    header = F,
    row.name = NULL
  data = subset(data, V1 == 2 | V1 == 3)
  return(data)
}
separate.classes <- function(x) {</pre>
  if (x == 3) {
    return(1)
  } else {
    return(-1)
  }
}
zip.train <- read.and.filter.data(training.file.name)</pre>
training.data <- data.matrix(zip.train[, -1])</pre>
training.targets <- data.matrix(zip.train[, 1])</pre>
for (i in 1:length(training.targets)) {
  training.targets[i] <- separate.classes(training.targets[i])</pre>
}
zip.test <- read.and.filter.data(test.file.name)</pre>
test.data <- data.matrix(zip.train[, -1])</pre>
test.targets <- data.matrix(zip.train[, 1])</pre>
for (i in 1:length(test.targets)) {
  test.targets[i] <- separate.classes(test.targets[i])</pre>
}
k.range <- 1:15 # simpler than doing just 1, 3, 5, 7, 15
determine.category <- function(x) {</pre>
  if(x >= 0) {
    return(1)
  } else {
    return(-1)
  }
}
#####################
# Linear Regression #
###################
least.squares <- function(x, y) {</pre>
  xt.x \leftarrow t(x) %% x
  xt.x.inv <- solve(xt.x)
  full.mat <- xt.x.inv %*% t(x)</pre>
  regression.weights <- full.mat %*% y
```

```
return(regression.weights)
}
linear.predict <- function(inputs, betas) {</pre>
  \# sometimes the formula is listed as x^t * b, but our 'inputs' data already has each
  # test vector in a row when we pass it as a matrix (for this script)
 result <- inputs %*% betas
  result <- lapply(result, function(x) {return(determine.category(x))})
 return(result)
digit.model.weights <- least.squares(x = training.data, y = training.targets)
linear.training.predictions <- list()</pre>
for(i in 1:length(training.targets)) {
  linear.training.predictions[i] <- linear.predict(inputs = training.data[i, ], betas = digit.model.wei
linear.test.predictions <- list()</pre>
for(i in 1:length(test.targets)) {
  linear.test.predictions[i] <- linear.predict(inputs = test.data[i, ], betas = digit.model.weights)</pre>
}
#######
# K-NN #
#######
euclidean.distance <- function(x, y) {</pre>
 diff \leftarrow x - y
  squares <- diff^2
  return(sqrt(sum(squares)))
}
my.knn.predict <- function(input.point, training.points, training.labels, k) {
  distances = NULL
  for (i in 1:length(training.points[, 1])) {
    distances[i] <- euclidean.distance(input.point, training.points[i, ])</pre>
  }
  category <- training.labels</pre>
  results <- data.frame(distances, category)</pre>
 results <- results[order(-distances), ]</pre>
  results <- results$category[1:k]
  return(determine.category(mean(results)))
}
knn.training.predictions <- matrix(0, length(k.range), length(training.targets))
knn.test.predictions <- matrix(0, length(k.range), length(test.targets))</pre>
for (k in k.range) {
  for (j in 1:length(training.targets)) {
    knn.training.predictions[k, j] <- my.knn.predict(</pre>
      input.point = training.data[j, ],
      training.points = training.data,
      training.labels = training.targets,
```

```
k = k
    )
 }
}
for (k in k.range) {
 for (j in 1:length(test.targets)) {
    knn.test.predictions[k, j] <- my.knn.predict(</pre>
      input.point = test.data[j, ],
      training.points = training.data,
      training.labels = training.targets,
      k = k
    )
 }
}
#############################
# Error Computations #
#########################
compute.error.rate <- function(targets, predictions) {</pre>
  incorrect.predictions = 0
  for (i in 1:length(targets)) {
    if (targets[1] != predictions[i]) {
      incorrect.predictions <- (incorrect.predictions + 1)</pre>
    }
  }
 return(incorrect.predictions / length(targets))
linear.training.error.rates <- compute.error.rate(training.targets, linear.training.predictions)</pre>
linear.training.error.rates <- rep(linear.training.error.rates, 15)</pre>
linear.test.error.rates <- compute.error.rate(test.targets, linear.test.predictions)</pre>
linear.test.error.rates <- rep(linear.test.error.rates, 15)</pre>
knn.training.error.rates <- NULL</pre>
for (k in k.range) {
 knn.training.error.rates[k] <- compute.error.rate(training.targets, knn.training.predictions[k, ])
}
knn.test.error.rates <- NULL</pre>
for (k in k.range) {
 knn.test.error.rates[k] <- compute.error.rate(test.targets, knn.test.predictions[k, ])
}
errors.for.plotting <- data.frame(</pre>
  "K" = 1:15,
  "Linear (Training)" = linear.training.error.rates,
  "Linear (Test)" = linear.test.error.rates,
  "KNN (Training)" = knn.training.error.rates,
  "KNN (Test)" = knn.test.error.rates
)
############
# Plotting #
```

## ###########

```
plot.data <- melt(errors.for.plotting, id = "K", variable_name = "Method")</pre>
ggplot(data = plot.data, aes(x = K, y = value, color = Method)) +
  geom_line() +
  ggtitle("A comparison of Linear Regression and k-NN Approaches") +
 xlab("K") +
  ylab("Error Rate") +
  ylim(0, 0.05) +
  xlim(0, 16) +
  scale_color_hue(
    name = "Method (Data)",
    labels = c(
      "Linear Regression (Training)",
      "Linear Regression (Test)",
      "k-NN (Training)",
      "k-NN (Test)"
    )
  )
ggsave('error_plot.png')
```